Preface

This book is an introduction to data structures and algorithms for functional languages, with a focus on proofs. It covers both functional correctness and running time analysis. It does so in a unified manner with inductive proofs about functional programs and their running time functions.

The unique feature of this book is that all the proofs have been machine-checked by the proof assistant Isabelle. That is, in addition to the text in the book, which requires no knowledge of proof assistants, there are the Isabelle definitions and proofs that can be accessed by following (in the PDF file) the links attached to section headings with a symbol. The structured nature of Isabelle proofs permits even novices to browse them and follow the high-level arguments.

This book has been classroom-tested for a number of years in a course for graduate and advanced undergraduate students. At the same time it is a reference for programmers and researchers who are interested in the details of some algorithm or proof.

Isabelle

Isabelle [Nipkow et al. 2002, Paulson 1989, Wenzel 2002] is a proof assistant for the logic HOL (= Higher-Order Logic), which is why the system is often called Isabelle/HOL. HOL is a generalization of first-order logic: functions can be passed as parameters and returned as results, just as in functional programming, and they can be quantified over. Isabelle’s version of HOL also supports a simple version of Haskell’s type classes.

The main strength of Isabelle and other proof assistants is their trustworthiness: all definitions, lemma statements, and inferences are checked. Beyond trustworthiness, formal proofs can also clarify arguments, by exposing and explaining difficult steps. Most Isabelle users will confirm that their pen-and-paper proofs have become more lucid, and more correct, after they subjected themselves to the discipline of formal proof.

As emphasized above, the reader need not be familiar with Isabelle or HOL in order to read this book. However, to take full advantage of our proof assistant approach, readers are encouraged to learn how to write Isabelle definitions and proofs themselves — and to solve some of the exercises in this book. To this end we recommend the
tutorial *Programming and Proving in Isabelle/HOL* [Nipkow], which is also Part I of the book *Concrete Semantics* [Nipkow and Klein 2014].

**Prerequisites**

We expect the reader to be familiar with

- the basics of discrete mathematics: propositional and first-order logic, sets and relations, proof principles including induction;
- a typed functional programming language like Haskell [Haskell], OCaml [OCaml] or Standard ML [Paulson 1996];
- simple inductive proofs about functional programs.

**Under Development**

This book is meant to grow. New chapters are meant to be added over time. The list of authors is meant to grow — you could become one of them!

**Colour**

For the quick orientation of the reader, definitions are displayed in coloured boxes:

- These boxes display functional programs.
- These boxes display auxiliary definitions.

From a logical point of view there is no difference between the two kinds of definitions except that auxiliary definitions need not be executable.

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# Contents

1 Basics 1

I Sorting and Selection 11

2 Sorting 13
3 Selection 31

II Search Trees 45

4 Binary Trees 47
5 Binary Search Trees 59
6 Abstract Data Types 77
7 2-3 Trees 85
8 Red-Black Trees 95
9 AVL Trees 105
10 Beyond Insert and Delete: $\cup$, $\cap$ and $-$ 117
11 Arrays via Braun Trees 127
12 Tries 149

III Priority Queues 161

13 Priority Queues 163
14 Leftist Heaps 167
15 Priority Queues via Braun Trees 175
16 Binomial Heaps 179

IV Advanced Design and Analysis Techniques 189
17 Dynamic Programming 191
18 Amortized Analysis 213
19 Queues 219
20 Splay Trees 231
21 Skew Heaps 239
22 Pairing Heaps 243

V Selected Topics 251
23 Knuth–Morris–Pratt String Search 255
24 Huffman’s Algorithm 267
25 Alpha-Beta Pruning 281

VI Appendix 303
A List Library 305
B Time Functions 309
C Notation 317
Bibliography 323
Authors 331
Index 332
1 Basics

Tobias Nipkow

In this chapter we describe the basic building blocks the book rests on.

**Programs:** The functional programming language we use is merely sketched because of its similarity with other well known functional languages.

**Predefined types and notation:** We introduce the basic predefined types and notations used in the book.

**Inductive proofs:** Although we do not explain proofs in general, we make an exception for certain inductive proofs.

**Running time:** We explain how we model running time by step counting functions.

### 1.1 Programs

The programs in this book are written in Isabelle's functional programming language which provides recursive algebraic data types (keyword: `datatype`), recursive function definitions and `let`, `if` and `case` expressions. The language is sufficiently close to a number of similar typed functional languages (SML [Paulson 1996], OCaml [OCaml], Haskell [Haskell]) to obviate the need for a detailed explanation. Moreover, Isabelle can generate SML, OCaml, Haskell and Scala code [Haftmann b]. What distinguishes Isabelle's functional language from ordinary programming languages is that all functions in Isabelle must terminate. Termination must be proved. For most of the functions in this book, termination is not difficult to see and Isabelle can prove it automatically. (For details on termination proofs, consult the function definition tutorial [Krauss].)

Isabelle's functional language is pure logic. All language elements have precise definitions. However, this book is about algorithms, not programming language semantics. A functional programmer's intuition suffices for reading it. (If you want to know more about the logical basis of recursive data types, recursive functions and code generation: see [Berghofer and Wenzel 1999, Haftmann and Nipkow 2010, Krauss 2006].)

A useful bit of notation: any infix operator can be turned into a function by enclosing it in parentheses, e.g. \((+\)).
1.2 Types

Type variables are denoted by ‘a, ‘b, etc. The function type arrow is \( \Rightarrow \). Type constructor names follow their argument types, e.g. ‘a list. The notation \( t :: \tau \) means that term \( t \) has type \( \tau \). The following types are predefined.

**Booleans** Type bool comes with the constants True and False and the usual operations. We mostly write \( = \) instead of \( \leftrightarrow \).

**Numbers** There are three numeric types: the natural numbers \( \text{nat} \) (0, 1, ...), the integers \( \text{int} \) and the real numbers \( \text{real} \). They correspond to the mathematical sets \( \mathbb{N} \), \( \mathbb{Z} \) and \( \mathbb{R} \) and not to any machine arithmetic. All three types come with the usual (overloaded) operations.

**Sets** The type ‘a set of sets (finite and infinite) over type ‘a comes with the standard mathematical operations. The minus sign “−”, unary or binary, can denote set complement or difference.

**Lists** The type ‘a list of lists whose elements are of type ‘a is a recursive data type:

```plaintext
datatype ‘a list = Nil | Cons ‘a (‘a list)
```

Constant Nil represents the empty list and Cons \( x \) \( xs \) represents the list with first element \( x \), the head, and rest list \( xs \), the tail. The following syntactic sugar is sprinkled on top;

\[
\begin{align*}
[] & \equiv \text{Nil} \\
x \ # \ xs & \equiv \text{Cons} \ x \ xs \\
[x_1, \ldots, x_n] & \equiv \ x_1 \ # \ldots \ # \ x_n \ # \ []
\end{align*}
\]

The \( \equiv \) symbol means that the left-hand side is merely an abbreviation of the right-hand side.

A library of predefined functions on lists is shown in Appendix A. The length of a list \( xs \) is denoted by \( |xs| \).

**Type ‘a option** The data type ‘a option is defined as follows:

```plaintext
datatype ‘a option = None | Some ‘a
```

**Pairs and Tuples** Pairs are written \( (a, b) \). Functions \( \text{fst} \) and \( \text{snd} \) select the first and second component of a pair: \( \text{fst} (a, b) = a \) and \( \text{snd} (a, b) = b \). The type unit contains only a single element (), the empty tuple.
1.2 Types

Functions 'a ⇒ 'b come with a predefined pointwise update operation with its own notation:

\[ f(a := b) = (\lambda x. \text{if } x = a \text{ then } b \text{ else } f) \]

1.2.1 Pattern Matching

Functions are defined by equations and pattern matching, for example over lists. Natural numbers may also be used in pattern-matching definitions:

\[ \text{fib } (n + 2) = \text{fib } (n + 1) + \text{fib } n \]

 Occasionally we use an extension of pattern matching where patterns can be named. For example, the defining equation

\[ f(x \# (y \# zs := ys)) = ys \@ zs \]

introduces a variable ys on the left that stands for y \# zs and can be referred to on the right. Logically it is just an abbreviation of

\[ f(x \# y \# zs) = (\text{let } ys = y \# zs \text{ in } ys \@ zs) \]

although it is suggestive of a more efficient interpretation. The general format is pattern =: variable.

1.2.2 Numeric Types and Coercions

The numeric types nat, int and real are all distinct. Converting between them requires explicit coercion functions, in particular the inclusion functions int :: nat ⇒ int and real :: nat ⇒ real that do not lose any information (in contrast to coercions in the other direction). We do not show inclusions unless they make a difference. For example, (m + n) :: real, where m, n :: nat, is mathematically unambiguous because real (m + n) = real m + real n. On the other hand, (m − n) :: real is ambiguous because real (m − n) ≠ real m − real n because (0::nat) − n = 0. In some cases we can also drop coercions that are not inclusions, e.g. nat :: int ⇒ nat, which coerces negative integers to 0: if we know that i ≥ 0 then we can drop the nat in nat i.

We prefer type nat over type real for ease of (Isabelle) proof. For example, for m, n :: nat we prefer m ≤ 2^n over lg m ≤ n. Function lg is the binary logarithm.

1.2.3 Multisets

Informally, a multiset is a set where elements can occur multiple times. Multisets come with the following operations:
{ } :: 'a multiset
( ∈ₐ) :: 'a ⇒ 'a multiset ⇒ bool
add_mset :: 'a ⇒ 'a multiset ⇒ 'a multiset
(+ ) :: 'a multiset ⇒ 'a multiset ⇒ 'a multiset
size :: 'a multiset ⇒ nat
mset :: 'a list ⇒ 'a multiset
set_mset :: 'a multiset ⇒ 'a set
image_mset :: ('a ⇒ 'b) ⇒ 'a multiset ⇒ 'b multiset
filter_mset :: ('a ⇒ bool) ⇒ 'a multiset ⇒ 'a multiset
sum_mset :: 'a multiset ⇒ 'a

Their meaning: { } is the empty multiset; ( ∈ₐ) is the element test; add_mset adds an
element to a multiset; (+ ) is the sum of two multisets, where multiplicities of elements
are added; size M, written |M|, is the number of elements in M, taking multiplicities
into account; mset converts a list into a multiset by forgetting about the order of
elements; set_mset converts a multiset into a set; image_mset applies a function to
all elements of a multiset; filter_mset removes all elements from a multiset that do
not satisfy the given predicate; sum_mset is the sum of the values of a multiset, the
iteration of (+ ) (taking multiplicity into account).

We use some additional suggestive syntax for some of these operations:

\{ x ∈ₐ M | P x\} ≡ filter_mset P M
\{ f x | x ∈ₐ M\} ≡ image_mset f M
∑ₓ M ≡ sum_mset M
∑ₓ∈ₐ M f x ≡ sum_mset (image_mset f M)

See Section C.3 in the appendix for an overview of such syntax.

1.3 Notation
We deviate from Isabelle’s notation in favour of standard mathematics in a number
of points:

- There is only one implication: \( \rightarrow \) is printed as \( \rightarrow \) and \( P \rightarrow Q \rightarrow R \) is
  printed as \( P \land Q \rightarrow R \).
- Multiplication is printed as \( x \cdot y \).
- Exponentation is uniformly printed as \( x^y \).
- We sweep under the carpet that type \( nat \) is defined as a recursive data type:
datatype nat = 0 | Suc nat. In particular, constructor Suc is hidden: Suc \( ^k \) 0 is
  printed as \( k \) and Suc \( ^k \) n (where \( n \) is not 0) is printed as \( n + k \).
1.4 Proofs

Proofs are the raison d'être of this book. Thus we present them in more detail than is customary in a book on algorithms. However, not all proofs:

- We omit proofs of simple properties of numbers, lists, sets and multisets, our predefined types. Obvious properties (e.g. \(|zs @ ys| = |zs| + |ys|\) or commutativity of \(\cup\)) are used implicitly without proof.
- With some exceptions, we only state properties if their proofs require induction, in which case we will say so, and we will always indicate which supporting properties were used.
- If a proposition is simply described as "inductive" or its proof is described by a phrase like "by an easy/automatic induction" it means that in the Isabelle proofs all cases of the induction were automatic, typically by simplification.

As a simple example of an easy induction consider the append function

\[
(\@) :: 'a list \Rightarrow 'a list \Rightarrow 'a list
\]

\[
[] @ ys = ys
\]

\[
(x @ xs) @ ys = x @ xs @ ys
\]

and the proof of \((xs @ ys) @ zs = xs @ ys @ zs\) by structural induction on \(xs\). (Note that \((\@)\) associates to the right.) The base case is trivial by definition: \([[] @ ys] @ zs = [] @ ys @ zs\). The induction step is easy:

\[
(x @ xs @ ys) @ zs
\]

\[
= x # (xs @ ys) @ zs \quad \text{by definition of \((\@)\)}
\]

\[
= x @ xs @ ys @ zs \quad \text{by IH}
\]

Note that \(\text{IH}\) stands for Induction Hypothesis, in this case \((xs @ ys) @ zs = xs @ ys @ zs\).

1.4.1 Computation Induction

Because most of our proofs are about recursive functions, most of them are by induction, and we say so explicitly. If we do not state explicitly what form the induction takes, it is by an obvious structural induction. The alternative and more general induction schema is computation induction where the induction follows
the terminating computation, but from the bottom up. For example, the terminating recursive definition for \( \text{gcd} :: \text{nat} \Rightarrow \text{nat} \Rightarrow \text{nat} \)

\[
\text{gcd} \ m \ n = (\text{if } n = 0 \text{ then } m \text{ else } \text{gcd} \ n \ (m \mod n))
\]
gives rise to the following induction schema:

If \((n \neq 0 \Rightarrow P n \ (m \mod n)) \Rightarrow P m \ n\) (for all \(m\) and \(n\)),
then \(P m \ n\) (for all \(m\) and \(n\)).

In general, let \(f :: \tau \Rightarrow \tau'\) be a terminating function of, for simplicity, one argument. Proving \(P(x :: \tau)\) by induction on the computation of \(f\) means proving

\[P \ r_1 \land \ldots \land P \ r_n \Rightarrow P \ e\]

for every defining equation

\[f \ e = \ldots \ f \ r_1 \ldots \ f \ r_n \ldots\]

where \(f \ r_1, \ldots, f \ r_n\) are all the recursive calls. For simplicity we have ignored the if and case contexts that a recursive call \(f \ r_i\) occurs in and that should be preconditions of the assumption \(P \ r_i\) as in the \(\text{gcd}\) example. If the defining equations for \(f\) overlap, the above proof obligations are stronger than necessary.

### 1.5 Running Time

Our approach to reasoning about the running time of a function \(f\) is very simple: we explicitly define a function \(T_f\) such that \(T_f x\) models the time the computation of \(f x\) takes. More precisely, \(T_f\) counts the number of non-primitive function calls in the computation of \(f\). It is not intended that \(T_f\) yields the exact running time but only that the running time of \(f\) is in \(O(T_f)\).

Given a function \(f :: \tau_1 \Rightarrow \ldots \Rightarrow \tau_n \Rightarrow \tau\) we define a (running) time function \(T_f :: \tau_1 \Rightarrow \ldots \Rightarrow \tau_n \Rightarrow \text{nat}\) by translating every defining equation for \(f\) into a defining equation for \(T_f\). The translation is defined by two functions: \(E\) translates defining equations for \(f\) to defining equations for \(T_f\) and \(T\) translates expressions that compute some value to expressions that computes the number of function calls. The unusual notation \(E[\_]\) and \(T[\_]\) emphasizes that they are not functions in the logic.

\[
\begin{align*}
E[f \ p_1 \ldots \ p_n = e] & = (T_f \ p_1 \ldots \ p_n - T[e] + 1) \\
T[g \ e_1 \ldots \ e_k] & = T[e_1] + \ldots + T[e_k] + T_g \ e_1 \ldots \ e_k
\end{align*}
\] (1.1)

This is the general idea. It requires some remarks and clarifications:

- This definition of \(T_f\) is an abstraction of a call-by-value semantics. Thus it is also correct for lazy evaluation but may be a very loose upper bound.
• Definition (1.1) is incomplete: if \( g \) is a variable or constructor function (e.g. \( \text{Nil} \) or \( \text{Cons} \)), then there is no defining equation and thus no \( T_g \). Conceptually we define \( T_g : = 0 \) if \( g \) is a variable, constructor function or predefined function on \( \text{bool} \) or numbers. That is, we count only user-defined function calls. This does not change \( O(T_f) \) for user-defined functions \( f \) (see Discussion below).

• \textbf{if, case} and \textbf{let} are treated specially:

\[
\begin{align*}
T[\text{if } b \text{ then } e_1 \text{ else } e_2] &= T[b] + (\text{if } b \text{ then } T[e_1] \text{ else } T[e_2]) \\
T[\text{case } e \text{ of } p_1 \Rightarrow e_1 | \ldots | p_k \Rightarrow e_k] &= T[e] + (\text{case } e \text{ of } p_1 \Rightarrow T[e_1] | \ldots | p_k \Rightarrow T[e_k]) \\
T[\text{let } x = e_1 \text{ in } e_2] &= T[e_1] + (\text{let } x = e_1 \text{ in } T[e_2])
\end{align*}
\]

• For simplicity we restrict ourselves to a first-order language above. Nevertheless we use a few basic higher-order functions like \texttt{map} in the book. Their running time functions are defined in Appendix B.1.

As an example consider the append function \((@)\) defined above. The defining equations for \( T_{\text{append}} :: \text{a list} \Rightarrow \text{a list} \Rightarrow \text{nat} \) are easily derived. The first equation translates like this:

\[
\begin{align*}
E[\text{[] @ } ys = ys] &= (T_{\text{append} \text{[]}} ys = T[ys] + 1) \\
&= (T_{\text{append} \text{[]}} ys = 1)
\end{align*}
\]

The right-hand side of the second equation translates like this:

\[
\begin{align*}
T[x \# xs @ ys] &= T[x] + T[ys @ xs] + T_{\text{Cons}} x (xs @ ys) \\
&= 0 + (T[xs] + T[ys] + T_{\text{append}} xs ys) + 1 \\
&= 0 + (0 + 0 + T_{\text{append}} xs ys) + 1
\end{align*}
\]

Thus the two defining equations for \( T_{\text{append}} \) are

\[
\begin{align*}
T_{\text{append} \text{[]}} ys &= 1 \\
T_{\text{append}} (x \# xs) ys &= T_{\text{append}} xs ys + 1
\end{align*}
\]

As a final simplification, we drop the +1 in the time functions for non-recursive functions (think inlining). In that case \( E[f x_1 \ldots x_n = e] = (T f x_1 \ldots x_n = T[e]) \). Again, this does not change \( O(T_f) \) (except in the trivial case where \( T[e] = 0 \)).
In the main body of the book we initially show the definition of each $T_f$. Once the principles above have been exemplified sufficiently, the time functions are relegated to Appendix B.

The definition of $T_f$ from the definition of $f$ has been automated in Isabelle.

1.5.1 Example: List Reversal

This section exemplifies not just the definition of time functions but also their analysis. The standard list reversal function $rev$ is defined in Appendix A. This is the corresponding time function:

$$T_{\text{rev}} :: 'a \text{ list} \Rightarrow \text{nat}$$

$$T_{\text{rev}} [] = 1$$

$$T_{\text{rev}} (x \# xs) = T_{\text{rev}} xs + T_{\text{append}} (\text{rev} \: xs) \; [x] + 1$$

A simple induction shows $T_{\text{append}} \: xs \: ys = |xs| + 1$. The precise formula for $T_{\text{rev}}$ is less immediately obvious (exercise!) but an upper bound is easy to guess and verify by induction:

$$T_{\text{rev}} \: xs \leq (|xs| + 1)^2$$

We will frequently prove upper bounds only.

Of course one can also reverse a list in linear time:

$$itrev :: 'a \text{ list} \Rightarrow 'a \text{ list} \Rightarrow 'a \text{ list}$$

$$itrev [] \: ys = ys$$

$$itrev (x \# xs) \: ys = itrev \: xs \: (x \# ys)$$

Function $itrev$ has linear running time: $T_{\text{itrev}} \: xs \: ys = |xs| + 1$. A simple induction yields $itrev \: xs \: ys = rev \: xs @ ys$. Thus $itrev$ implements $rev :: rev \: xs = itrev \: xs \: []$.

1.5.2 Discussion

Analysing the running time of a program requires a precise cost model. For imperative programs the standard model is the Random Access Machine (RAM) where each instruction takes one time unit. For functional programs a standard measure is the
number of function calls. We follow Sands [1990, 1995] by counting only non-primitive function calls. One could also count variable accesses, primitive and constructor function calls. This would not change $O(T_f)$ because it would only add a constant to each defining equation for $T_f$. However, it would make reasoning about $T_f$ more tedious.

A full proof that the execution time of our functional programs is in $O(T_f)$ on some actual soft- and hardware is a major undertaking: one would need to formalize the full stack of compiler, runtime system and hardware. We do not offer such a proof. Thus our formalization of “time” should be seen as conditional: given a stack that satisfies our basic assumptions in the definition of $E$ and $T$, our analyses are correct for that stack. Below we argue that these assumptions are reasonable (on a RAM) provided we accept that both the address space and numbers have a fixed size and cannot grow arbitrarily. Of course this means that actual program execution may abort if the resources are exhausted.

To simplify our argument, we assume that $T$ counts all function calls and variable accesses (which does not change $O(T_f)$ as we argued above). Thus our basic assumption is that function calls take constant time. This is reasonable (on a RAM) because we just need to allocate, initialize and later deallocate a stack frame of constant size. It is of constant size because all parameters are references or numbers and thus of fixed size. We also assumed that variable access takes constant time. This is a standard RAM assumption. Assuming that constructor functions take constant time is reasonable because the memory manager could simply employ a single reference to the first free memory cell and increment that with each constructor function call. How to account for garbage collection is less clear. In the worst case we have to assume that garbage collection is switched off, which simply exhausts memory more quickly. Finally we assume that operations on $bool$ and numbers take constant time. The former is obvious, the latter follows from our assumption that we have fixed-size numbers.

In the end, we are less interested in a specific model of time and more in the principle that time (and other resources) can be analyzed just as formally as functional correctness once the ground rules (e.g. $T$) have been established.

### 1.5.3 Asymptotic Notation

The above approach to running time analysis is nicely concrete and avoids the more sophisticated machinery of asymptotic notation, $O(.)$ and friends. Thus we have intentionally lowered the entry barrier to the book for readers who want to follow the Isabelle formalization: we require no familiarity with Isabelle’s real analysis library and in particular with the existing formalization of and automation for asymptotic notation [Eberl 2017b]. Of course this comes at a price: one has to come up with and reason about somewhat arbitrary constants in the analysis of individual functions.
Moreover we rarely appeal to the “master theorem” (although Eberl [2017b] provides a generalized version) but prove solutions to recurrence relations correct by induction. Again, this is merely to reduce the required mathematical basis and to show that it can be done. In informal explanations, typically when considering inessential variations, we do use standard mathematical notation and write, for example, $O(n \lg n)$.
Part I

Sorting and Selection
In this chapter we define and verify the following sorting functions: insertion sort, quicksort, and three variations of merge sort. We also analyze their running times (except for quicksort, whose running time analysis is beyond the scope of this book).

Sorting involves an ordering. We assume such an ordering to be provided by comparison operators ≤ and < defined on the underlying type.

Sortedness of lists is defined as follows:

\[
\text{sorted} :: (\forall a \cdot \text{linorder}(a)) \Rightarrow \text{list}(a) \Rightarrow \text{bool}
\]
\[
\text{sorted}[] = \text{True}
\]
\[
\text{sorted}(x \# ys) = ((\forall y \in \text{set}(ys). x \leq y) \land \text{sorted}(ys))
\]

That is, every element is ≤ to all elements to the right of it: the list is sorted in increasing order.

The notation \(\forall a \cdot \text{linorder}(a)\) restricts the type variable \(a\) to linear orders, which means that \text{sorted} is only applicable if a binary predicate \((\leq) :: \forall a \Rightarrow \forall a \Rightarrow \text{bool}\) is defined and \((\leq)\) is a linear order, i.e. the following properties are satisfied:

- Reflexivity: \(x \leq x\)
- Transitivity: \(x \leq y \land y \leq z \rightarrow x \leq z\)
- Antisymmetry: \(a \leq b \land b \leq a \rightarrow a = b\)
- Linearity/Totality: \(x \leq y \lor y \leq x\)

Moreover, the binary predicate \((<)\) must satisfy

\(x < y \leftrightarrow x \leq y \land x \neq y\).

On the numeric types \text{nat}, \text{int} and \text{real}, \((\leq)\) is a linear order.

Note that \text{linorder} is a specific predefined example of a \text{type class} [Haftmann a]. We will not explain type classes any further because we do not require the general concept. In fact, we will mostly not even show the \text{linorder} restriction in types: you can assume that if you see \(\leq\) or \(<\) on a generic type \(a\) in this book, \(a\) is implicitly restricted to \text{linorder}, unless we explicitly say otherwise.
2.1 Specification of Sorting Functions

A sorting function \( \text{sort} :: 'a \text{ list} \Rightarrow 'a \text{ list} \) (where, as usual, \( 'a::\text{linorder} \)) must obviously satisfy the following property:

\[
\text{sorted} (\text{sort} \; \text{xs})
\]

However, this is not enough — otherwise, \( \text{wrong_sort} \; \text{xs} = [] \) would be a correct sorting function. The set of elements in the output must be the same as in the input, and each element must occur the same number of times. This is most readily captured with the notion of a multiset (see Section 1.2.3). The second property that a sorting function \( \text{sort} \) must satisfy is

\[
\text{mset} (\text{sort} \; \text{xs}) = \text{mset} \; \text{xs}
\]

where function \( \text{mset} \) converts a list into its corresponding multiset.

2.2 Insertion Sort

Insertion sort is well-known for its intellectual simplicity and computational inefficiency. Its simplicity makes it an ideal starting point for this book. Below, it is implemented by the function \( \text{insort} \) with the help of the auxiliary function \( \text{insort1} \) that inserts a single element into an already sorted list.

\[
\begin{align*}
\text{insort1} &:: 'a \Rightarrow 'a \text{ list} \Rightarrow 'a \text{ list} \\
\text{insort1} \; x \; [] & = [x] \\
\text{insort1} \; x \; (y \# \; ys) & = (\text{if} \; x \leq y \; \text{then} \; x \# y \# \; ys \; \text{else} \; y \# \; \text{insort1} \; x \; ys)
\end{align*}
\]

\[
\begin{align*}
\text{insort} &:: 'a \text{ list} \Rightarrow 'a \text{ list} \\
\text{insort} \; [] & = [] \\
\text{insort} \; (x \# \; xs) & = \text{insort1} \; x \; (\text{insort} \; xs)
\end{align*}
\]

2.2.1 Functional Correctness

We start by proving the preservation of the multiset of elements:

\[
\begin{align*}
\text{mset} (\text{insort1} \; x \; \text{xs}) & = \{x\} + \text{mset} \; \text{xs} \quad (2.1) \\
\text{mset} (\text{insort} \; \text{xs}) & = \text{mset} \; \text{xs} \quad (2.2)
\end{align*}
\]

Both properties are proved by induction; the proof of (2.2) requires (2.1).

Now we turn to sortedness. Because the definition of \( \text{sorted} \) involves \( \text{set} \), it is frequently helpful to prove multiset preservation first (as we have done above) because that yields preservation of the set of elements. That is, from (2.1) we obtain:
2.2 Insertion Sort

\[ \text{set } (\text{insort} \ 1 \ x \ xs) = \{x\} \cup \text{set} \ xs \]  
(2.3)

Two inductions prove

\[ \text{sorted } (\text{insort} \ 1 \ a \ xs) = \text{sorted} \ xs \]  
(2.4)
\[ \text{sorted } (\text{insort} \ xs) \]  
(2.5)

where the proof of (2.4) uses (2.3) and the proof of (2.5) uses (2.4).

2.2.2 Running Time Analysis

These are the running time functions (according to Section 1.5):

\[
T_{\text{insort}1} :: 'a \Rightarrow 'a \ list \Rightarrow \text{nat} \\
T_{\text{insort}1} \ [] = 1 \\
T_{\text{insort}1} \ x \ (y \ # \ ys) = (\text{if } x \leq y \ \text{then} \ 0 \ \text{else} \ T_{\text{insort}1} \ x \ ys) + 1
\]

\[
T_{\text{insort}} :: 'a \ list \Rightarrow \text{nat} \\
T_{\text{insort}} \ [] = 1 \\
T_{\text{insort}} \ (x \ # \ xs) = T_{\text{insort}} \ xs + T_{\text{insort}1} \ x \ (\text{insort} \ xs) + 1
\]

A dismal quadratic upper bound for the running time of insertion sort is proved readily:

**Lemma 2.1.** \( T_{\text{insort}} \ xs \leq (|xs| + 1)^2 \)

**Proof.** The following properties are proved by induction on \( xs \):

\[ T_{\text{insort}1} \ x \ xs \leq |xs| + 1 \]  
(2.6)
\[ |\text{insort}1 \ x \ xs| = |xs| + 1 \]  
(2.7)
\[ |\text{insort} \ xs| = |xs| \]  
(2.8)

The proof of (2.8) needs (2.7). The proof of \( T_{\text{insort}} \ xs \leq (|xs| + 1)^2 \) is also by induction on \( xs \). The base case is trivial. The induction step is easy:

\[
T_{\text{insort}} \ (x \ # \ xs) = T_{\text{insort}} \ xs + T_{\text{insort}1} \ x \ (\text{insort} \ xs) + 1 \\
\leq (|xs| + 1)^2 + T_{\text{insort}1} \ x \ (\text{insort} \ xs) + 1 \quad \text{by IH} \\
\leq (|xs| + 1)^2 + |xs| + 1 + 1 \quad \text{using (2.6) and (2.8)} \\
\leq (|x \ # \ xs| + 1)^2
\]

Exercise 2.1 asks you to show that \text{insort} actually has quadratic running time on all lists \([n, n-1, \ldots, 0]\).
2.3 Quicksort

Quicksort [Hoare 1961] is a divide-and-conquer algorithm that sorts a list as follows:

- pick a pivot element from the list;
- partition the remaining list into those elements that are smaller and those that are greater than the pivot (equal elements can go into either sublist);
- sort these sublists recursively and append the results.

A particularly simple version of this approach, where the first element is chosen as the pivot, and the equal elements are put into the second sublist, looks like this:

\[
\text{quicksort} :: \text{'}a\text{ list} \Rightarrow \text{'}a\text{ list}
\]

\[
\text{quicksort} [] = []
\]

\[
\text{quicksort} (x \neq xs)
= \text{quicksort} (\text{filter} \ (\lambda y. \ y < x) \ xs) \@ [x] \@ \text{quicksort} (\text{filter} \ (\lambda y. \ y \geq x) \ xs)
\]

2.3.1 Functional Correctness

Preservation of the multiset of elements

\[
\text{mset} \ (\text{quicksort} \ xs) = \text{mset} \ xs
\]

is proved by computation induction using these lemmas:

\[
\text{mset} \ (\text{filter} \ P \ xs) = \text{filter}_\text{mset} \ P \ (\text{mset} \ xs)
\]

\[
(\forall x. P x = (\neg Q x)) \rightarrow \text{filter}_\text{mset} \ P \ M + \text{filter}_\text{mset} \ Q \ M = M
\]

A second computation induction proves sortedness

\[
\text{sorted} \ (\text{quicksort} \ xs)
\]

using the lemmas

\[
\text{sorted} \ (xs \@ \ ys) = (\text{sorted} \ xs \land \text{sorted} \ ys \land (\forall x \in \text{set} \ xs. \forall y \in \text{set} \ ys. x \leq y))
\]

\[
\text{set} \ (\text{quicksort} \ xs) = \text{set} \ xs
\]

where the latter one is an easy consequence of (2.9).

We do not analyze the running time of quicksort. It is well known that in the worst case it is quadratic (exercise!) but that the average-case running time (in a certain sense) is \(O(n \log n)\). If the pivot is chosen randomly instead of always choosing the first element, the expected running time is also \(O(n \log n)\). The necessary probabilistic analysis is beyond the scope of this text but can be found elsewhere [Eberl 2017a, Eberl et al. 2018].
2.4 Top-Down Merge Sort

Merge sort is another prime example of a divide-and-conquer algorithm, and one whose running time is guaranteed to be $O(n \lg n)$. We will consider three variants and start with the simplest one: split the list into two halves, sort the halves separately and merge the results.

merge :: 'a list ⇒ 'a list ⇒ 'a list
merge [] ys = ys
merge xs [] = xs
merge (x # xs) (y # ys) = (if x ≤ y then x # merge xs (y # ys) else y # merge (x # xs) ys)

msort :: 'a list ⇒ 'a list
msort xs = (let n = j xs j in if n ≤ 1 then xs else merge (msort (take (n div 2) xs)) (msort (drop (n div 2) xs)))

2.4.1 Functional Correctness

We start off with multisets and sets of elements:

\[ mset (merge xs ys) = mset xs + mset ys \tag{2.10} \]
\[ set (merge xs ys) = set xs \cup set ys \tag{2.11} \]

Proposition (2.10) is proved by induction on the computation of \( merge \) and (2.11) is an easy consequence.

**Lemma 2.2.** \( mset (msort xs) = mset xs \)

*Proof* by induction on the computation of \( msort \). Let \( n = |xs| \). The base case \((n ≤ 1)\) is trivial. Now assume \( n > 1 \) and let \( ys = \text{take } (n \text{ div } 2) \text{ xs} \) and \( zs = \text{drop } (n \text{ div } 2) \text{ xs} \).

\[
\begin{align*}
  mset (msort xs) &= mset (msort ys) + mset (msort zs) \quad \text{by (2.10)} \\
  &= mset ys + mset zs \quad \text{by IH} \\
  &= mset (ys @ zs) \\
  &= mset xs \\
\end{align*}
\]

Now we turn to sortedness. An induction on the computation of \( merge \), using (2.11), yields
Chapter 2  

\[ \text{sorted (merge } xs \ ys) = (\text{sorted } xs \land \text{sorted } ys) \]  

(2.12)

Lemma 2.3. \( \text{sorted (msort } xs) \) 

The proof is an easy induction on the computation of \( msort \). The base case \((n \leq 1)\) follows because every list of length \( \leq 1 \) is sorted. The induction step follows with the help of (2.12).

2.4.2 Running Time Analysis 

To simplify the analysis, and in line with the literature, we only count the number of comparisons:

\[ C_{\text{merge}} :: \text{a list } \Rightarrow \text{a list } \Rightarrow \text{nat} \]

\[ C_{\text{merge}} [] [] = 0 \]

\[ C_{\text{merge}} [] [] = 0 \]

\[ C_{\text{merge}} (x \# xs) (y \# ys) = 1 + (\text{if } x \leq y \text{ then } C_{\text{merge}} xs (y \# ys) \text{ else } C_{\text{merge}} (x \# xs) ys) \]

\[ C_{\text{msort}} :: \text{a list } \Rightarrow \text{nat} \]

\[ C_{\text{msort}} xs = (\text{let } n = |xs|; \]

\[ ys = \text{take } (n \text{ div } 2) xs; \]

\[ zs = \text{drop } (n \text{ div } 2) xs \]

\[ \text{in if } n \leq 1 \text{ then } 0 \]

\[ \text{else } C_{\text{msort}} ys + C_{\text{msort}} zs + C_{\text{merge}} (\text{msort } ys) (\text{msort } zs) \]

By computation inductions we obtain:

\[ |\text{merge } xs \ ys| = |xs| + |ys| \]  

(2.13)

\[ |\text{msort } xs| = |xs| \]  

(2.14)

\[ C_{\text{merge}} xs \ ys \leq |xs| + |ys| \]  

(2.15)

where the proof of (2.14) uses (2.13).

To simplify technicalities, we prove the \( n \cdot \log n \) bound on the number of comparisons in \( msort \) only for \( n = 2^k \), in which case the bound becomes \( k \cdot 2^k \).

Lemma 2.4. \(|xs| = 2^k \longrightarrow C_{\text{msort } xs} \leq k \cdot 2^k \)

Proof by induction on \( k \). The base case is trivial and we concentrate on the step. Let \( n = |xs|, ys = \text{take } (n \text{ div } 2) xs \) and \( zs = \text{drop } (n \text{ div } 2) xs \). The case \( n \leq 1 \) is trivial. Now assume \( n > 1 \).
2.5 Bottom-Up Merge Sort

Bottom-up merge sort starts by turning the input \([x_1, \ldots, x_n]\) into the list \([[[x_1], \ldots, [x_n]]]\). Then it passes over this list of lists repeatedly, merging pairs of adjacent lists on every pass until at most one list is left.

\[
C_{\text{msort}} \text{zs} = C_{\text{msort}} \text{ys} + C_{\text{msort}} \text{zs} + C_{\text{merge}} (\text{msort} \text{ys}) (\text{msort} \text{zs})
\]

\[
\leq C_{\text{msort}} \text{ys} + C_{\text{msort}} \text{zs} + |\text{ys}| + |\text{zs}|
\]

using (2.15) and (2.14) by IH

\[
= k \cdot 2^k + k \cdot 2^k + |\text{ys}| + |\text{zs}|
\]

\[
= (k + 1) \cdot 2^k + 1
\]

by assumption \(|\text{zs}| = 2^k + 1\)

### 2.5.1 Functional Correctness

We introduce the abbreviation \(\text{mset}_\text{mset} :: \text{'a list list} \Rightarrow \text{'a multiset}:

\[
\text{mset}_\text{mset} \text{xss} \equiv \sum_{\text{y}} (\text{image}_\text{mset} \text{mset} (\text{mset} \text{xss}))
\]

These are the key properties of the functions involved:

\[
\text{mset}_\text{mset} (\text{merge}_\text{adj}2 \text{acc xss}) = \text{mset}_\text{mset} \text{acc} + \text{mset}_\text{mset} \text{xss}
\]
\[ mset (\text{merge\_all} \, \text{ss}) = mset \, mset \, \text{ss} \quad (2.17) \]
\[ mset (\text{msort\_bu} \, \text{xs}) = mset \, \text{xs} \]
\[ (\forall \, \text{zs} \in \text{set} \, \text{zs}. \, \text{sorted} \, \text{zs}) \rightarrow (\forall \, \text{zs} \in \text{set} \, (\text{merge\_adj} \, \text{ss}). \, \text{sorted} \, \text{zs}) \]
\[ (\forall \, \text{zs} \in \text{set} \, \text{zs}. \, \text{sorted} \, \text{zs}) \rightarrow \text{sorted} \, (\text{merge\_all} \, \text{ss}) \quad (2.18) \]
\[ \text{sorted} \, (\text{msort\_bu} \, \text{xs}) \]

The third and the last proposition prove functional correctness of \text{msort\_bu}. The proof of each proposition may use the preceding proposition and the propositions (2.10) and (2.12). The propositions about \text{merge\_adj} and \text{merge\_all} are proved by computation inductions.

### 2.5.2 Running Time Analysis

Again, we count only comparisons:

\[ C_{\text{merge\_adj}} :: \forall \text{a list list} \Rightarrow \text{nat} \]
\[ C_{\text{merge\_adj}} [] = 0 \]
\[ C_{\text{merge\_adj}} [\_] = 0 \]
\[ C_{\text{merge\_adj}} (\text{xs} \# \text{ys} \# \text{zs}) = C_{\text{merge} \, \text{xs} \, \text{ys}} + C_{\text{merge\_adj} \, \text{zs}} \]

\[ C_{\text{merge\_all}} :: \forall \text{a list list} \Rightarrow \text{nat} \]
\[ C_{\text{merge\_all}} [] = 0 \]
\[ C_{\text{merge\_all}} [\text{xs}] = 0 \]
\[ C_{\text{merge\_all}} \, \text{ss} = C_{\text{merge\_adj} \, \text{ss}} + C_{\text{merge\_all}} (\text{merge\_adj} \, \text{ss}) \]

\[ C_{\text{msort\_bu}} :: \forall \text{a list} \Rightarrow \text{nat} \]
\[ C_{\text{msort\_bu}} \, \text{xs} = C_{\text{merge\_all}} (\text{map} (\lambda \, x. \, [x]) \, \text{xs}) \]

By simple computation inductions we obtain:

\[ \text{even} \, |\text{ss}| \land (\forall \, \text{zs} \in \text{set} \, \text{zs}. \, |\text{zs}| = m) \rightarrow (\forall \, \text{zs} \in \text{set} \, (\text{merge\_adj} \, \text{ss}). \, |\text{zs}| = 2 \cdot m) \]
\[ (\forall \, \text{zs} \in \text{set} \, \text{zs}. \, |\text{zs}| = m) \rightarrow C_{\text{merge\_adj} \, \text{ss}} \leq m \cdot |\text{ss}| \quad (2.19) \]
\[ C_{\text{merge\_all}} \, \text{ss} \leq m \cdot k \cdot 2^k \quad (2.20) \]

using (2.13) for (2.19) and (2.15) for (2.20).

**Lemma 2.5.** \( (\forall \, \text{zs} \in \text{set} \, \text{zs}. \, |\text{zs}| = m) \land |\text{ss}| = 2^k \rightarrow C_{\text{merge\_all}} \, \text{ss} \leq m \cdot k \cdot 2^k \)

**Proof** by induction on the computation of \text{merge\_all}. We concentrate on the nontrivial recursive case arising from the third equation. We assume \(|\text{ss}| > 1\),
∀xs ∈ set xss. |xs| = m and |xss| = 2^k. Clearly k ≥ 1 and thus even |xss|. Thus (2.19) implies ∀xs ∈ set (merge_adj xss). |xs| = 2 · m. Also note

|merge_adj xss|
= ([xss] + 1) div 2
= 2^k - 1

using |xss| = 2^k and k ≥ 1 by arithmetic

Let yss = merge_adj xss. We can now prove the lemma:

\[ C_{merge_all} xss = C_{merge_adj} xss + C_{merge_all} yss \]
\[ ≤ m · 2^k + C_{merge_all} yss \]
\[ ≤ m · 2^k + 2 · m · (k - 1) · 2^k - 1 \]
\[ = m · k · 2^k \]

by IH using ∀xs ∈ set yss. |xs| = 2 · m and |yss| = 2^k - 1

Setting m = 1 we obtain the same upper bound as for top-down merge sort in Lemma 2.4:

Corollary 2.6. |xs| = 2^k → C_{msort_bu} xs ≤ k · 2^k

2.6 Natural Merge Sort

A disadvantage of all the sorting functions we have seen so far (except insertion sort) is that even in the best case they do not improve upon the \( n \cdot \log n \) bound. For example, given the sorted input [1, 2, 3, 4, 5], msort_bu will, as a first step, create [[1], [2, 3], [4, 5]] and then merge this list of lists recursively.

A slight variation of bottom-up merge sort, sometimes referred to as “natural merge sort,” first partitions the input into its constituent ascending and descending subsequences (collectively referred to as runs) and only then starts merging. In the above example we would get merge_all [[1, 2, 3, 4, 5]], which returns immediately with the result [1, 2, 3, 4, 5]. Assuming that obtaining runs is of linear complexity, this yields a best-case performance that is linear in the number of list elements.

Function runs computes the initial list of lists; it is defined mutually recursively with asc and desc, which gather ascending and descending runs in accumulating parameters:

\[
\begin{align*}
\text{runs} &:: \ 'a \ list \Rightarrow \ 'a \ list \ list \\
\text{runs} \ (a \# b \# xs) &= (\text{if} \ \text{b} < \text{a} \ \text{then} \ \text{desc} \ b \ [a] \ xss \ \text{else} \ \text{asc} \ b \ ((\#) \ a) \ xss) \\
\text{runs} \ [x] &= [[x]] \\
\text{runs} \ [] &= [] \\
\text{asc} &:: \ 'a \Rightarrow \ ('a \ list \Rightarrow \ 'a \ list) \Rightarrow \ 'a \ list \Rightarrow \ 'a \ list \ list
\end{align*}
\]
\[
\text{asc } a \text{ as } (b \# bs)
= (\text{if } b < a \text{ then } \text{asc } b \text{ (as } o \text{ (}) a \text{ bs } \textbf{else} \text{ as } [a] \# \text{ runs } (b \# bs))
\text{asc } a \text{ as } [] = [\text{as } [a]]
\]

\[
\text{desc :: 'a } \Rightarrow \text{ 'a list } \Rightarrow \text{ 'a list } \Rightarrow \text{ 'a list list}
\]
\[
\text{desc } a \text{ as } (b \# bs)
= (\text{if } b < a \text{ then } \text{desc } b \text{ (a } \# \text{ as) bs } \textbf{else} \text{ (a } \# \text{ as) } \# \text{ runs } (b \# bs))
\text{desc } a \text{ as } [] = [\text{a } \# \text{ as]}
\]

Function \text{desc} needs to reverse the descending run it collects. Therefore a natural choice for the type of its accumulator \text{as} is \text{list}, since recursively prepending elements (using \((#)) ultimately yields a reversed list.

Function \text{asc} collects an ascending run and is slightly more complicated than \text{desc}. If we used lists, we could accumulate the elements similarly to \text{desc} but using \text{as } @ [a] instead of \text{a } \# \text{ as}. This would take quadratic time in the number of appended elements. Therefore the “standard” solution is to accumulate elements using \((#)) and to reverse the accumulator in linear time (as shown in Section 1.5.1) at the end. However, another interesting option (that yields better performance for some functional languages, like Haskell) is to use difference lists. This is the option we chose for \text{asc}.

In the functional programming world, difference lists are a well-known trick to append lists in constant time by representing lists as functions of type \text{'}a list } \Rightarrow \text{ 'a list}. For difference lists, we have the following correspondences: empty list [] \approx \lambda x. x, singleton list [x] \approx (\#) x, and list append \text{xs } @ \text{ys } \approx \text{xs } \circ \text{ys} (\text{taking constant time}). Moreover, transforming a difference list \text{xs} into a normal list is as easy as \text{xs } [] (\text{taking linear time}).

Note that, due to the mutually recursive definitions of \text{runs}, \text{asc}, and \text{desc}, whenever we prove a property of \text{runs}, we simultaneously have to prove suitable properties of \text{asc} and \text{desc} using mutual induction.

Natural merge sort is the composition of \text{merge\_all} and \text{runs}:

\[
\text{nmsort :: 'a list } \Rightarrow \text{ 'a list}
\]
\[
\text{nmsort } xs = \text{merge\_all } (\text{runs } xs)
\]

### 2.6.1 Functional Correctness

We have
2.6 Natural Merge Sort

\begin{align*}
& (\forall xs\; ys.\; f\; (xs \odot ys) = f\; xs \odot ys) \rightarrow \\
& mset\_mset\; (\text{asc}\; x\; f\; ys) = \{ x \} + mset\; (f\; []) + mset\; ys \tag{2.21} \\
& mset\_mset\; (\text{desc}\; x\; xs\; ys) = \{ x \} + mset\; xs + mset\; ys \tag{2.22} \\
& mset\_mset\; (\text{runs}\; xs) = mset\; xs \tag{2.23} \\
& mset\; (nmsort\; xs) = mset\; xs \tag{2.24}
\end{align*}

where (2.23), (2.21), and (2.22) are proved simultaneously. The assumption of (2.21) on \( f \) ensures that \( f \) is a difference list. We use (2.23) together with (2.17) in order to show (2.24). Moreover, we have

\begin{align*}
& \forall x \in \text{set}\; (\text{runs}\; xs).\; \text{sorted}\; x \tag{2.25} \\
& \text{sorted}\; (\text{nmsort}\; xs) \tag{2.26}
\end{align*}

where we use (2.25) together with (2.18) to obtain (2.26).

2.6.2 Running Time Analysis

Once more, we only count comparisons:

\begin{align*}
&C_{\text{runs}} :: \forall a\; \text{list} \Rightarrow \text{nat} \\
&C_{\text{runs}} (a \# b \# xs) = 1 + \left( \begin{array}{l}
\text{if } b < a \text{ then } C_{\text{desc}} b\; xs \text{ else } C_{\text{asc}} b\; xs \\
\end{array} \right) \\
&C_{\text{runs}} [] = 0 \\
&C_{\text{runs}} [\_] = 0 \\

&C_{\text{asc}} :: \forall a \Rightarrow \forall \text{list} \Rightarrow \text{nat} \\
&C_{\text{asc}} a (b \# bs) = 1 + \left( \begin{array}{l}
\text{if } b < a \text{ then } C_{\text{asc}} b\; bs \text{ else } C_{\text{runs}} (b \# bs) \\
\end{array} \right) \\
&C_{\text{asc}} [\_] = 0 \\

&C_{\text{desc}} :: \forall a \Rightarrow \forall \text{list} \Rightarrow \text{nat} \\
&C_{\text{desc}} a (b \# bs) = 1 + \left( \begin{array}{l}
\text{if } b < a \text{ then } C_{\text{desc}} b\; bs \text{ else } C_{\text{runs}} (b \# bs) \\
\end{array} \right) \\
&C_{\text{desc}} [\_] = 0 \\

&C_{\text{nmsort}} :: \forall \text{list} \Rightarrow \text{nat} \\
&C_{\text{nmsort}} xs = C_{\text{runs}} xs + C_{\text{merge\_all}} (\text{runs}\; xs)
\end{align*}

Again note the mutually recursive definitions of \( C_{\text{runs}} \), \( C_{\text{asc}} \), and \( C_{\text{desc}} \). Hence the remark on proofs about \text{runs} also applies to proofs about \( C_{\text{runs}} \).

Before talking about \( C_{\text{nmsort}} \), we need a variant of Lemma 2.5 that also works for lists whose lengths are not powers of two (since the result of \text{runs} will usually not satisfy this property).
To this end, we will need the following two results, which we prove by two simple computation inductions using (2.15) and (2.13):

\[ C_{\text{merge}_{\text{adj}}} xss \leq |\text{concat } xss| \quad (2.27) \]
\[ |\text{concat} (\text{merge}_{\text{adj}} xss)| = |\text{concat } xss| \quad (2.28) \]

**Lemma 2.7.** \( C_{\text{merge}_{\text{all}}} xss \leq |\text{concat } xss| \cdot [\lg |xss|] \)

**Proof.** by induction on the computation of \( C_{\text{merge}_{\text{all}}} \). We concentrate on the nontrivial recursive case arising from the third equation. It follows that \( xss \) is of the form \( xs \# ys \# zss \).

Further note that for all \( n :: \text{nat} \):

\[ 2 \leq n \rightarrow [\lg n] = [\lg ((n - 1) \div 2 + 1)] + 1 \quad (2.29) \]

Now, let \( m = |\text{concat } xss| \). Then we have

\[
\begin{align*}
C_{\text{merge}_{\text{all}}} xss &= C_{\text{merge}_{\text{adj}}} xss + C_{\text{merge}_{\text{all}}} (\text{merge}_{\text{adj}} xss) \\
&\leq m + C_{\text{merge}_{\text{all}}} (\text{merge}_{\text{adj}} xss) \quad \text{using (2.27)} \\
&\leq m + |\text{concat} (\text{merge}_{\text{adj}} xss)| \cdot [\lg |\text{merge}_{\text{adj}} xss|] \quad \text{by IH} \\
&= m + m \cdot [\lg |\text{merge}_{\text{adj}} xss|] \\
&= m + m \cdot [\lg ((|xss| + 1) \div 2)] \quad \text{by (2.16)} \\
&= m + m \cdot [\lg ((|xss| + 1) \div 2 + 1)] \\
&= m \cdot (\lfloor \lg (\lfloor |xss| + 1 \rfloor \div 2 + 2) \rfloor + 1) \quad \text{(2.29)} \\
&= m \cdot [\lg (\lfloor |xss| + 2 \rfloor)] \quad \square
\end{align*}
\]

Three simple computation inductions, each performed simultaneously for the corresponding mutually recursive definitions, yield:

\[
(\forall xs ys. f (xs @ ys) = f xs @ ys) \rightarrow \\
|\text{concat} (\text{asc } a f ys)| = 1 + |f []| + |ys|, \\
|\text{concat} (\text{desc } a xs ys)| = 1 + |xs| + |ys|, \\
|\text{concat} (\text{runs } xs)| = |xs| \quad (2.30)
\]

\[
(\forall xs ys. f (xs @ ys) = f xs @ ys) \rightarrow |\text{asc } a f ys| \leq 1 + |ys|, \\
|\text{desc } a xs ys| \leq 1 + |ys|, |\text{runs } xs| \leq |xs| \quad (2.31)
\]

\[
C_{\text{asc } a } x ys \leq |ys|, C_{\text{desc } a } ys \leq |ys|, C_{\text{runs } xs} \leq |xs| - 1 \quad (2.32)
\]

At this point we obtain an upper bound on the number of comparisons required by \( C_{\text{nmsort}} \).

**Lemma 2.8.** \( |xs| = n \rightarrow C_{\text{nmsort}} x xs \leq n + n \cdot [\lg n] \)

**Proof.** Note that
2.7 Uniqueness of Sorting

We have seen many different sorting functions now and it may come as a surprise that they are all the same in the sense that they are all extensionally equal: they have the same input/output behaviour (but of course not the same running time).

A more abstract formulation of this is that the result of sorting a list is uniquely determined by the specification of sorting. This is what we call the uniqueness of sorting: Consider lists whose elements are sorted w.r.t. some linear order. Then any two such lists with the same multiset of elements are equal. Formally:

**Theorem 2.9** (Uniqueness of sorting).

\[ \text{mset} \; \text{xs} = \text{mset} \; \text{ys} \land \text{sorted} \; \text{xs} \land \text{sorted} \; \text{ys} \rightarrow \text{xs} \; = \; \text{ys} \]

*Proof* by induction on \( \text{xs} \) (for arbitrary \( \text{ys} \)). The base case is trivial. In the induction step, \( \text{xs} = x \# \; \text{xs}' \). Thus \( \text{ys} \) must also be of the form \( y \# \; \text{ys}' \) (otherwise their multisets could not be equal).

Thus we now have to prove \( x \# \; \text{xs}' = y \# \; \text{ys}' \), and the facts that we have available to do this are

\[ \text{mset} \; (x \# \; \text{xs}') = \text{mset} \; (y \# \; \text{ys}') \quad (2.33) \]
\[ \text{sorted} \; (x \# \; \text{xs}') \land \text{sorted} \; (y \# \; \text{ys}') \quad (2.34) \]

and the induction hypothesis

\[ \forall \; \text{ys}'. \; \text{mset} \; \text{xs}' = \text{mset} \; \text{ys}' \land \text{sorted} \; \text{xs}' \land \text{sorted} \; \text{ys}' \rightarrow \text{xs}' \; = \; \text{ys}' \]  \hfill (IH)

Our first objective now is to show that \( x = y \). Either \( x \leq y \) or \( x \geq y \) must hold. Let us first prove \( x = y \) for the case \( x \leq y \). From (2.33), we have \( x \in_{a} \text{mset} \; (x \# \; \text{xs}') = \text{mset} \; (y \# \; \text{ys}') \). Thus \( x \) is contained somewhere in the list \( y \# \; \text{ys}' \). Since \( y \# \; \text{ys}' \) is sorted, all elements of \( y \# \; \text{ys}' \) are \( \geq y \); in particular we then have \( x \geq y \). Together with \( x \leq y \), we obtain \( x = y \) as desired. The case \( x \geq y \) is completely analogous.
26 Chapter 2 Sorting

Now that we know that \(x = y\), the rest of the proof is immediate: From (2.33) we obtain \(mset\,xs' = mset\,ys'\), and with that and (2.34), the (IH) tells us that \(xs' = ys'\) and we are done. \qed

This theorem directly implies the extensional equality of all sorting functions that we alluded to earlier. That is, any two functions that satisfy the specification from Section 2.1 are extensionally equal.

**Corollary 2.10** (All sorting functions are extensionally equal). If \(f\) and \(g\) are functions of type \((\alpha :: \text{linorder}) \Rightarrow \alpha \text{ list}\) such that

\[
\forall\,zs.\,\text{sorted}\,(f\,zs) \land mset\,(f\,zs) = mset\,zs
\]
\[
\forall\,zs.\,\text{sorted}\,(g\,zs) \land mset\,(g\,zs) = mset\,zs
\]

then \(\forall\,zs.\,f\,zs = g\,zs\); or, equivalently: \(f = g\)

**Proof.** We use Theorem 2.9 with the instantiations \(xs = f\,zs\) and \(ys = g\,zs\). \qed

Note that for both of these theorems, the \text{linorder} constraint on the element type is crucial: if we have an order \(\preceq\) that is not linear, then there are elements \(x, y\) with \(x \preceq y\) and \(y \preceq x\) but \(x \neq y\). Consequently, the lists \([x, y]\) and \([y, x]\) are not equal, even though they are both sorted w.r.t. \(\preceq\) and contain the same elements.

### 2.8 Stability

A sorting function is called stable if the order of equal elements is preserved. However, this only makes a difference if elements are not identified with their keys, as we have done so far. Let us assume instead that sorting is parameterized with a key function \(f :: \alpha \Rightarrow \text{'k}\) that maps an element to its key and that the keys \(\text{'k}\) are linearly ordered, not the elements. This is the specification of a sorting function \(\text{sort\_key}\):

\[
\text{mset}\,(\text{sort\_key}\,f\,xs) = mset\,xs
\]
\[
\text{sorted}\,(\text{map}\,f\,(\text{sort\_key}\,f\,xs))
\]

Assuming (for simplicity) we are sorting pairs of keys and some attached information, stability means that sorting \([(2, x), (1, z), (1, y)]\) yields \([(1, z), (1, y), (2, x)]\) and not \([(1, y), (1, z), (2, x)]\). That is, if we extract all elements with the same key after sorting \(xs\), they should be in the same order as in \(xs\):

\[
\text{filter}\,(\lambda y.\,f\,y = k)\,(\text{sort\_key}\,f\,xs) = \text{filter}\,(\lambda y.\,f\,y = k)\,xs
\]

We will now define insertion sort adapted to keys and verify its correctness and stability.
The proofs of the functional correctness properties

\[
\text{mset } (\text{insort\_key } f \ x s) = \text{mset } x s
\]

\[
\text{sorted } (\text{map } f \ (\text{insort\_key } f \ x s))
\]  \hspace{2cm} (2.35)

are completely analogous to their counterparts for plain insort.

The proof of stability uses three auxiliary properties:

\[
(\forall x \in \text{set } x s. \ f \ a \leq f x) \rightarrow \text{insort\_key } f \ a \ x s = a \# x s \]  \hspace{2cm} (2.36)

\[
\neg P x \rightarrow \text{filter } P \ (\text{insort\_key } f \ x s) = \text{filter } P \ x s
\]  \hspace{2cm} (2.37)

\[
\text{sorted } (\text{map } f \ x s) \land P x \rightarrow
\text{filter } P \ (\text{insort\_key } f \ x s) = \text{insort\_key } f \ x \ (\text{filter } P \ x s)
\]  \hspace{2cm} (2.38)

The first one is proved by a case analysis on \( x s \). The other two are proved by induction on \( x s \), using (2.36) in the proof of (2.38).

Lemma 2.11 (Stability of insort\_key).

\[\text{filter } (\lambda y. \ f \ y = k) \ (\text{insort\_key } f \ x s) = \text{filter } (\lambda y. \ f \ y = k) \ x s\]

**Proof**  by induction on \( x s \). The base case is trivial. In the induction step we consider the list \( a \# x s \) and perform a case analysis. If \( f \ a \neq k \) the claim follows by IH using (2.37). Now assume \( f \ a = k \):

\[
\text{filter } (\lambda y. \ f \ y = k) \ (\text{insort\_key } f \ (a \# x s))
\]  \hspace{2cm} using \( f \ a = k \), (2.38), (2.35)

\[
= \text{filter } (\lambda y. \ f \ y = k) \ (\text{insort\_key } f \ a \ (\text{insort\_key } f \ x s))
\]  \hspace{2cm} \text{by IH}

\[
= \text{insort\_key } f \ a \ (\text{filter } (\lambda y. \ f \ y = k) \ (\text{insort\_key } f \ x s))
\]  \hspace{2cm} \text{using } f \ a = k \text{ and } (2.36)

\[
= \text{filter } (\lambda y. \ f \ y = k) \ (a \# x s)
\]  \hspace{2cm} \text{using } f \ a = k

As exercises we recommend to adapt some of the other sorting algorithms above to sorting with keys and to prove their correctness and stability.
2.9 Exercises

Exercise 2.1. Show that $T_{\text{insort}}$ achieves its optimal value of $2 \cdot n + 1$ for sorted lists, and its worst-case value of $(n + 1) \cdot (n + 2) \div 2$ for the list rev [0..<n].

Exercise 2.2. Function $\text{quicksort}$ appends the lists returned from the recursive calls. This is expensive because the running time of (@) is linear in the length of its first argument. Define a function $\text{quicksort2} :: 'a list \Rightarrow 'a list \Rightarrow 'a list$ that avoids (@) but accumulates the result in its second parameter via (#) only. Prove $\text{quicksort2} \; xs \; ys = \text{quicksort} \; xs @ ys$.

Exercise 2.3. There is one obvious optimisation to the version of quicksort that we studied before: instead of partitioning the list into those elements that are smaller than the pivot and those that are at least as big as the pivot, we can use three-way partitioning:

\[
\begin{align*}
\text{partition3} :: 'a \Rightarrow 'a list \Rightarrow 'a list \times 'a list \times 'a list \\
\text{partition3} \; x \; xs &= (\text{filter} \; (\lambda y. \; y < x) \; xs, \; \text{filter} \; (\lambda y. \; y = x) \; xs, \; \text{filter} \; (\lambda y. \; y > x) \; xs) \\
\text{quicksort3} :: 'a list \Rightarrow 'a list \\
\text{quicksort3} \; [] &= [] \\
\text{quicksort3} \; (x \# xs) &= (\text{let} \; (ls, \; es, \; gs) = \text{partition3} \; x \; xs \\
& \quad \text{in} \; \text{quicksort3} \; ls @ x \# es @ \text{quicksort3} \; gs)
\end{align*}
\]

Prove that this version of quicksort also produces the correct results.

Exercise 2.4. In this exercise, we will examine the worst-case behaviour of Quicksort, which is e.g. achieved if the input list is already sorted. Consider the time function for Quicksort:

\[
\begin{align*}
T_{\text{quicksort}} :: 'a list \Rightarrow \text{nat} \\
T_{\text{quicksort}} \; [] &= 1 \\
T_{\text{quicksort}} \; (x \# xs) &= T_{\text{quicksort}} \; (\text{filter} \; (\lambda y. \; y < x) \; xs) + \\
& \quad T_{\text{quicksort}} \; (\text{filter} \; (\lambda y. \; y \geq x) \; xs) + \\
& \quad 2 \cdot T_{\text{filter}} \; (\lambda x. \; 1) \; xs + 1
\end{align*}
\]

1. Show that Quicksort takes quadratic time on sorted lists by proving
2.9 Exercises

\[ \text{sorted } xs \rightarrow T_{\text{quicksort}} xs = a \cdot |xs|^2 + b \cdot |xs| + c \]
for suitable values \(a, b, c\).

2. Show that this is the worst-case running time by proving
\[ T_{\text{quicksort}} xs \leq a \cdot |xs|^2 + b \cdot |xs| + c \]
for the values of \(a, b, c\) you determined in the previous step.

Exercise 2.5. The definition of \(m\text{sort}\) is inefficient in that it calls \(\text{length}\), \(\text{take}\) and \(\text{drop}\) for each list. Instead we can split the list into two halves by traversing it only once and putting its elements alternately on two piles, for example \(\text{halve } [2, 3, 4] \rightarrow ([0], [1]) = ([4, 2, 0], [3, 1])\). Define \(\text{halve}\) and \(m\text{sort}_2\)

\[
\begin{align*}
m\text{sort}_2 [] &= [] \\
m\text{sort}_2 [x] &= [x] \\
m\text{sort}_2 xs &= (\text{let} \ (ys_1, ys_2) = \text{halve } xs ([], []) \text{ in } \text{merge } (m\text{sort}_2 ys_1) (m\text{sort}_2 ys_2))
\end{align*}
\]
and prove \(\text{mset } (m\text{sort}_2 xs) = \text{mset } xs\) and \(\text{sorted } (m\text{sort}_2 xs)\). (Hint for Isabelle users: The definition of \(m\text{sort}_2\) is tricky because its termination relies on suitable properties of \(\text{halve}\).)

Exercise 2.6. Define a tail-recursive variant

\[
\text{merge}_\text{adj2} :: \text{'a list list} \Rightarrow \text{'a list list} \Rightarrow \text{'a list list}
\]
of \(\text{merge}_\text{adj}\) and define new variants \(\text{merge}_\text{all2}\) and \(\text{msort}_\text{bu2}\) of \(\text{merge}_\text{all}\) and \(\text{msort}_\text{bu}\) that utilize \(\text{merge}_\text{adj2}\). Prove functional correctness:

\[
\text{mset } (\text{msort}_\text{bu2 } xs) = \text{mset } xs \quad \text{sorted } (\text{msort}_\text{bu2 } xs)
\]
Note that \(\text{merge}_\text{adj2 } []\ xss = \text{merge}_\text{adj } xss\) is not required.
A topic that is somewhat related to that of sorting is selection: given a list $xs$ of length $n$ with some linear order defined on its elements and a natural number $k < n$, return the $k$-th smallest number in the list (starting with $k = 0$ for the minimal element). If $xs$ is sorted, this is exactly the $k$-th element of the list.

The defining properties of the selection operation are as follows:

$$k < |xs| \implies |\{ y \in_x mset xs \mid y < \text{select} \ k \ xs \}| \leq k$$
$$\land |\{ y \in_x mset xs \mid y > \text{select} \ k \ xs \}| < |xs| - k$$

(3.1)

In words: $\text{select} \ k \ xs$ has the property that at most $k$ elements in the list are strictly smaller than it and at most $n - k$ are strictly bigger.

These properties fully specify the selection operation, as shown by the following theorem:

**Theorem 3.1 (Uniqueness of the selection operation).**

If $k < |xs|$ and

$$|\{ z \in_x mset xs \mid z < x \}| \leq k$$
$$|\{ z \in_x mset xs \mid z > y \}| \leq k$$

then $x = y$.

**Proof.** Suppose $x \neq y$ and then w.l.o.g. $x < y$. This implies:

$$\{ z \in_x mset xs \mid z < x \} \subseteq \{ z \in_x mset xs \mid z < y \}$$

(3.3)

From this we can prove the contradiction $|xs| < |xs|$: 

$$|xs| = |\{ z \in_x mset xs \mid z \leq x \}| + |\{ z \in_x mset xs \mid z > x \}|$$
$$\leq |\{ z \in_x mset xs \mid z < y \}| + |\{ z \in_x mset xs \mid z > x \}|$$
$$< k + (|xs| - k)$$
$$= |xs|$$

using (3.2), (3.3) □
An equivalent, more concrete definition is the following:

\[
\text{select} :: \text{nat} \Rightarrow \text{a list} \Rightarrow \text{a} \\
\text{select} k \ xs = \text{sort} \ xs \ ! k
\] (3.4)

**Theorem 3.2.** *select* as defined by Equation (3.4) satisfies the conditions (3.1).

*Proof.* If \(ys\) is sorted, a straightforward induction on \(ys\) shows the following:

\[
\begin{align*}
\{ x \in_s \text{mset} \ ys \mid x < ys ! k \} & \subseteq \text{mset} \ (\text{take} \ k \ ys) \\
\{ x \in_s \text{mset} \ ys \mid x > ys ! k \} & \subseteq \text{mset} \ (\text{drop} \ (k + 1) \ ys)
\end{align*}
\]

Taking the size of the multisets on both sides, we obtain:

\[
\begin{align*}
\| \{ x \in_s \text{mset} \ ys \mid x < ys ! k \} \| & \leq k \\
\| \{ x \in_s \text{mset} \ ys \mid x > ys ! k \} \| & < |ys| - k
\end{align*}
\]

Now, for an arbitrary list \(xs\), we instantiate the above with \(ys := \text{sort} \ xs\) and obtain:

\[
\begin{align*}
k & \geq \| \{ x \in_s \text{mset} \ (\text{sort} \ xs) \mid x < \text{sort} \ xs ! k \} \|
= \| \{ x \in_s \text{mset} \ xs \mid x < \text{sort} \ xs ! k \} \|
\quad \text{using mset (sort xs) = mset xs}
= \| \{ x \in_s \text{mset} \ xs \mid x < \text{select} k \ xs \} \|
\quad \text{using (3.4)}
\end{align*}
\]

and analogously for the elements greater than \(\text{select} k \ xs\). \(\square\)

We will frequently need another important fact about \(\text{sort}\) and \(\text{select}\), namely that they are invariant under permutation of the input list:

**Lemma 3.3.** Let \(xs\) and \(ys\) be lists with \(\text{mset} \ xs = \text{mset} \ ys\). Then:

\[
\begin{align*}
\text{sort} \ xs & = \text{sort} \ ys \quad (3.5) \\
\text{select} k \ xs & = \text{select} k \ ys \quad (3.6)
\end{align*}
\]

*Proof.* Equation (3.5) follows directly from Theorem 2.9 (the uniqueness of the \(\text{sort}\) operation), and (3.6) then follows from (3.5) and our definition of \(\text{select}\). \(\square\)

The definition using \(\text{sort} \ xs \ ! k\) already gives us a straightforward \(O(n \lg n)\) algorithm for the selection operation: sort the list with one of our \(O(n \lg n)\) sorting algorithms and then return the \(k\)-th element of the resulting sorted list. It is also fairly easy to come up with an algorithm that has running time \(O(kn)\), i.e. that runs in linear time in \(n\) for any fixed \(k\) (see Exercise 3.3).

In the remainder of this chapter, we will look at a selection algorithm that achieves \(O(n)\) running time *uniformly for all* \(k < n\) [Blum et al. 1973]. Since a selection algorithm must inspect every element at least once (see Exercise 3.4), this running time is asymptotically optimal.
3.0.1 Exercises

Exercise 3.1. A simple special case of selection is \( \text{select} \ 0 \ \text{xs} \), i.e. the minimum. Implement a linear-time function \( \text{select0} \) such that

\[
x \neq [] \rightarrow \text{select0} \ xs = \text{select} \ 0 \ xs
\]

and prove this. This function should be tail-recursive and traverse the list exactly once. You need not prove the linear running time (it should be obvious).

Exercise 3.2. How can your \( \text{select0} \) algorithm be modified to obtain an analogous algorithm \( \text{select1} \) such that

\[
|xs| > 1 \rightarrow \text{select1} \ xs = \text{select} \ 1 \ xs
\]

Do not try to prove the correctness yet; it gets somewhat tedious and you will be able to prove it more easily after the next exercise.

Exercise 3.3.

1. Based on the previous two exercises, implement and prove correct an algorithm \( \text{select\textunderscore fixed} \) that fulfills

\[
k < |xs| \rightarrow \text{select\textunderscore fixed} \ k \ xs = \text{select} \ k \ xs
\]

The algorithm must be tail-recursive with running time \( O(kn) \) and traverse the list exactly once.

Hint: one approach is to first define a function \( \text{take\textunderscore sort} \) that computes \( \text{take} \ m \ (\text{sort} \ xs) \) in time \( O(mn) \).

2. Prove your \( \text{select1} \) from the previous exercise correct by showing that it is equivalent to \( \text{select\textunderscore fixed} \ 1 \).

3. Define a suitable time function for your \( \text{select\textunderscore fixed} \). Prove that this time function is \( O(kn) \), i.e. that

\[
T_{\text{select\textunderscore fixed}} \ k \ xs \leq C_1 \cdot k \cdot |xs| + C_2 \cdot |xs| + C_3 \cdot k + C_4
\]

for all \( k < |xs| \) for some constants \( C_1 \) to \( C_4 \).

If you have trouble finding the concrete values for these constants, try proving the result with symbolic constants first and observe what conditions need to be fulfilled in order to make the induction step go through.

Exercise 3.4. Show that if \( xs \) is a list of integers with no repeated elements, an algorithm computing the result of \( \text{select} \ k \ xs \) must examine every single element, i.e. for any index \( i < |xs| \), the \( i \)-th element can be replaced by some other number such that the result changes. Formally:

\[
k < |xs| \land i < |xs| \land \text{distinct} \ xs \rightarrow \\
(\exists \ z. \ \text{select} \ k \ (xs[i := z]) \neq \text{select} \ k \ xs)
\]
Here, the notation $\text{xs}[i := z]$ denotes the list $\text{xs}$ where the $i$-th element has been replaced with $z$ (the first list element, as always, having index 0).

Hint: a lemma you might find useful is that $\lambda k. \text{select } k \text{ xs}$ is injective if $\text{xs}$ has no repeated elements.

### 3.1 A Divide-and-Conquer Approach

As a first step in our attempt to derive an efficient algorithm for selection, recall what we did with the function $\text{partition3}$ in the threeway quicksort algorithm in Exercise 2.3: we picked some pivot value $x$ from $\text{xs}$ and partitioned the input list $\text{xs}$ into the sublists $\text{ls}$, $\text{es}$, and $\text{gs}$ of the elements smaller, equal, and greater than $x$, respectively.

If we do the same for $\text{select } k \text{ xs}$, there are three possible cases:

- If $k < |\text{ls}|$, the element we are looking for is located in $\text{ls}$. To be more precise, it is the $k$-th smallest element of $\text{ls}$, i.e. $\text{select } k \text{ ls}$.
- If $k < |\text{ls}| + |\text{es}|$, the element we are looking for is located in $\text{es}$ and must therefore be equal to $x$.
- Otherwise, the element we are looking for must be located in $\text{gs}$. More precisely, it is the $k' = k - |\text{ls}| - |\text{es}|$-th smallest element of $\text{gs}$.

This gives us a straightforward recursive divide-and-conquer algorithm for selection.

To prove this formally, we first prove the following lemma about the behaviour of $\text{select}$ applied to a list of the form $\text{xs} @ \text{ys}$:

**Lemma 3.4.**

$$k < |\text{xs}| + |\text{ys}| \rightarrow (\forall x \in \text{set } \text{xs}. \forall y \in \text{set } \text{ys}. x \leq y) \rightarrow$$

$$\text{select } k \ (\text{xs} @ \text{ys}) = \begin{cases} \text{select } k \text{ xs} & \text{if } k < |\text{xs}| \\ \text{select } (k - |\text{ys}|) \text{ ys} & \text{otherwise} \end{cases}$$ (3.7)

**Proof.** The assumptions imply that $\text{sort } \text{xs} @ \text{sort } \text{ys}$ is sorted, so that due to the uniqueness of the $\text{sort}$ operation, we have:

$$\text{sort } (\text{xs} @ \text{ys}) = \text{sort } \text{xs} @ \text{sort } \text{ys}$$ (3.8)

Then:

$$\text{select } k \ (\text{xs} @ \text{ys})$$

$$= \text{sort } (\text{xs} @ \text{ys})! k \quad \text{using (3.4)}$$

$$= (\text{sort } \text{xs} @ \text{sort } \text{ys})! k \quad \text{using (3.8)}$$

$$= \begin{cases} \text{sort } \text{xs}! k & \text{if } k < |\text{xs}| \\ \text{sort } \text{ys}! (k - |\text{xs}|) & \text{otherwise} \end{cases}$$

$$= \begin{cases} \text{select } k \text{ xs} & \text{if } k < |\text{xs}| \\ \text{select } (k - |\text{ys}|) \text{ ys} & \text{otherwise} \end{cases} \quad \text{using (3.4)}$$

Now the recurrence outlined before is a direct consequence:
Theorem 3.5 (A recurrence for select). Let $k < |xs|$ and $x$ arbitrary. Then:

\[
\text{select } k \ text{ } xs = \begin{cases} 
\text{let } (ls, es, gs) = \text{partition } 3 \ x \ xs \\
\text{in } \begin{cases} 
\text{if } k < |ls| \ then \ \text{select } k \ ls \\
\text{else if } k < |ls| + |es| \ then \ x \\
\text{else } \ \text{select } (k - |ls| - |es|) \ gs
\end{cases}
\end{cases}
\]

Proof. We have $mset \ xs = mset \ ls + mset \ es + mset \ gs$ and $|xs| = |ls| + |es| + |gs|$. Then:

\[
\text{select } k \ text{ } xs = \begin{cases} 
\text{select } k \ (ls \ @ \ es \ @ \ gs) \\
\text{if } k < |ls| \ then \ \text{select } k \ ls \\
\text{else if } k - |ls| < |es| \ then \ \text{select } (k - |ls|) \ es \\
\text{else } \ \text{select } (k - |ls| - |es|) \ gs
\end{cases}
\]

Clearly, $k - |ls| < |es| \iff k < |ls| + |es|$ and $\text{select } (k - |ls|) \ es = x$ since $\text{select } (k - |ls|) es \in \text{set } es$ and $\text{set } es = \{x\}$ by definition. \qed

Note that this holds for any pivot $x$. Indeed, $x$ need not even be in the list itself. Therefore, the algorithm (which is also known as Quicksort [Hoare 1961] due to its similarities with Quicksort) is partially correct no matter what pivot we choose.

However, like with Quicksort, the number of recursive calls (and thereby the running time) depends strongly on the pivot choice:

- If we always choose a pivot that is smaller than any element in the list or bigger than any element in the list, the algorithm does not terminate at all.

- If we choose the smallest element in the list as a pivot every time, only one element is removed from the list in every recursion step so that we get $n$ recursive calls in total. Since we do a linear amount of work in every step, this leads to a running time of $\Theta(n^2)$.

- If we choose pivots from the list at random, the worst-case running time is again $\Theta(n^2)$, but the expected running time can be shown to be $\Theta(n)$, similarly to the situation in Quickselect. Indeed, it can also be shown that it is very unlikely that the running time is “significantly worse than linear” [Karp 1994, Section 2.5].

- If we choose a pivot that cuts the list in half every time (i.e. at most $\frac{n}{2}$ elements are strictly smaller than the pivot and at most $\frac{n}{2}$ are strictly bigger), we get a recursion depth of at most $\lceil \lg n \rceil$ and, by the Master Theorem, a running time of $\Theta(n)$ (assuming we can find such a pivot in linear time).

Clearly, the last case is the most desirable one. An element that cuts the list in half is called a median (a concept widely used in statistics).
For lists of odd length, there is a unique element in that list that achieves this, whereas for lists of even length there are two such elements (e.g. for the list [1,2,3,4], both 2 and 3 work). In general, a median need also not necessarily be an element of the list itself.

For our purposes, it is useful to pick one of the list elements as a canonical median and refer to it as the median of that list. If the list has even length, we use the smaller of the two medians. This leads us to the following formal definition:

\[
\text{median} :: 'a\ \text{list} \Rightarrow 'a
\]
\[
\text{median } xs = \text{select } (\lfloor|xs| - 1\rfloor + 2) xs
\]

Unfortunately, computing the median of a list is no easier than selection (see Exercise 3.5), so it seems that, for now, this does not really help us.

Exercise 3.5. Show that computing \(\text{select } k xs\) can be reduced in linear time to computing the median of a list, i.e. give a linear-time function \(\text{reduce}\_\text{select}\_\text{median}\) that satisfies

\[
xs \neq [] \land k < |xs| \rightarrow \\
\text{reduce}\_\text{select}\_\text{median } k xs \neq [] \land \\
\text{median } (\text{reduce}\_\text{select}\_\text{median } k xs) = \text{select } k xs
\]

and prove it.

3.2 The Median of Medians

We have seen that computing a true median in every recursive step is just as hard as the general selection problem, so using the median as a pivot is not going to work. The natural question now is: is there something that is almost as good as a median but easier to compute?

This is indeed the case, and this is where the ingenuity of the algorithm lies: instead of computing the median of all the list elements, compute the median of only a small fraction of list elements. To be precise, we do the following:

- chop the list into groups of 5 elements each (possibly with one smaller group at the end if \(n\) is not a multiple of 5)
- compute the median of each of the \(\lceil \frac{n}{5} \rceil\) groups (which can be done in constant time for each group using e.g. insertion sort, since their sizes are bounded by 5)
- compute the median \(M\) of these \(\lceil \frac{n}{5} \rceil\) elements (which can be done by a recursive call to the selection algorithm)
3.2 The Median of Medians

We call $M$ the **median of medians**. $M$ is not quite as good a pivot as the true median, but it is still fairly decent:

**Theorem 3.6 (Pivoting bounds for the median of medians).**

Let $xs$ be a list and let $< \text{ or } >$. Let

$$M \coloneqq \text{median}\left(\text{map}\ \text{median}\ (\text{chop}\ 5\ xs)\right)$$

where the chop function cuts a list into groups of a given size as described earlier:

$$\text{chop} :: \text{nat} \implies \text{a list} \implies \text{a list list}$$

$$\text{chop}\ 0\ _ = []$$

$$\text{chop}\ \_\ [] = []$$

$$\text{chop}\ s\ xs = \text{take}\ s\ xs \#\ \text{chop}\ s\ (\text{drop}\ s\ xs)$$

Then: $\| y \in_s\ mset\ xs \mid y < M \| \leq [0.7 \cdot n + 3]$

**Proof.** The result of chop $5\ xs$ is a list of $\lfloor n / 5 \rfloor$ chunks, each of size at most 5, i.e. $|\text{chop}\ 5\ xs| = \lfloor n / 5 \rfloor$. Let us split these chunks into two groups according to whether their median is $< M$ or $\geq M$:

$$Y_\prec := \{ y \in_s\ mset\ (\text{chop}\ 5\ xs) \mid \text{median}\ ys < M\}$$

$$Y_\succ := \{ y \in_s\ mset\ (\text{chop}\ 5\ xs) \mid \text{median}\ ys \geq M\}$$

We clearly have

$$\text{mset}\ xs = (\sum_{y \in\ \text{chop}\ 5\ xs}\ \text{mset}\ ys) \quad (3.9)$$

$$\text{mset}\ (\text{chop}\ 5\ xs) = Y_\prec + Y_\succ \quad (3.10)$$

$$\lfloor n / 5 \rfloor = |Y_\prec| + |Y_\succ| \quad (3.11)$$

and since $M$ is the median of the medians of the groups, we also know that:

$$|Y_\prec| < \frac{1}{2} \cdot \lfloor n / 5 \rfloor \quad (3.12)$$

The core idea of the proof is that any group $ys \in_s\ Y_\prec$ can have at most 2 elements that are $< M$:

$$\| y \in_s\ mset\ ys \mid y < M \| \leq \| y \in_s\ mset\ ys \mid y < \text{median}\ ys \|$$

$$\leq |ys| \div 2 \quad \text{because} \ y \in_s\ Y_\succ \quad \text{using (3.1)}$$

$$\leq 5 \div 2 = 2$$

And of course, since each group has size at most 5, any group in $ys \in_s\ Y_\succ$ can contribute at most 5 elements. In summary, we have:

$$\forall ys\in_s\ Y_\prec. \| y \in_s\ mset\ ys \mid y < M \| \leq 5$$

$$\forall ys\in_s\ Y_\succ. \| y \in_s\ mset\ ys \mid y < M \| \leq 2 \quad (3.13)$$
With this, we can begin our estimate of the number of elements \( M \):

\[
\{ y \in x \mid y < M \}
\]

\[
= \{ y \in \sum_{y \leftarrow \text{chop} 5 x} \mid y < M \}
\]

\[
= \sum_{y \leftarrow \text{chop} 5 x} \{ y \in \text{set y} \mid y < M \}
\]

\[
= \sum_{y \in \text{set}(X_\prec + Y_\succ)} \{ y \in \text{set y} \mid y < M \}
\]

using (3.9)

Taking the size of both sides, we have

\[
\| \{ y \in x \mid y < M \} \|
\]

\[
\leq \sum_{y \in \text{set}(Y_\prec + Y_\succ)} \| \{ y \in \text{set y} \mid y < M \} \|
\]

\[
= \sum_{y \in \text{set} Y_\prec} \| \{ y \in \text{set y} \mid y < M \} \| + \sum_{y \in \text{set} Y_\succ} \| \{ y \in \text{set y} \mid y < M \} \|
\]

\[
\leq (\sum_{y \in \text{set} Y_\prec} 5) + (\sum_{y \in \text{set} Y_\succ} 2)
\]

using (3.13)

\[
= 5 \cdot |Y_\prec| + 2 \cdot |Y_\succ|
\]

\[
= 2 \cdot \left[ \frac{n}{5} \right] + 2 \cdot |Y_\succ|
\]

\[
= 2 \cdot \left[ \frac{n}{5} \right] + 3 \cdot |Y_\prec| + 3 \cdot |Y_\succ|
\]

\[
= 2 \cdot \left[ \frac{n}{5} \right] + 3 \cdot \left[ \frac{n}{5} \right]
\]

\[
\leq 3.5 \cdot \left[ \frac{n}{5} \right]
\]

\[
\leq 0.7 \cdot n + 3
\]

The delicate arithmetic reasoning about rounding in the end can thankfully be done fully automatically by Isabelle’s linarith method.

\[\Box\]

### 3.3 Selection in Linear Time

We now have all the ingredients to write down our algorithm: the base cases (i.e. sufficiently short lists) can be handled using the naive approach of performing insertion sort and then returning the \( k \)-th element. For bigger lists, we perform the divide-and-conquer approach outlined in Theorem 3.5 using \( M \) as a pivot. We have two recursive calls: one on a list with exactly \( \lfloor 0.2 \cdot n \rfloor \) elements to compute \( M \), and one on a list with at most \( \lfloor 0.7 \cdot n + 3 \rfloor \).

We will still need to show later that this actually leads to a linear-time algorithm, but the fact that \( 0.7 + 0.2 < 1 \) is at least encouraging: intuitively, the “work load” is reduced by at least 10% in every recursive step, so we should reach the base case in a logarithmic number of steps.

The full algorithm looks like this:

```plaintext
chop :: nat ⇒ 'a list ⇒ 'a list list
chop 0 _ = []
chop _ [] = []
```
3.3 Selection in Linear Time

\[
\text{chop } s \ x s = \text{take } s \ x s \neq \text{chop } s \ (\text{drop } s \ x s)
\]

\[
\begin{align*}
\text{slow\_select} &: \text{nat} \Rightarrow 'a \ \text{list} \Rightarrow 'a \\
\text{slow\_select} \ k \ x s &= \text{insort} \ x s \ ! \ k \\
\text{slow\_median} &: 'a \ \text{list} \Rightarrow 'a \\
\text{slow\_median} \ x s &= \text{slow\_select} ((|x s| - 1) \div 2) \ x s \\
\text{mom\_select} &: \text{nat} \Rightarrow 'a \ \text{list} \Rightarrow 'a \\
\text{mom\_select} \ k \ x s &= \begin{cases}
\text{if } |x s| \leq 20 & \text{then } \text{slow\_select} \ k \ x s \\
\text{else let } M &= \text{mom\_select} \ ((|x s| / 5) - 1) \div 2) \text{\Big(} \text{map slow\_median (chop 5 x s)}; \text{\Big)}; \\
\text{\quad (ls, es, gs) = partition3 M x s} \\
\text{\quad in if } k < |ls| & \text{then } \text{mom\_select} \ k \ ls \\
\text{\quad else if } k < |ls| + |es| & \text{then } M \\
\text{\quad else } \text{mom\_select} \ (k - |ls| - |es|) \ gs \end{cases}
\end{align*}
\]

Correctness and termination are easy to prove:

**Theorem 3.7** (Partial Correctness of \(\text{mom\_select}\)). Let \(x s\) be a list and \(k < |x s|\). Then if \(\text{mom\_select} \ k \ x s\) terminates, we have

\[
\text{mom\_select} \ k \ x s = \text{select} \ k \ x s.
\]

**Proof.** Straightforward computation induction using Theorem 3.5. \(\square\)

**Theorem 3.8** (Termination of \(\text{mom\_select}\)). Let \(x s\) be a list and \(k < |x s|\). Then \(\text{mom\_select} \ k \ x s\) terminates.

**Proof.** We use \(|x s|\) as a termination measure. We need to show that it decreases in each of the two recursive calls under the precondition \(|x s| > 20\). This is easy to see:

- The list in the first recursive call has length \(|x s| / 5|\), which is strictly less than \(|x s|\) if \(|x s| > 1\).
- The length of the list in the second recursive call is at most \(|x s| - 1\): by induction hypothesis, the first recursive call terminates, so by Theorem 3.7 we know that

\[
M = \text{median} \ (\text{map median} \ (\text{chop 5 x s})) \text{ and thus:}
\]

\[
M \in \text{set} \ (\text{map median} \ (\text{chop 5 x s}))
\]

\[
= \{\text{median } y s \mid y s \in \text{set} \ (\text{chop 5 x s})\}.
\]
Chapter 3  Selection

\[ \subseteq \bigcup_{ys \in \text{set } (\text{chop} 5 \text{ } xs)} \text{set } ys = \text{set } xs \]

Hence, \( M \in \text{set } xs \) but \( M \notin \text{set } ls \) and \( M \notin \text{set } gs \) by construction. Since \( \text{set } xs \) and \( \text{set } ys \) are subsets of \( \text{set } xs \), this implies that \( |ls| < |zs| \) and \( |gs| < |zs| \).

So in either of the two cases for the second recursive call, the length decreases by at least 1.

Of course, we will later see that it actually decreases by quite a bit more than that, but this very crude estimate is sufficient to show termination.

Exercise 3.6. The recursive definition of \textit{mom\_select} handles the cases \( |zs| \leq 20 \) through the naive algorithm using insertion sort. The constant 20 here seems somewhat arbitrary. Find the smallest constant \( n_0 \) for which the algorithm still works. Why do you think 20 was chosen?

Note that in practice it may be sensible to choose a much larger cut-off size than 20 and handle shorter lists with a more direct approach that empirically works well for such short lists.

3.4  Time Functions

It remains to show now that this indeed leads to a linear-time algorithm. The time function for our selection algorithm is as follows:

\[
T_{\text{mom\_select}} :: \text{n} \Rightarrow \text{'}a \text{ list} \Rightarrow \text{n} \\
T_{\text{mom\_select}} \ k \ xs \\
= T_{\text{length}} \ xs + \\
(\text{if } |xs| \leq 20 \text{ then } T_{\text{slow\_select}} \ k \ xs \\
\text{else let } xss = \text{chop} \ 5 \ xs; \quad \ms = \text{map } \text{slow\_median} \ xss; \\
\quad \idx = (\lfloor |xs| / 5 \rfloor - 1) \text{ div } 2; \\
\quad x = \text{mom\_select } \idx \ ms; \\
\quad (ls, es, gs) = \text{partition3 } x \ xs \\
\quad \text{in } T_{\text{mom\_select}} \idx \ ms + T_{\text{chop}} \ 5 \ xs + T_{\text{map}} \ T_{\text{slow\_median}} \ xss + \\
\quad T_{\text{partition3}} \ x \ xs + T_{\text{length}} \ ls + 1 + \\
\quad (\text{if } k < |ls| \text{ then } T_{\text{mom\_select}} \ k \ ls \\
\quad \text{else if } k < |ls| + |es| \text{ then } T_{\text{length}} \ es \\
\quad \text{else } T_{\text{mom\_select}} (k - |ls| - |es|) \ gs + T_{\text{length}} \ es)))
\]

We can then prove
3.5 “Akra–Bazzi Light”

3.5 “Akra–Bazzi Light”

The function $T'_{\text{mom\_select}}$ (let us write it as $f$ for now) satisfies the recurrence

$$ n > 20 \rightarrow f(n) = f(0.2 \cdot n) + f(0.7 \cdot n + 3) + 19 \cdot n + 54 $$

Such divide-and-conquer recurrences are beyond the “normal” Master Theorem, but a generalisation, the Akra–Bazzi Theorem [Akra and Bazzi 1998, Eberl 2017b, Leighton 1996], does apply to them. Let us first abstract the situation a bit and consider the recurrence

$$ n > 20 \rightarrow f(n) = f(a \cdot n + b) + f(c \cdot n + d) + C_1 \cdot n + C_2 $$

where $0 < a, b < 1$ and $C_1, C_2 > 0$. The Akra–Bazzi Theorem then tells us that such a function is $O(n)$ if (and only if) $a + b < 1$. We will prove the relevant direction of this particular case of the theorem now — “Akra–Bazzi Light”, so to say.

Instead of presenting the full theorem statement and its proof right away, let us take a more explorative approach. What we want to prove in the end is that there are real constants $C_3 > 0$ and $C_4$ such that $f(n) \leq C_3 \cdot n + C_4$ for all $n$. Suppose

where the upper bound $T'_{\text{mom\_select}}$ is defined as follows:

$$ T'_{\text{mom\_select}} :: \text{nat} \Rightarrow \text{nat} $$

$$ T'_{\text{mom\_select}} n = (\text{if } n \leq 20 \text{ then } 482 \text{ else } T'_{\text{mom\_select}} [0.2 \cdot n] + T'_{\text{mom\_select}} [0.7 \cdot n + 3] + 19 \cdot n + 54) $$

The time functions of the auxiliary functions used here can be found in Section B.2 in the appendix. The proof is a simple computation induction using Theorem 3.6 and the time bounds for the auxiliary functions from Chapter B in the appendix.

The next section will be dedicated to showing that $T'_{\text{mom\_select}} \in O(n)$. 

Exercise 3.7. Show that the upper bound $[0.7 \cdot n + 3]$ is fairly tight by giving an infinite family $(xs_i)_{i \in \mathbb{N}}$ of lists with increasing lengths for which more than 70% of the elements are larger than the median of medians (with chopping size 5). In Isabelle terms: define a function $f :: \text{nat} \Rightarrow \text{nat\ list}$ such that

$\{ y \in \text{mset\ (f n)} | y > \text{mom\ (f n)} \} > 0.7$

where $\text{mom\ xs} = \text{median\ (map\ median\ (chop\ 5\ xs))}$.
we already knew such constants and now wanted to prove that the inequality holds. For the sake of simplicity of the presentation, we assume \( b, d \geq 0 \), but note that these assumptions are unnecessary and the proof still works for negative \( b \) and \( d \) if we replace \( b \) and \( d \) with \( \max 0 \ b \) and \( \max 0 \ d \).

The obvious approach to show this is by induction on \( n \), following the structure of the recurrence above. To do this, we use strong induction (i.e. the induction hypothesis holds for all \( m < n \))\(^1\) and a case analysis on \( n > n_1 \) (where \( n_1 \) is some constant we will determine later).

The two cases we have to show in the induction are then:

**Base case:** \( \forall n \leq n_1. \ f n \leq C_3 \cdot n + C_4 \)

**Step:** \( \forall n > n_1. (\forall m < n. f m \leq C_3 \cdot m + C_4) \rightarrow f n \leq C_3 \cdot n + C_4 \)

We can see that in order to even be able to apply the induction hypothesis in the induction step, we need \([a \cdot n + b] < n\). We can make the estimate\(^2\)

\[ [a \cdot n + b] \leq a \cdot n + b + 1 \leq n \]

and then solve for \( n \), which gives us \( n \geq \frac{b + 1}{1 - a} \). If we do the same for \( c \) and \( d \) as well, we get the conditions

\[ n_1 \geq \frac{b + 1}{1 - a} \quad \text{and} \quad n_1 \geq \frac{d + 1}{1 - c} \]

However, it will later turn out that these are implied by the other conditions we will have accumulated anyway.

Now that we have ensured that the basic structure of our induction will work out, let us continue with the two cases.

The base cases \((n \leq n_1)\) is fairly uninteresting: we can simply choose \( C_4 \) to be big enough to satisfy the equality for all \( n \leq n_1 \), whatever \( n_1 \) is.

In the recursive step, unfolding one step of the recurrence and applying the induction hypothesis leaves us with the proof obligation

\[ (C_3 \cdot [a \cdot n + b] + C_4) + (C_3 \cdot [c \cdot n + d] + C_4) + C_1 \cdot n + C_2 \leq C_3 \cdot n + C_4 , \]

or, equivalently,

\[ C_3 \cdot ([a \cdot n + b] + [c \cdot n + d] - n) + C_1 \cdot n + C_2 + C_4 \leq 0 \]

We estimate the left-hand side like this:

\(^1\)In Isabelle, the corresponding rule is called `less_induct`:

\[(\forall n. (\forall k < n. P k) \rightarrow P n) \rightarrow P n \ (\text{where } n :: \text{nat})\]

\(^2\)The notation \( \leq \) stands for “must be less than”. It emphasises that this inequality is not a consequence of what we have shown so far, but something that we still need to show, or in this case something that we need to ensure by adding suitable preconditions.
\[ C_3 \cdot (a \cdot n + b) + [c \cdot n + d] - n = C_1 \cdot n + C_2 + C_4 \]
\[ \leq C_3 \cdot ((b + d + 1) + (c \cdot n + d + 1) - n) = C_1 \cdot n + C_2 + C_4 \]
\[ = C_3 \cdot (b + d + 2) + C_2 + C_4 - (C_3 \cdot (1 - a - c) - C_1) \cdot n \]
\[ \leq C_3 \cdot (b + d + 2) + C_2 + C_4 - (C_3 \cdot (1 - a - c) - C_1) \cdot n_1 \]
\[ \leq 0 \]

The step from (\( \ast \)) to (\( \dagger \)) uses the fact that \( n > n_1 \) and requires the factor \( C_3 \cdot (1 - a - c) - C_1 \) in front of the \( n \) to be positive, i.e. we need to add the assumption
\[ C_3 > \frac{C_1}{1 - a - c}. \]  

(3.15)

The term (\( \dagger \)) (which we want to be \( \leq 0 \)) is now a constant. If we solve that inequality for \( C_3 \), we get the following two additional conditions:
\[ n_1 > \frac{b + d + 2}{1 - a - c} \quad \text{and} \quad C_3 \geq \frac{C_1 \cdot n_1 + C_2 + C_4}{(1 - a - c) \cdot n_1 - b - d - 2} \]

(3.16)

The former of these directly implies our earlier conditions (3.14), so we can safely discard those now.

Now all we have to do is to find a combination of \( n_1 \), \( C_3 \), and \( C_4 \) that satisfies (3.15) and (3.16). This is straightforward:
\[ n_1 := \max_{n_0} \left( \left\lfloor \frac{b + d + 2}{1 - a - c} \right\rfloor + 1 \right) \quad C_4 := \text{Max} \{f \mid n \leq n_1\} \]
\[ C_3 := \max \left( \frac{C_1}{1 - a - c} \right) \left( \frac{C_1 \cdot n_1 + C_2 + C_4}{(1 - a - c) \cdot n_1 - b - d - 2} \right) \]

And with that, the induction goes through and we get the following theorem:

Theorem 3.9 (Akra Bazzi Light).
\[ a > 0 \wedge c > 0 \wedge a + c < 1 \wedge C_1 \geq 0 \wedge \]
\[ (\forall n > n_0. \ f n = f [a \cdot n + b] + f [c \cdot n + d] + C_1 \cdot n + C_2) \quad \rightarrow \]
\[ (\exists C_3 \ C_4. \ \forall n. \ f n \leq C_3 \cdot n + C_4) \]

3.6 Conclusion

In this chapter, we have seen how to find the \( k \)-th largest element in a list containing \( n \) elements in time \( O(n) \), uniformly for all \( k \). Of course, we did not really talk about the constant coefficients that are hidden by the \( O(n) \) and which determine how efficient that algorithm is in practice. Although median-of-medians selection is guaranteed to run in worst-case linear time and therefore asymptotically time-optimal, other approaches with a worse worst-case running time like \( O(n \log n) \) or even \( O(n^2) \) may perform better in most situations in practice.
One solution to remedy this is to take a hybrid approach: we can use a selection algorithm that performs well in most situations (e.g. the divide-and-conquer approach from Section 3.1 with a fixed or a random pivot) and only resort to the guaranteed-linear-time algorithm if we notice that we are not making much progress. This is the approach taken by Musser’s Introselect algorithm [Musser 1997].

Exercise 3.8.

1. Suppose that instead of groups of 5, we now chop into groups of size \( l \geq 1 \). Prove a corresponding generalisation of Theorem 3.6.
2. Examine (on paper only): how does this affect correctness and running time of our selection algorithm? Why do you think \( l = 5 \) was chosen?
Part II

Search Trees
4

Binary Trees

Tobias Nipkow

Binary trees are defined as a recursive data type:

```
datatype 'a tree = Leaf | Node ('a tree) 'a ('a tree)
```

The following syntactic sugar is sprinkled on top:

```
⟨⟩ ≡ Leaf
⟨l, x, r⟩ ≡ Node l x r
```

The trees \( l \) and \( r \) are the left and right children of the node \( ⟨l, x, r⟩ \).

Because most of our trees will be binary trees, we drop the “binary” most of the time and have also called the type merely \( \text{tree} \).

When displaying a tree in the usual graphical manner we show only the \( \text{Nodes} \).
For example, \( ⟨⟨⟨⟩, 3⟩, ⟨⟩⟩, 9, ⟨⟨⟩, 7, ⟨⟩⟩⟩ \) is displayed like this:

```
\[
\begin{array}{c}
\text{9} \\
\text{3} \\
\text{7}
\end{array}
\]
```

The (label of the) root node is 9. The depth (or level) of some node (or leaf) in a tree is the distance from the root. The left spine of a tree is the sequence of nodes starting from the root and following the left child until that is a leaf. Dually for the right spine. We use these concepts only informally.

4.1 Basic Functions

Two canonical functions on data types are \( \text{set} \) and \( \text{map} \):

```
set_tree :: 'a tree ⇒ 'a set
set_tree ⟨⟩ = ∅
```

Chapter 4  Binary Trees

\[ \text{set_tree}(l, x, r) = \text{set_tree} l \cup \{x\} \cup \text{set_tree} r \]

\[ \text{map_tree} :: (\alpha \Rightarrow \beta) \Rightarrow \alpha \text{ tree} \Rightarrow \beta \text{ tree} \]
\[ \text{map_tree} f \; () = () \]
\[ \text{map_tree} f \; (l, x, r) = (\text{map_tree} f \; l, f \; x, \text{map_tree} f \; r) \]

The inorder, preorder and postorder traversals (we omit the latter) list the elements in a tree in a particular order:

\[ \text{inorder} :: \alpha \text{ tree} \Rightarrow \alpha \text{ list} \]
\[ \text{inorder} \; () = [] \]
\[ \text{inorder} \; (l, x, r) = \text{inorder} \; l @ [x] @ \text{inorder} \; r \]

\[ \text{preorder} :: \alpha \text{ tree} \Rightarrow \alpha \text{ list} \]
\[ \text{preorder} \; () = [] \]
\[ \text{preorder} \; (l, x, r) = x \# \text{preorder} \; l @ \text{preorder} \; r \]

These two size functions count the number of Nodes and Leafs in a tree:

\[ \text{size} :: \alpha \text{ tree} \Rightarrow \text{nat} \]
\[ |()| = 0 \]
\[ |(l, \_ , r)| = |l| + |r| + 1 \]

\[ \text{size1} :: \alpha \text{ tree} \Rightarrow \text{nat} \]
\[ |()|_1 = 1 \]
\[ |(l, \_ , r)|_1 = |l|_1 + |r|_1 \]

The syntactic sugar \(|t|\) for size \(t\) and \(|t|_1\) for size1 \(t\) is only used in this text, not in the Isabelle theories.

Induction proves a convenient fact that explains the name size1:

\[ |t|_1 = |t| + 1 \]

The height and the minimal height of a tree are defined as follows:

\[ \text{height} :: \alpha \text{ tree} \Rightarrow \text{nat} \]
\[ h \; () = 0 \]
4.1 Basic Functions

You can think of them as the longest and shortest (cycle-free) path from the root to a leaf. The real names of these functions are \textit{height} and \textit{min\_height}. The abbreviations \( h \) and \( mh \) are only used in this text, not in the Isabelle theories.

The obvious properties \( h t \leq |t| \) and \( mh t \leq h t \) and the following classical properties have easy inductive proofs:

\[ 2^{mh t} \leq |t|_1 \quad |t|_1 \leq 2^{h t} \]

We will simply use these fundamental properties without referring to them by a name or number.

The set of subtrees of a tree is defined as follows:

\[
\text{subtrees} :: \ 'a tree \Rightarrow 'a tree set \\
\text{subtrees} () = \{()\} \\
\text{subtrees} (l, a, r) = \{\langle l, a, r \rangle\} \cup \text{subtrees} l \cup \text{subtrees} r
\]

Note that every tree is a subtree of itself.

4.1.1 Exercises

Exercise 4.1. Function \textit{inorder} has quadratic complexity because the running time of \((\@)\) is linear in the length of its first argument. Define a function \textit{inorder2} :: \( 'a tree \Rightarrow 'a list \Rightarrow 'a list \) that avoids \((\@)\) but accumulates the result in its second parameter via \((\#)\) only. Its running time should be linear in the size of the tree. Prove \textit{inorder2} \( t \, xs = \text{inorder} \, t \, @ \, xs \).

Exercise 4.2. Write a function \textit{enum\_tree} :: \( 'a list \Rightarrow 'a tree list \) such that \( \text{set} \, (\text{enum\_tree} \, xs) = \{t \mid \text{inorder} \, t \, = \, xs\} \) and prove this proposition. You could also prove that \textit{enum\_tree} produces lists of \textit{distinct} elements, although that is likely to be harder.

Exercise 4.3. Although we focus on binary trees, arbitrarily branching trees can be defined just as easily:
Chapter 4  Binary Trees

4.2 Complete Trees

A complete tree is one where all the leaves are on the same level. An example is shown in Figure 4.1. The predicate `complete` is defined recursively:

```haskell
datatype 'a rtree = Nd 'a ("a rtree list)

Such trees are often called rose trees. Define a function `mir :: 'a rtree ⇒ 'a rtree` that mirrors a rose tree and prove `mir (mir t) = t`.

4.2 Complete Trees

A complete tree is one where all the leaves are on the same level. An example is shown in Figure 4.1. The predicate `complete` is defined recursively:

```haskell
complete :: 'a tree ⇒ bool
complete () = True
complete (l, _, r) = (h l = h r ∧ complete l ∧ complete r)
```

This recursive definition is equivalent with the above definition that all leaves must have the same distance from the root. Formally:

**Lemma 4.1.** `complete t ↝ mh t = h t`

**Proof** by induction and case analyses on `min` and `max`.

The following classic property of complete trees is easily proved by induction:

**Lemma 4.2.** `complete t → |t|₁ = 2^h t`

It turns out below that this is in fact a defining property of complete trees.

For complete trees we have `2^{mh t} ≤ |t|₁ = 2^h t`. For incomplete trees both ≤ and = become < as the following two lemmas prove:

**Lemma 4.3.** `¬ complete t → |t|₁ < 2^h t`

**Proof** by induction. We focus in the induction step where `t = ⟨l, x, r⟩`. If `t` is incomplete, there are a number of cases and we prove `|t|₁ < 2^h t` in each case. If `h l ≠ h r`, consider the case `h l < h r` (the case `h r < h l` is symmetric). From `2^{h l} <`
4.3 Almost Complete Trees

Almost complete trees are important because among all the trees with the same number of nodes they have minimal height.
Lemma 4.6. acomplete \( s \wedge |s| \leq |t| \rightarrow h \ s \leq h \ t \\
Proof by cases. If complete \( s \) then, by Lemma 4.2, \( 2^h \ s = |s|_1 \leq |t|_1 \leq 2^h \ t \) and thus \( h \ s \leq h \ t \). Now assume \( \neg \) complete \( s \). Then Lemma 4.4 yields \( 2^{mh} \ s < |s|_1 \leq |t|_1 \leq 2^h \ t \) and thus \( mh \ s < h \ t \). Furthermore we have \( h \ s = mh \ s \leq 1 \) (from acomplete \( s \)), \( h \ s \neq mh \ s \) (from Lemma 4.1) and \( mh \ s \leq h \ s \), which together imply \( mh \ s + 1 = h \ s \). With \( mh \ s < h \ t \) this implies \( h \ s \leq h \ t \).

This is relevant for search trees because their height determines the worst case running time. Almost complete trees are optimal in that sense.

The following lemma yields an explicit formula for the height of an almost complete tree:

Lemma 4.7. acomplete \( t \rightarrow h \ t = \lfloor \log |t|_1 \rfloor \)

Proof by cases. If \( t \) is complete, the claim follows from Lemma 4.2. Now assume \( t \) is incomplete. Then \( h \ t = mh \ t + 1 \) because acomplete \( t \), \( mh \ t \leq h \ t \) and complete \( t \) \( mh \ t = h \ t \) (Lemma 4.1). Together with \( |t|_1 \leq 2^h \ t \) this yields \( |t|_1 \leq 2^{mh} \ t + 1 \) and thus \( \log |t|_1 \leq mh \ t + 1 \). By Lemma 4.4 we obtain \( mh \ t < \log |t|_1 \). These two bounds for \( \log |t|_1 \) together imply the claimed \( h \ t = \lfloor \log |t|_1 \rfloor \).

In the same manner we also obtain:

Lemma 4.8. acomplete \( t \rightarrow mh \ t = \lfloor \log |t|_1 \rfloor \)

4.3.1 Converting a List into an Almost Complete Tree

We will now see how to convert a list \( xs \) into an almost complete tree \( t \) such that inorder \( t = xs \). If the list is sorted, the result is an almost complete binary search tree (see the next chapter). The basic idea is to cut the list in two halves, turn them into almost complete trees recursively and combine them. But cutting up the list in two halves explicitly would lead to an \( n \cdot \log n \) algorithm but we want a linear one.
Therefore we use an additional \texttt{nat} parameter to tell us how much of the input list should be turned into a tree. The remaining list is returned with the tree:

\begin{verbatim}
bal :: nat \Rightarrow 'a list \Rightarrow 'a tree \times 'a list

bal n xs
= (if n = 0 then ((), xs)
   else let m = n \div 2;
         (l, ys) = bal m xs;
         (r, zs) = bal (n - 1 - m) (tl ys)
         in ((l, hd ys, r), zs))
\end{verbatim}

The trick is not to chop \texttt{xs} in half but \texttt{n} because we assume that arithmetic is constant-time. Hence \texttt{bal} runs in linear time (see Exercise 4.6). Figure 4.3 shows the result of \texttt{bal 10 [0..9]}.

Balancing some prefix or all of a list or tree is easily derived:

\begin{verbatim}
bal_list :: nat \Rightarrow 'a list \Rightarrow 'a tree
bal_list n xs = fst (bal n xs)

balance_list :: 'a list \Rightarrow 'a tree
balance_list xs = bal_list \mid xs\mid xs

bal_tree :: nat \Rightarrow 'a tree \Rightarrow 'a tree
bal_tree n t = bal_list n (inorder t)
\end{verbatim}
Chapter 4  Binary Trees

balance_tree :: 'a tree ⇒ 'a tree
balance_tree t = bal_tree n t

4.3.1.1 Correctness

The following lemma clearly expresses that \( \text{bal} \ n \ \text{xs} \) turns the prefix of length \( n \) of \( \text{xs} \) into a tree and returns the corresponding suffix of \( \text{xs} \):

Lemma 4.9. \( n \leq |\text{xs}| \land \text{bal} \ n \ \text{xs} = (t, \text{zs}) \rightarrow \text{xs} = \text{inorder} \ t @ \text{zs} \land |t| = n \)

Proof  by complete induction on \( n \), assuming that the proposition holds for all values below \( n \). If \( n = 0 \) the claim is trivial. Now assume \( n \neq 0 \) and let \( m = n \div 2 \) and \( m' = n - 1 - m \) (and thus \( m, m' < n \)). From \( \text{bal} \ m \ \text{xs} = (l, \text{ys}) \), \( \text{bal} \ m' \ (tl \ ys) = (r, \text{zs}) \) and \( t = \langle l, \text{hd} \ ys, r \rangle \).

Because \( m < n \leq |\text{xs}| \) the induction hypothesis implies \( \text{xs} = \text{inorder} \ l @ \text{ys} \land |l| = m \ (**\). This in turn implies \( m' \leq |tl \ ys| \) and thus the induction hypothesis implies \( tl \ ys = \text{inorder} \ r @ \text{zs} \land |r| = m' \ (**\). Properties (**\) and (**\) together with \( t = \langle l, \text{hd} \ ys, r \rangle \) imply the claim \( \text{xs} = \text{inorder} \ t @ \text{zs} \land |t| = n \) because \( ys \neq [] \).

The corresponding correctness properties of the derived functions are easy consequences:

\[
\begin{align*}
n \leq |\text{xs}| & \quad \rightarrow \quad \text{inorder} \ (\text{bal_list} \ n \ \text{xs}) = \text{take} \ n \ \text{xs} \\
& \quad \text{inorder} \ (\text{balance_list} \ \text{xs}) = \text{xs} \\
n \leq |t| & \quad \rightarrow \quad \text{inorder} \ (\text{bal_tree} \ n \ t) = \text{take} \ n \ (\text{inorder} \ t) \\
& \quad \text{inorder} \ (\text{balance_tree} \ t) = \text{inorder} \ t
\end{align*}
\]

To prove that \( \text{bal} \) returns an almost complete tree we determine its height and minimal height.

Lemma 4.10. \( n \leq |\text{xs}| \land \text{bal} \ n \ \text{xs} = (t, \text{zs}) \rightarrow h \ t = \lfloor \lg \ (n + 1) \rfloor \)

Proof. The proof structure is the same as for Lemma 4.9 and we reuse the variable names introduced there. In the induction step we obtain the simplified induction hypotheses \( h \ l = \lfloor \lg \ (m + 1) \rfloor \) and \( h \ r = \lfloor \lg \ (m' + 1) \rfloor \). This leads to

\[
\begin{align*}
h \ t & = \max \ (h \ l) \ (h \ r) + 1 \\
& = h \ l + 1 \\
& = \lfloor \lg \ (m + 1) \rfloor + 1 \\
& = \lfloor \lg \ (n + 1) \rfloor \quad \text{by } (2.29)
\end{align*}
\]

The following complementary lemma is proved in the same way:

Lemma 4.11. \( n \leq |\text{xs}| \land \text{bal} \ n \ \text{xs} = (t, \text{zs}) \rightarrow mh \ t = \lfloor \lg \ (n + 1) \rfloor \)
4.4  Augmented Trees

By definition of complete and because \(|x| - |y| \leq 1\) we obtain that \(\text{bal}\) (and consequently the functions that build on it) returns an almost complete tree:

**Corollary 4.12.** \(n \leq |xs| \wedge \text{bal} n xs = (t, ys) \rightarrow \text{acomp}l e t\)

4.3.2 Exercises

**Exercise 4.5.** Find a formula \(B\) such that \(\text{acomp}l e t (l, x, \_r) = B\) where \(B\) may only contain the functions \(\text{acomp}l e t, \text{com}l e t, \text{heigh}t, \text{arithmetic and Boolean operations, l and r, but in particular not min\_heigh}t\) or \(\text{Node} = (\_, \_, \_\_))\). Prove \(\text{acomp}l e t (l, x, \_r) = B\).

**Exercise 4.6.** Prove that the running time of function \(\text{bal}\) is linear in its first argument. (Isabelle hint: you need to refer to \(\text{bal}\) as \(\text{Balance. bal}\).)

4.4  Augmented Trees

A tree of type \('<a\ tree\>'\) only stores elements of type \('a\). However, it is frequently necessary to store some additional information of type \('<b\)'\) in each node too, often for efficiency reasons. Typical examples are:

- The size or the height of the tree. Because recomputing them requires traversing the whole tree.
- Lookup tables where each key of type \('<a\)'\) is associated with a value of type \('<b\)'\).

In this case we simply work with trees of type \('<a \times b\ tree\>'\) and call them augmented trees. As a result we need to redefine a few functions that should ignore the additional information. For example, function \textit{inorder}, when applied to an augmented tree, should return an \'<a list\>'. Thus we redefine it in the obvious way:

\[
\text{inorder} :: (<a \times b\) \rightarrow [a] \\
\text{inorder} () = [] \\
\text{inorder} (l, (a, _), r) = \text{inorder} l @ a # \text{inorder} r
\]

Another example is \textit{set_tree} :: (<a \times b\) \rightarrow set. In general, if a function \(f\) is originally defined on type \('<a\ tree\>'\) but should ignore the \('<b\)'-values in an \('<a \times b\) tree\') then we assume that there is a corresponding revised definition of \(f\) on augmented trees that focuses on the \('<a\)'-values just like \textit{inorder} above does. Of course functions that do not depend on the information in the nodes, e.g. size and height, stay unchanged.

Note that there are two alternative redefinitions of \textit{inorder} (and similar functions):

\textit{map} \textit{fst} \circ \textit{Tree.inorder} or \textit{Tree.inorder} \circ \textit{map_tree} \textit{fst} where \textit{Tree.inorder} is the original function.
4.4.1 Maintaining Augmented Trees

Maintaining the 'b-values in a tree can be hidden inside a suitable smart version of Node that has only a constant time overhead. Take the example of augmentation by size:

\[
\begin{align*}
    sz :: ('a \times \text{nat}) \text{tree} & \Rightarrow \text{nat} \\
    sz \langle \rangle &= 0 \\
    sz \langle \_, (_, n), \_ \rangle &= n \\
    node_sz :: ('a \times \text{nat}) \text{tree} & \Rightarrow 'a \Rightarrow ('a \times \text{nat}) \text{tree} \Rightarrow ('a \times \text{nat}) \text{tree} \\
    node_sz \ l \ a \ r &= \langle l, (a, sz\ l + sz\ r + 1), r \rangle
\end{align*}
\]

A ('a \times \text{nat}) \text{tree} satisfies \text{invar\_sz} if the size annotation of every node is computed from its children as specified in node\_sz:

\[
\begin{align*}
    \text{invar\_sz} :: ('a \times \text{nat}) \text{tree} & \Rightarrow \text{bool} \\
    \text{invar\_sz} \langle \rangle &= \text{True} \\
    \text{invar\_sz} \langle l, (_, n), r \rangle &= (n = sz\ l + sz\ r + 1 \land \text{invar\_sz} \ l \land \text{invar\_sz} \ r)
\end{align*}
\]

This predicate is preserved by node\_sz (i.e. \text{invar\_sz} \ l \land \text{invar\_sz} \ r \rightarrow \text{invar\_sz} (node\_sz \ l \ a \ r)) and it guarantees that sz returns the size:

\[
\text{invar\_sz} \ t \rightarrow sz\ t = |t|
\]

We can generalize this example easily. Assume we have a constant zero :: 'b and a function \( f :: 'b \Rightarrow 'a \Rightarrow 'b \Rightarrow 'b \) which we iterate over the tree:

\[
\begin{align*}
    F :: ('a \times 'b) \text{tree} & \Rightarrow 'b \\
    F \langle \rangle &= \text{zero} \\
    F \langle l, (a, \_), r \rangle &= f (F\ l)\ a\ (F\ r)
\end{align*}
\]

This generalizes the definition of size. Let node\_f compute the 'b-value from the 'b-value of its children via \( f \):

\[
\begin{align*}
    b\_val :: ('a \times 'b) \text{tree} & \Rightarrow 'b \\
    b\_val \langle \rangle &= \text{zero}
\end{align*}
\]
4.4 Augmented Trees

\[
b_{\text{val}} \langle \_, \_, b \rangle = b
\]

\[
node_f :: (\langle a \times b \rangle) \text{tree} \Rightarrow \langle a \times b \rangle \text{tree} \Rightarrow (\langle a \times b \rangle) \text{tree}
\]

\[
node_f \text{ } l \text{ } a \text{ } r = \langle l, (a, f (b_{\text{val}} l) a (b_{\text{val}} r)), r \rangle
\]

If all 'b-values are computed as in \( node_f \)

\[
invar_f :: (\langle a \times b \rangle) \text{tree} \Rightarrow \text{bool}
\]

\[
invar_f \langle \rangle = \text{True}
\]

\[
invar_f \langle l, (a, b), r \rangle = (b = f (b_{\text{val}} l) a (b_{\text{val}} r) \land invar_f l \land invar_f r)
\]

then all 'b-values equal \( F \): \( invar_f t \rightarrow b_{\text{val}} t = F t \).

4.4.2 Exercises

Exercise 4.7. Augment trees by a pair of a Boolean and something else where the Boolean indicates whether the tree is complete or not. Define \( ch, node_{\text{ch}} \) and \( invar_{\text{ch}} \) as in Section 4.4.1 and prove the following properties:

\[
invar_{\text{ch}} t \rightarrow ch t = (\text{complete } t, ?)
\]

\[
invar_{\text{ch}} l \land invar_{\text{ch}} r \rightarrow invar_{\text{ch}} (node_{\text{ch}} l a r)
\]

Exercise 4.8. Assume type 'a is of class \text{linorder} and augment each \text{Node} with the maximum value in that tree. Following Section 4.4.1 (but mind the \text{option} type!) define \( mx :: (\langle a \times b \rangle) \text{tree} \Rightarrow 'b \text{option}, node_{\text{mx}} \) and \( invar_{\text{mx}} \) and prove

\[
invar_{\text{mx}} t \rightarrow
\]

\[
mx t = (\text{if } t = \langle \rangle \text{ then None else Some (Max (set_tree t)))}
\]

where \text{Max} is the predefined maximum operator on finite, non-empty sets.
The purpose of this chapter is threefold: to introduce binary search trees (BSTs), to discuss their correctness proofs, and to provide a first example of an abstract data type, a notion discussed in more detail in the next chapter.

Search trees are a means for storing and accessing collections of elements efficiently. In particular they can support sets and maps. We concentrate on sets. We have already seen function set_tree that maps a tree to the set of its elements. This is an example of an abstraction function that maps concrete data structures to the abstract values that they represent.

BSTs require a linear ordering on the elements in the tree (as in Chapter Sorting). For each node, the elements in the left child are smaller than the root and the elements in the right child are bigger:

\[
\text{bst} : (\forall \alpha \in \text{linorder}) \quad \text{tree} \Rightarrow \text{bool}
\]

\[
\text{bst} (\lambda) = \text{True}
\]

\[
\text{bst} (l, a, r) = ((\forall x \in \text{set_tree} l. \ x < a) \land (\forall x \in \text{set_tree} r. \ a < x) \land \text{bst} l \land \text{bst} r)
\]

This is an example of a (coincidentally almost complete) BST:

It is obvious how to search for an element in a BST by comparing the element with the root and descending into one of the two children if you have not found it yet. In the worst case this takes time proportional to the height of the tree. In later chapters we discuss a number of methods for ensuring that the height of the tree is logarithmic in its size. For now we ignore all efficiency considerations and permit our BSTs to degenerate. Thus we call them unbalanced.
Exercise 5.1. The above recursive definition of bst is not a direct translation of the description “For each node” given in the text. For a more direct translation define a function

\[ \text{nodes} :: \texttt{a tree} \Rightarrow (\texttt{a tree} \times \texttt{a} \times \texttt{a tree}) \text{ set} \]

that collects all the nodes as triples \((l, a, r)\). Now define \(\text{bst\_nodes} t = (\forall (l,a,r) \in \text{nodes} t. \ldots)\) and prove \(\text{bst\_nodes} t = \text{bst} t\).

5.1 Interface
Trees are concrete data types that provide the building blocks for realizing abstract data types like sets. The abstract type has a fixed interface, i.e. set of operations, through which the values of the abstract type can be manipulated. The interface hides all implementation detail. In the Search Tree part of the book we focus on the abstract type of sets with the following interface:

\[
\begin{align*}
\text{empty} :: & \texttt{'}s \\
\text{insert} :: & \texttt{'}a \Rightarrow \texttt{'}s \Rightarrow \texttt{'}s \\
\text{delete} :: & \texttt{'}a \Rightarrow \texttt{'}s \Rightarrow \texttt{'}s \\
\text{isin} :: & \texttt{'}s \Rightarrow \texttt{'}a \Rightarrow \texttt{bool}
\end{align*}
\]

where \(\texttt{'}s\) is the type of sets of elements of type \(\texttt{'}a\). Most of our implementations of sets will be based on variants of BSTs and will require a linear order on \(\texttt{'}a\), but the general interface does not require this. The correctness of an implementation of this interface will be proved by relating it back to HOL’s type \(\texttt{'}a\ \text{set}\) via an abstraction function, e.g. \texttt{set\_tree}.

5.2 Implementing Sets via unbalanced BSTs
So far we have compared elements via \(=\), \(\leq\) and \(<\). Now we switch to a comparator-based approach:

\[
\begin{align*}
\text{datatype} \ \texttt{cmp\_val} & = \texttt{LT} | \texttt{EQ} | \texttt{GT} \\
\text{cmp} :: (\texttt{'}a :: \text{linorder}) \Rightarrow \texttt{'}a \Rightarrow \texttt{'}a \Rightarrow \texttt{cmp\_val} \\
\text{cmp} x y & = (\textbf{if} \ x < y \ \textbf{then} \ \texttt{LT} \ \textbf{else if} \ x = y \ \textbf{then} \ \texttt{EQ} \ \textbf{else} \ \texttt{GT})
\end{align*}
\]

We will frequently phrase algorithms in terms of \(\texttt{cmp}, \texttt{LT}, \texttt{EQ} \text{ and } \texttt{GT}\) instead of \(<, =\) and \(>\). This leads to more symmetric code. If some type comes with its own primitive \(\texttt{cmp}\) function this can yield a speed-up over the above generic \(\texttt{cmp}\) function.

Below you find an implementation of the set interface in terms of BSTs. Functions \texttt{isin} and \texttt{insert} are self-explanatory. Deletion is more interesting.
5.2 Implementing Sets via unbalanced BSTs

empty :: 'a tree
empty = ()

isin :: 'a tree ⇒ 'a ⇒ bool
isin () _ = False
isin (l, a, r) x
   = (case cmp x a of LT ⇒ isin l x | EQ ⇒ True | GT ⇒ isin r x)

insert :: 'a ⇒ 'a tree ⇒ 'a tree
insert x () = ((), x, ())
insert x (l, a, r) = (case cmp x a of
   LT ⇒ (insert x l, a, r) |
   EQ ⇒ (l, a, r) |
   GT ⇒ (l, a, insert x r))

delete :: 'a ⇒ 'a tree ⇒ 'a tree
delete x () = ()
delete x (l, a, r) = (case cmp x a of
   LT ⇒ (delete x l, a, r) |
   EQ ⇒ if r = () then l else let (a', r') = split_min r in (l, a', r') |
   GT ⇒ (l, a, delete x r))

split_min :: 'a tree ⇒ 'a × 'a tree
split_min (l, a, r)
   = (if l = () then (a, r) else let (x, l') = split_min l in (x, (l', a, r)))

5.2.1 Deletion

Function delete deletes a from (l, a, r) (where r ≠ ()) by replacing a with a' and r by r' where

  a' is the leftmost (least) element of r, also called the inorder successor of a,
  r' is the remainder of r after removing a'.

We call this deletion by replacing. Of course one can also obtain a' as the inorder predecessor of a in l.

An alternative is to delete a from (l, a, r) by "joining" l and r:
Chapter 5  Binary Search Trees

We call this **deletion by joining**. The characteristic property of **join** is \(!inorder (join l r) = inorder l @ inorder r\).

The definition of **join** may appear needlessly complicated. Why not this much simpler version:

We focus on **delete**, deletion by replacing, in the rest of the chapter.

### Correctness

Why is the above implementation correct? Roughly speaking, because the implementations of **empty**, **insert**, **delete** and **isin** on type **'a tree** simulate the behaviour of
\{\}, \cup, \neg \text{ and } \in \text{ on type 'a set}. \text{ Taking the abstraction function into account we can formulate the simulation precisely:}

\begin{align*}
\text{set_tree empty} &= \{\} \\
\text{set_tree (insert x t)} &= \text{set_tree t} \cup \{x\} \\
\text{set_tree (delete x t)} &= \text{set_tree t} \setminus \{x\} \\
\text{isin t x} &= (x \in \text{set_tree t})
\end{align*}

However, the implementation only works correctly on BSTs. Therefore we need to add the precondition \text{bst t} to all but the first proposition. But why are we permitted to assume this precondition? Only because \text{bst} is an invariant of this implementation: \text{bst} holds for \text{empty}, and both \text{insert} and \text{delete} preserve \text{bst}. Therefore every tree that can be manufactured through the interface is a BST. Of course this adds another set of proof obligations for correctness, invariant preservation:

\begin{align*}
\text{bst empty} \\
\text{bst t} \rightarrow \text{bst (insert x t)} \\
\text{bst t} \rightarrow \text{bst (delete x t)}
\end{align*}

When looking at the abstract data type of sets from the user (or 'client') perspective, we would call the collection of all proof obligations for the correctness of an implementation the specification of the abstract type.

Exercise 5.3. Verify the implementation in Section 5.2 by showing all the proof obligations above, without the detour via sorted lists explained below.

### 5.4 Correctness Proofs

It turns out that direct proofs of the properties in the previous section are cumbersome — at least for \text{delete} and for proof assistants like Isabelle. Yet the correctness of the implementation is quite obvious to most (functional) programmers. Which is why most algorithm texts do not spend any time on functional correctness of search trees and concentrate on non-obvious structural properties that imply the logarithmic height of the trees — of course our simple BSTs do not guarantee the latter.

We will now present how the vague notion of “obvious” can be concretized and automated to such a degree that we do not need to discuss functional correctness of search tree implementations again in this book. This is because our approach is quite generic: it works not only for the BSTs in this chapter but also for the more efficient variants discussed in later chapters. The remainder of this section can be skipped if one is not interested in proof automation.
5.4.1 The Idea
The key idea [Nipkow 2016] is to express \( \text{bst} \) and \( \text{set\_tree} \) via \( \text{inorder} \):
\[
\text{bst} \ t = \text{sorted} \ (\text{inorder} \ t) \quad \text{and} \quad \text{set\_tree} \ t = \text{set} \ (\text{inorder} \ t)
\]
where
\[
\text{sorted} :: \ 'a \ list \Rightarrow \ \text{bool}
\]
\[
\text{sorted} \ [\] = \ True
\]
\[
\text{sorted} \ [\_] = \ True
\]
\[
\text{sorted} \ (x \ # \ y \ # \ zs) = (x < y \ \&\& \ \text{sorted} \ (y \ # \ zs))
\]
Note that this is “sorted w.r.t. \((<)\)” whereas in the chapter on sorting \( \text{sorted} \) was defined as “sorted w.r.t. \((\leq)\).

Instead of showing directly that BSTs implement sets, we show that they implement an intermediate specification based on lists (and later that the list-based specification implies the set-based one). We can assume that the lists are \( \text{sorted} \) because they are abstractions of BSTs. Insertion and deletion on sorted lists can be defined as follows:

\[
\text{ins\_list} :: \ 'a \Rightarrow \ 'a \ list \Rightarrow \ 'a \ list
\]
\[
\text{ins\_list} \ x \ [\] = [x]
\]
\[
\text{ins\_list} \ x \ (a \ # \ xs) = (\text{if} \ x < a \ \text{then} \ x \ # \ a \ # \ xs \ \text{else if} \ x = a \ \text{then} \ a \ # \ xs \ \text{else} \ a \ # \ \text{ins\_list} \ x \ xs)
\]
\[
\text{del\_list} :: \ 'a \Rightarrow \ 'a \ list \Rightarrow \ 'a \ list
\]
\[
\text{del\_list} \ _ \ [\] = [\]
\]
\[
\text{del\_list} \ x \ (a \ # \ xs) = (\text{if} \ x = a \ \text{then} \ xs \ \text{else} \ a \ # \ \text{del\_list} \ x \ xs)
\]

The abstraction function from trees to lists is function \( \text{inorder} \). The specification in Figure 5.1 expresses that \( \text{empty} \), \( \text{insert} \), \( \text{delete} \) and \( \text{isin} \) implement \([\] \), \( \text{ins\_list} \), \( \text{del\_list} \) and \( \lambda xs. \ x \in \text{set} \ xs \). One nice aspect of this specification is that it does not require us to prove invariant preservation explicitly: it follows from the fact (proved below) that \( \text{ins\_list} \) and \( \text{del\_list} \) preserve \( \text{sorted} \).

5.4.2 BSTs Implement Sorted Lists — A Framework
We present a library of lemmas that automate the functional correctness proofs for the BSTs in this chapter and the more efficient variants in later chapters. This library
5.4 Correctness Proofs

\[ inorder\ empty = [] \]

\[ sorted\ (inorder\ t) \rightarrow inorder\ (insert\ x\ t) = ins\_list\ x\ (inorder\ t) \]

\[ sorted\ (inorder\ t) \rightarrow inorder\ (delete\ x\ t) = del\_list\ x\ (inorder\ t) \]

\[ sorted\ (inorder\ t) \rightarrow isin\ t\ x = (x \in\ set\ (inorder\ t)) \]

**Figure 5.1** List-based Specification of BSTs

is motivated by general considerations concerning the shape of formulas that arise during verification.

As a motivating example we examine how to prove

\[ sorted\ (inorder\ t) \rightarrow inorder\ (insert\ x\ t) = ins\_list\ x\ (inorder\ t) \]

The proof is by induction on \( t \) and we consider the case \( t = \langle l, a, r \rangle \) such that \( x < a \). Ideally the proof looks like this:

\[
\begin{align*}
inorder\ (insert\ x\ t) &= inorder\ (insert\ x\ l) \circ a \# inder\ r \\
&= ins\_list\ x\ (inorder\ l) \circ a \# inorder\ r \\
&= ins\_list\ x\ (inorder\ l \circ a \# inorder\ r) = ins\_list\ x\ t
\end{align*}
\]

The first and last step are by definition, the second step by induction hypothesis, and the third step by lemmas in Figure 5.2: (5.1) rewrites the assumption \( sorted\ (inorder\ t) \) to \( sorted\ (inorder\ l \circ [a]) \land sorted\ (a \# inorder\ r) \), thus allowing (5.5) to rewrite \( ins\_list\ x\ (inorder\ l \circ a \# inorder\ r) \) to \( ins\_list\ x\ (inorder\ l) \circ a \# inorder\ r \).

The lemma library in Figure 5.2 helps to prove the properties in Figure 5.1. These proofs are by induction on \( t \) and lead to (possibly nested) tree constructor terms like \( \langle t_1, a_1, t_2 \rangle \) where the \( t_i \) and \( a_i \) are variables. Evaluating \( inorder\ ) of such a tree leads to a list of the following form:

\[ inorder\ t_1 \circ a_1 \# inorder\ t_2 \circ a_2 \# \ldots \# inorder\ t_n \]

Now we discuss the lemmas in Figure 5.2 that simplify the application of \( sorted\ ), \( ins\_list\ ) and \( del\_list\ ) to such terms.

Terms of the form \( sorted\ (x_1 \circ a_1 \# x_2 \circ a_2 \# \ldots \# x_n) \) are decomposed into the following basic formulas

\[
\begin{align*}
sorted\ (x \circ [a]) & \quad \text{(simulating } \forall x \in set.\ x < a) \\
sorted\ (a \# xs) & \quad \text{(simulating } \forall x \in set.\ a < x) \\
a < b &
\end{align*}
\]

by the rewrite rules (5.1)–(5.2). Lemmas (5.3)–(5.4) enable deductions from basic formulas.
**Chapter 5  Binary Search Trees**

\[
\text{sorted } (xs @ y # ys) = (\text{sorted } (xs @ [y]) \land \text{sorted } (y # ys)) \quad (5.1)
\]

\[
\text{sorted } (x # xs @ y # ys) = (\text{sorted } (x # xs) \land x < y \land \text{sorted } (xs @ [y]) \land \text{sorted } (y # ys)) \quad (5.2)
\]

\[
\text{sorted } (x # xs) \longrightarrow \text{sorted } xs \quad (5.3)
\]

\[
\text{sorted } (xs @ [y]) \longrightarrow \text{sorted } xs \quad (5.4)
\]

\[
\text{sorted } (xs @ [a]) \implies \text{ins_list } x (xs @ a # ys) = \begin{cases} 
\text{if } x < a \text{ then } \text{ins_list } x xs @ a # ys \text{ else } xs @ \text{ins_list } x (a # ys) 
\end{cases} \quad (5.5)
\]

\[
\text{sorted } (xs @ a # ys) \implies \text{del_list } x (xs @ a # ys) = \begin{cases} 
\text{if } x < a \text{ then } \text{del_list } x xs @ a # ys \text{ else } xs @ \text{del_list } x (a # ys) 
\end{cases} \quad (5.6)
\]

\[
\text{sorted } (x # xs) = ((\forall y \in \text{set } xs. x < y) \land \text{sorted } xs) \quad (5.7)
\]

\[
\text{sorted } (xs @ [x]) = (\text{sorted } xs \land (\forall y \in \text{set } xs. y < x)) \quad (5.8)
\]

---

**Figure 5.2**  Lemmas for \textit{sorted}, \textit{ins_list}, \textit{del_list}

Terms of the form \(\text{ins_list } x (xs_1 @ a_1 # xs_2 @ a_2 # \ldots # xs_n)\) are rewritten with (5.5) (and the defining equations for \textit{ins_list}) to push \textit{ins_list} inwards. Terms of the form \(\text{del_list } x (xs_1 @ a_1 # xs_2 @ a_2 # \ldots # xs_n)\) are rewritten with (5.6) (and the defining equations for \textit{del_list}) to push \textit{del_list} inwards. The \textit{isin} property in Figure 5.1 can be proved with the help of (5.1), (5.7) and (5.8).

The lemmas in Figure 5.2 form the complete set of basic lemmas on which the automatic proofs of almost all search trees in the book rest; only splay trees (see Chapter 20) need additional lemmas.

### 5.4.3 Sorted Lists Implement Sets

It remains to be shown that the list-based specification (Figure 5.1) implies the set-based correctness properties in Section 5.3. Because \(\text{bst } t = \text{sorted } (\text{inorder } t)\), the latter correctness properties become

\[
\text{set_tree empty} = \{\}
\]

\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{set_tree } (\text{insert } x t) = \text{set_tree } t \cup \{x\}
\]

\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{set_tree } (\text{delete } x t) = \text{set_tree } t - \{x\}
\]

\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{isin } t x = (x \in \text{set_tree } t)
\]

\[
\text{sorted } (\text{inorder empty})
\]

\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{sorted } (\text{inorder } (\text{insert } x t))
\]

\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{sorted } (\text{inorder } (\text{delete } x t))
\]
They are proved directly by composing the list-based specification (Figure 5.1, proved above) with the correctness of the sorted list implementation of sets

\[
\begin{align*}
\text{set } (\text{ins}_\text{list} \ x \ \text{xs}) &= \text{set } \text{xs} \cup \{x\} \\
\text{sorted } \text{xs} &\rightarrow \text{set } (\text{del}_\text{list} \ x \ \text{xs}) = \text{set } \text{xs} - \{x\} \\
\text{sorted } \text{xs} &\rightarrow \text{sorted } (\text{ins}_\text{list} \ x \ \text{xs}) \\
\text{sorted } \text{xs} &\rightarrow \text{sorted } (\text{del}_\text{list} \ x \ \text{xs})
\end{align*}
\]

(which have easy inductive proofs) using set_tree \( t \equiv \text{set } (\text{inorder } t) \).

5.5 Tree Rotations

As discussed in the introduction to this chapter, the BST on the left is better than the one on the right, which has degenerated to a list:

On average, searching for a random key is faster in the left than in the right BST, assuming that all keys are equally likely. In later chapters, a number of balancing schemas will be presented that guarantee logarithmic height (in the number of nodes) of trees balanced according to those schemas. The basic balancing mechanisms are rotations, local tree transformations that preserve inorder but modify the shape:

\[
\begin{align*}
\text{right-rotation} &\\
\text{left-rotation}
\end{align*}
\]

We will now show that any two trees \( t_1 \) and \( t_2 \) with the same inorder can be transformed into each other by a linear number of rotations. The basic idea is simple. Transform \( t_1 \) into a list-like tree \( l \) by right-rotations. In order to transform \( l \) into \( t_2 \), note that we can transform \( t_2 \) into \( l \) (because inorder \( t_1 = \text{inorder } t_2 \)). Hence we merely need to reverse the transformation of \( t_2 \) into \( l \).

We call a tree in list-form if it is of the form
Formally:

\[
\text{is\_list :: 'a tree } \Rightarrow \text{ bool}
\]

\[
\text{is\_list } \langle l, \_ , r \rangle = (l = \langle \rangle \land \text{is\_list } r)
\]

\[
\text{is\_list } \langle \rangle = \text{ True}
\]

A tree is in list-form iff no right-rotation is applicable anywhere in the tree. The following function performs right-rotations in a top-down manner along the right spine of a tree by replacing subtrees of the form \( \langle A, a, B, b, C \rangle \) by \( A, a, \langle B, b, C \rangle \):

\[
\text{list\_of :: 'a tree } \Rightarrow \text{ 'a tree}
\]

\[
\text{list\_of } \langle \langle A, a, B \rangle, b, C \rangle = \text{ list\_of } \langle A, a, \langle B, b, C \rangle \rangle
\]

\[
\text{list\_of } \langle \langle \rangle, a, A \rangle = \langle \langle \rangle, a, \text{ list\_of } A \rangle
\]

\[
\text{list\_of } \langle \rangle = \langle \rangle
\]

The termination of this function may not be obvious. The problem is the first equation because the of size of \( \langle A, a, B \rangle \) and \( \langle A, a, \langle B, b, C \rangle \rangle \) are the same. However, the right spine has become one longer, which must end when all nodes of the tree are on the right spine. This suggests the measure function \( \lambda t. |t| - \text{rlen } t \) where

\[
\text{rlen :: 'a tree } \Rightarrow \text{ nat}
\]

\[
\text{rlen } \langle \rangle = 0
\]

\[
\text{rlen } \langle \_ , \_ , r \rangle = \text{ rlen } r + 1
\]

This works for the first \( \text{list\_of } \) equation but not for the second one: \( |\langle \langle \rangle, a, A \rangle| - \text{rlen } \langle \langle \rangle, a, A \rangle = |A| - \text{rlen } A \). Luckily the measure function \( \lambda t. 2 \cdot |t| - \text{rlen } t \) decreases with every recursive call, thus proving termination.

The correctness of \( \text{list\_of } \) is easily expressed

\[
\text{is\_list } (\text{list\_of } t)
\]

\[
\text{inorder } (\text{list\_of } t) = \text{ inorder } t
\]

and proved by computation induction.

The claim that only a linear number of rotations is needed cannot be proved from function \( \text{list\_of } \) because it does not count the rotations (but see Exercise 5.4). More problematic is the fact that we cannot formalize the second step of our overall proof,
namely the idea of reversing the sequence of rotations that list_of performs because the rotations are hidden inside list_of. Thus we abandon this formalization and restart by introducing an explicit notion of position (type pos) in a tree:

```plaintext
datatype dir = L | R

type_synonym pos = dir list
```

The position of a node in a tree is a sequence of left/right directions. They encode how to reach that node from the root by turning left or right at each successive node. For example, the position of ⟨⟨⟩⟩, 1, ⟨⟩⟩ in ⟨⟨⟨⟩⟩, 0, ⟨⟩⟩, 1, ⟨⟩⟩, 2, ⟨⟩⟩, 3, ⟨⟩⟩⟩ is [L, R].

Function rotR_pos is the analogue of list_of but whereas list_of returns the rotated tree, rotR_pos produces the list of positions where the rotations should be applied:

```plaintext
rotR_pos :: 'a tree ⇒ pos list
rotR_pos ⟨⟨A, a, B⟩⟩, b, C⟩ = [] # rotR_pos ⟨⟨A, a, ⟨B, b, C⟩⟩
rotR_pos ⟨⟨⟩⟩, _, A⟩ = map ((#) R) (rotR_pos A)
rotR_pos ⟨⟩ = []
```

Termination is again proved with the help of the measure function \(\lambda t . 2 \cdot |t| - \text{rlen } t\).

Functions apply_at and apply_ats perform a transformation at a (list of) position(s):

```plaintext
apply_at :: ('a tree ⇒ 'a tree) ⇒ pos ⇒ 'a tree ⇒ 'a tree
apply_at f [] t = f t
apply_at f (L # ds) (l, a, r) = (apply_at f ds l, a, r)
apply_at f (R # ds) (l, a, r) = (l, a, apply_at f ds r)

apply_ats :: ('a tree ⇒ 'a tree) ⇒ pos list ⇒ 'a tree ⇒ 'a tree
apply_ats _ [] t = t
apply_ats f (p # ps) t = apply_ats f ps (apply_at f p t)
```

We are interested in left and right rotations:

```plaintext
rotR :: 'a tree ⇒ 'a tree
```
Chapter 5  Binary Search Trees

rotR \langle \langle A, a, B \rangle, b, C \rangle = \langle A, a, \langle B, b, C \rangle \rangle

rotL :: 'a tree \Rightarrow 'a tree
rotL \langle A, a, \langle B, b, C \rangle \rangle = \langle \langle A, a, B \rangle, b, C \rangle \rangle

rotRs \equiv \text{apply}_ats \text{rotR}
rotLs \equiv \text{apply}_ats \text{rotL}

Now we can prove by computation induction that rotRs (rotR\_poss \ t) transforms \ t into list-form and preserves inorder
\[
is_list (\text{rotRs} (\text{rotR\_poss} \ t)) \quad \text{(5.9)}
\]
\[
inorder (\text{rotRs} (\text{rotR\_poss} \ t)) = inorder \ t \quad \text{(5.10)}
\]
using the inductive lemma
\[
\text{apply}_ats f (\text{map} ((\#) R) p s) \langle l, a, r \rangle = \langle l, a, \text{apply}_ats f \ p s \ r \rangle \quad \text{(5.11)}
\]
Moreover, we can now express and prove how many right-rotations are required:
\[
|\text{rotR\_poss} \ t| = |\ t| - \text{rlen} \ t \quad \text{(5.12)}
\]
The reason: each right-rotation moves one more node onto the right spine. The proof is by computation induction and uses an easy inductive fact: \( \text{rlen} \ t \leq |\ t| \).

Thus the number of right-rotations to reach list-form is upper-bounded by \( |\ t| \). In fact, (5.12) implies an upper bound of \( |\ t| - 1 \) because \( |\ t| - \text{rlen} \ t \leq |\ t| - 1 \) (why?). This upper bound is tight: any tree with only one node on the right spine needs that many right-rotations because each right-rotation increases \( \text{rlen} \) only by one.

At last we return to the original question, how to transform any tree into any other tree by rotations. The key lemma, which we can express at last, is that reversing the transformation to list-form takes us back to the original tree:
\[
\text{rotLs} (\text{rev} (\text{rotR\_poss} \ t)) (\text{rotRs} (\text{rotR\_poss} \ t)) \ t = \ t \quad \text{(5.13)}
\]
The proof is an easy computation induction using (5.11), the fact that \( \text{map} \) and \( \text{rev} \) commute and the easy inductive fact
\[
\text{apply}_ats f (\ p s_1 @ p s_2) \ t = \text{apply}_ats f \ p s_2 (\text{apply}_ats f \ p s_1 \ t)
\]
With this easy inductive proposition
\[
is_list \ t_1 \land is_list \ t_2 \land inorder \ t_1 = inorder \ t_2 \rightarrow t_1 = t_2 \quad \text{(5.14)}
\]
we can finally transform any \( t_1 \) into any \( t_2 \) by rotations if \( inorder \ t_1 = inorder \ t_2 \). First observe that
5.6 Case Study: Interval Trees

In this section we study binary trees for representing a set of intervals, called interval trees. In addition to the usual insertion and deletion functions of standard BSTs, interval trees support a function for determining whether a given interval overlaps with some interval in the tree.

5.6.1 Augmented BSTs

The efficient implementation of the search for an overlapping interval relies on an additional piece of information in each node. Thus interval trees are another example of augmented trees as introduced in Section 4.4. We reuse the modified definitions of set_tree and inorder from that section. Moreover we use a slightly adjusted version of isin:

\[
isin :: ('a \times 'b) \text{tree} \Rightarrow 'a \Rightarrow \text{bool}
\]

\[
isin \langle \rangle \_ = \text{False}
\]

\[
isin \langle l, (a, \_), r \rangle x
= (\text{case } \text{cmp} x a \text{ of } \text{LT} \Rightarrow isin \text{ l x} \mid \text{EQ} \Rightarrow \text{True} \mid \text{GT} \Rightarrow isin \text{ r x})
\]

This works for any kind of augmented BST, not just interval trees.
5.6.2 Intervals
An interval ‘a ivl is simply a pair of lower and upper bound, accessed by functions low and high, respectively. Intuitively, an interval represents the closed set between low and high. The standard mathematical notation is \([l, h]\), the Isabelle notation is \(\{l..h\}\). We restrict ourselves to non-empty intervals:

\[
\text{low } p \leq \text{high } p
\]

Type ‘a can be any linearly ordered type with a minimum element \(\bot\) (for example, the natural numbers or the real numbers extended with \(-\infty\)). Intervals can be linearly ordered by first comparing low, then comparing high. The definitions are as follows:

\[
\begin{align*}
(x < y) &= (\text{low } x < \text{low } y \lor \text{low } x = \text{low } y \land \text{high } x < \text{high } y) \\
(x \leq y) &= (\text{low } x < \text{low } y \lor \text{low } x = \text{low } y \land \text{high } x \leq \text{high } y)
\end{align*}
\]

Two intervals overlap if they have at least one point in common:

\[
\text{overlap } x y = (\text{low } y \leq \text{high } x \land \text{low } x \leq \text{high } y)
\]

The readers should convince themselves that overlap does what it is supposed to do:

\[
\text{overlap } x y = (\{\text{low } x..\text{high } x\} \cap \{\text{low } y..\text{high } y\} \neq \{\})
\]

We also define the concept of an interval overlapping with some interval in a set:

\[
\text{has_overlap } S y = (\exists x \in S. \text{overlap } x y)
\]

5.6.3 Interval Trees
An interval tree associates to each node a number max_hi, which records the maximum high value of all intervals in the subtrees. This value is updated during insert and delete operations, and will be crucial for enabling efficient determination of overlap with some interval in the tree.

\[
\begin{align*}
\text{type_synonym } 'a \text{ ivl\_tree} &= ('a \text{ ivl} \times 'a) \text{ tree} \\
\text{max\_hi} :: 'a \text{ ivl\_tree} \Rightarrow 'a \\
\text{max\_hi} \bot &= \bot \\
\text{max\_hi} (_, (_, m), _) &= m
\end{align*}
\]

If the max_hi value of every node in a tree agrees with max3.
5.6 Case Study: Interval Trees

5.6.4 Implementing Sets of Intervals via Interval Trees

Interval trees can implement sets of intervals via unbalanced BSTs as in Section 5.2. Of course empty = (). Function isin was already defined in Section 5.6.1. Insertion and deletion are also very close to the versions in Section 5.2, but the value of max_hi must be computed (by max3) for each new node. We follow Section 4.4 and introduce a smart constructor node for that purpose and replace (l, a, r) by node l a r (on the right-hand side):

inv_max_hi :: 'a ivl_tree ⇒ bool
inv_max_hi () = True
inv_max_hi (l, (a, m), r)
= (m = max3 a (max_hi l) (max_hi r) ∧ inv_max_hi l ∧
  inv_max_hi r)

max3 :: 'a ivl ⇒ 'a ⇒ 'a ⇒ 'a
max3 a m n = max (high a) (max m n)

it follows by induction that max_hi is the maximum value of high in the tree and comes from some node in the tree:

Lemma 5.1. inv_max_hi t ∧ a ∈ set_tree t → high a ≤ max_hi t
Lemma 5.2. inv_max_hi t ∧ t ≠ () → (∃a ∈ set_tree t. high a = max_hi t)

node :: 'a ivl_tree ⇒ 'a ivl ⇒ 'a ivl_tree ⇒ 'a ivl_tree
node l a r = (l, (a, max3 a (max_hi l) (max_hi r)), r)

insert :: 'a ivl ⇒ 'a ivl_tree ⇒ 'a ivl_tree
insert x () = (x, high x), ()
insert x (l, (a, m), r) = (case cmp x a of
  LT ⇒ node (insert x l) a r |
  EQ ⇒ (l, (a, m), r) |
  GT ⇒ node l a (insert x r))

split_min :: 'a ivl_tree ⇒ 'a ivl × 'a ivl_tree
split_min (l, (a, _), r)
= (if l = () then (a, r)
else let \((x, l') = \text{split\_min } l\text{ in } (x, \text{node } l' a r))

\[
delete :: 'a \text{ ivl} \Rightarrow 'a \text{ ivl\_tree} \Rightarrow 'a \text{ ivl\_tree}
\]
\[
delete _\ell () = ()
\]
\[
delete x (l, (a, _), r) = (\text{case } \text{cmp } x a \text{ of}
\]
\[
\quad LT \Rightarrow \text{node } (\text{delete } x l) a r |
\]
\[
\quad EQ \Rightarrow \text{if } r = () \text{ then } l \text{ else let } (x, y) = \text{split\_min } r \text{ in } \text{node } l x y |
\]
\[
\quad GT \Rightarrow \text{node } l a (\text{delete } x r))
\]

The correctness proofs for insertion and deletion cover two aspects. Functional correctness and preservation of the invariant \(\text{sorted } \circ \text{inorder}\) (the BST property) are proved exactly as in Section 5.3 for ordinary BSTs. Preservation of the invariant \(\text{inv\_max\_hi}\) can be proved by a sequence of simple inductive properties. In the end the main correctness properties are

\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{inorder } (\text{insert } x t) = \text{ins\_list } x (\text{inorder } t)
\]
\[
\text{sorted } (\text{inorder } t) \longrightarrow \text{inorder } (\text{delete } x t) = \text{del\_list } x (\text{inorder } t)
\]
\[
\text{inv\_max\_hi } t \longrightarrow \text{inv\_max\_hi } (\text{insert } x t)
\]
\[
\text{inv\_max\_hi } t \longrightarrow \text{inv\_max\_hi } (\text{delete } x t)
\]

Defining \(\text{invar } t = (\text{inv\_max\_hi } t \land \text{sorted } (\text{inorder } t))\) we obtain the following top-level correctness corollaries:

\[
\text{invar } s \longrightarrow \text{set\_tree } (\text{insert } x s) = \text{set\_tree } s \cup \{x\}
\]
\[
\text{invar } s \longrightarrow \text{set\_tree } (\text{delete } x s) = \text{set\_tree } s - \{x\}
\]
\[
\text{invar } s \longrightarrow \text{invar } (\text{insert } x s)
\]
\[
\text{invar } s \longrightarrow \text{invar } (\text{delete } x s)
\]

The above insertion function allows overlapping intervals to be added into the tree and deletion supports only deletion of whole intervals. This is appropriate for the computational geometry application sketched below in Subsection 5.6.6. Other applications may require a different design.

### 5.6.5 Searching for an Overlapping Interval

The added functionality of interval trees over ordinary BSTs is function \text{search} that searches for an overlapping rather than identical interval:
search :: 'a ivl_tree ⇒ 'a ivl ⇒ bool

search (\) _ = False
search (l, (a, _), r) x
  = (if overlap x a then True
     else if l \ 6\ (\) ∧ low x ≤ max_hi l then search l x else search r x)

The following theorem expresses the correctness of search assuming the same invariants as before; bst t would work just as well as sorted (inorder t).

Theorem 5.3. inv_max_hi t ∧ sorted (inorder t) \arrow search t x = has_overlap (set_tree t) x

Proof. The result is clear when t is (\). Now suppose t is in the form (l, (a, m), r), where m is the value of max_hi at root. If a overlaps with x, search returns True as expected. Otherwise, there are two cases.

- If l \ 6\ (\) and low x ≤ max_hi l, the search goes to the left child. If there is an interval in the left child overlapping with x, then the search returns True as expected. Otherwise, we show there is also no interval in the right child overlapping with x. Since l \ 6\ (\), Lemma 5.2 yields a node p in the left child such that high p = max_hi l. Since low x ≤ max_hi l, we have low x ≤ high p. Since p does not overlap with x, we must have high x < low p. But then, for every interval rp in the right child, low p ≤ low rp, so that high x < low rp, which implies that rp does not overlap with x.

- Now we consider the case where either l = (\) or max_hi l < low x. In this case, the search goes to the right. We show there is no interval in the left child that overlaps with x. This is clear if l = (\). Otherwise, for each interval lp, we have high lp ≤ max_hi l by Lemma 5.1, so that high lp < low x, which means lp does not overlap with x.

Exercise 5.7. Define a function that determines if a given point is in some interval in a given interval tree. Starting with

\[
in_{ivl} :: 'a ⇒ 'a ivl ⇒ bool\]
\[
in_{ivl} \ x \ iv = (\text{low} \ iv \ ≤ \ x \ ∧ \ x \ ≤ \ \text{high} \ iv)\]

write a recursive function

\[
\text{search1} :: 'a ivl_tree ⇒ 'a ⇒ bool\]

(without using search) such that search1 x t is True iff there is some interval iv in t such that in_{ivl} x iv. Prove

\[
\text{inv_max_hi t ∧ bst t \arrow search1 t x = (∃ iv ∈ set_tree t. in_{ivl} x iv)}\]
5.6.6 Application

While this section demonstrated how to augment an ordinary binary tree with intervals, any of the balanced binary trees (such as red-black tree) can be augmented in a similar manner. We leave this as exercises.

Interval trees have many applications in computational geometry. As a basic example, consider a set of rectangles whose sides are aligned to the $x$ and $y$-axes. We wish to efficiently determine whether any pair of rectangles in the set intersect each other (i.e. sharing a point, including boundaries). This can be done using a "sweep line" algorithm as follows. For each rectangle $[x_l, x_h] \times [y_l, y_h]$, we create two events: insert interval $[x_l, x_h]$ at $y$-coordinate $y_l$ and delete interval $[x_l, x_h]$ at $y$-coordinate $y_h$. Perform the events, starting from an empty interval tree, in ascending order of $y$-coordinates, with insertion events performed before deletion events. At each insertion, check whether the interval to be inserted overlaps with any of the existing intervals in the tree. If yes, we have found an intersection between two rectangles. If no overlap of intervals is detected throughout the process, then no pair of rectangles intersect. When using an interval tree based on a balanced binary tree, the time complexity of this procedure is $O(n \lg n)$, where $n$ is the number of rectangles.

5.7 Chapter Notes

Tree Rotations and Distance Culik II and Wood [1982] defined the rotation distance of two trees $t_1$ and $t_2$ with the same number of nodes $n$ as the minimum number of rotations needed to transform $t_1$ into $t_2$ and showed that it is upper-bounded by $2n - 2$. This result was improved by Sleator et al. [1986] and Pournin [2014] who showed that for $n \geq 11$ the maximum rotation distance is exactly $2n - 6$. The complexity of computing the rotation distance is open: it is in NP but it is currently not known if it is NP-complete.

Interval Trees We refer to Cormen et al. [2009, Section 14.3] for another exposition on interval trees and their applications. Interval trees, together with the application of finding rectangle intersection, have been formalized by Zhan [2018].
Abstract Data Types

Tobias Nipkow

In the previous chapter we looked at a very specific example of an abstract data type, namely sets. In this chapter we consider abstract data types in general and in particular the model-oriented approach to the specification of abstract data types. This will lead to a generic format for such specifications. As a second example we consider the abstract data type of maps.

6.1 Abstract Data Types

Abstract data types (ADTs) can be summarized by the following slogan:

\[ \text{ADT} = \text{interface} + \text{specification} \]

where the interface lists the operations supported by the ADT and the specification describes the behaviour of these operations. For example, our set ADT has the following interface:

- empty :: 's
- insert :: 'a ⇒ 's ⇒ 's
- delete :: 'a ⇒ 's ⇒ 's
- isin :: 's ⇒ 'a ⇒ bool

The purpose of an ADT is to be able to write applications based on this ADT that will work with any implementation of the ADT. To this end one can prove properties of the application that are solely based on the specification of the ADT. That is, one can write generic algorithms and prove generic correctness theorems about them in the context of the ADT specification.

6.2 Model-Oriented Specification

We follow the model-oriented style of specification advocated by Jones [1990]. In that style, an abstract type is specified by giving an abstract model for it. For simplicity we assume that each ADT describes one type of interest \( T \). In the set interface \( T \) is \( 's \). This type \( T \) must be specified by some existing HOL type \( A \), the abstract model. In the case of sets this is straightforward: the model for sets is simply the HOL type \( 'a \ set \). The motto is that \( T \) should behave like \( A \). In order to bridge the gap between the two types, the specification needs an
• abstraction function  \( \alpha :: T \Rightarrow A \)

that maps concrete values to their abstract counterparts. Moreover, in general only some elements of \( T \) represent elements of \( A \). For example, in the set implementation in the previous chapter not all trees but only BSTs represent sets. Thus the specification should also take into account an

• invariant  \( \text{invar} :: T \Rightarrow \text{bool} \)

Note that the abstraction function and the invariant are not part of the interface, but they are essential for specification and verification purposes.

As an example, the ADT of sets is shown in Figure 6.1 with suggestive keywords and a fixed mnemonic naming schema for the labels in the specification. This is

**ADT Set** =

**interface**

\( \text{empty} :: \text{'s} \)

\( \text{insert} :: \text{'a} \Rightarrow \text{'s} \Rightarrow \text{'s} \)

\( \text{delete} :: \text{'a} \Rightarrow \text{'s} \Rightarrow \text{'s} \)

\( \text{isin} :: \text{'s} \Rightarrow \text{'a} \Rightarrow \text{bool} \)

**abstraction**  \( \text{set} :: \text{'s} \Rightarrow \text{'a set} \)

**invariant**  \( \text{invar} :: \text{'s} \Rightarrow \text{bool} \)

**specification**

\( \text{set empty} = \{\} \)

\( \text{invar empty} \)

\( \text{invar s} \rightarrow \text{set(insert x s)} = \text{set s} \cup \{x\} \)

\( \text{invar s} \rightarrow \text{invar(insert x s)} \)

\( \text{invar s} \rightarrow \text{set(delete x s)} = \text{set s} - \{x\} \)

\( \text{invar s} \rightarrow \text{invar(delete x s)} \)

\( \text{invar s} \rightarrow \text{isin s x} = (x \in \text{set s}) \)

**Figure 6.1**

ADT Set

the template for ADTs that we follow throughout the book. We have intentionally refrained from showing the Isabelle formalization using so-called locales and have opted for a more intuitive textual format that is not Isabelle-specific, in accordance with the general philosophy of this book. The actual Isabelle text can of course be found in the source files, and locales are explained in a dedicated manual [Ballarin].

We conclude this section by explaining what the specification of an arbitrary ADT looks like. We assume that for each function \( f \) of the interface there is a corresponding
function $f_A$ in the abstract model $A$. For a uniform treatment we extend $\alpha$ and $\text{invar}$ to arbitrary types by setting $\alpha x = x$ and $\text{invar} x = \text{True}$ for all types other than $T$. Each function $f$ of the interface gives rise to two properties in the specification: preservation of the invariant and simulation of $f_A$. The precondition is shared:

\[
\begin{align*}
\text{invar } x_1 \land \ldots \land \text{invar } x_n & \rightarrow \\
\text{invar}(f x_1 \ldots x_n) & \quad \text{(f-inv)} \\
\alpha(f x_1 \ldots x_n) &= f_A(\alpha x_1) \ldots (\alpha x_n) & \quad \text{(f)}
\end{align*}
\]

To understand how the specification of ADT Set is the result of this uniform schema one has to take two things into account:

- Precisely which abstract operations on type 'a set model the functions in the interface of the ADT Set? This correspondence is implicit in the specification: empty is modeled by $\{\}$, insert is modeled by $\lambda x. s \cup \{x\}$, delete is modeled by $\lambda x. s - \{x\}$ and isin is modeled by $\lambda s. x \in s$.
- Because of the artificial extension of $\alpha$ and $\text{invar}$ the above uniform format often collapses to something simpler where some $\alpha$'s and $\text{invar}$'s disappear.

6.3 Implementing ADTs

An implementation of an ADT consists of definitions for all the functions in the interface. For the correctness proof, you also need to provide an abstraction function and the invariant. The latter two need not be executable unless they also occur in the interface and the implementation is meant to be executable. Finally you need to prove all propositions in the specification of the ADT, of course replacing the function names in the ADT by their implementations.

For Isabelle users: because ADTs are formalized as locales, an implementation of an ADT is an interpretation of the corresponding locale.

Exercise 6.1. Sets of natural numbers can be implemented as lists of intervals, where an interval is simply a pair of numbers. For example, the set $\{2, 3, 5, 7, 8, 9\}$ can be represented by the list $\{(2, 3), (5, 5), (7, 9)\}$.

\[
\begin{align*}
\text{type_synonym} & \quad \text{interval} = \text{nat} \times \text{nat} \\
\text{type_synonym} & \quad \text{intervals} = \text{interval list}
\end{align*}
\]

Define an abstraction function and invariant

\[
\begin{align*}
\text{set_of} :: \text{intervals} & \Rightarrow \text{nat set} \\
\text{invar} :: \text{intervals} & \Rightarrow \text{bool}
\end{align*}
\]
The invariant should enforce that all intervals are non-empty, they are sorted in ascending order and they do not overlap. Then define two functions for adding and deleting numbers to and from `intervals`:

\[
\begin{align*}
isin & :: \text{intervals} \Rightarrow \text{nat} \Rightarrow \text{bool} \\
addl & :: \text{nat} \Rightarrow \text{intervals} \Rightarrow \text{intervals} \\
dell & :: \text{nat} \Rightarrow \text{intervals} \Rightarrow \text{intervals}
\end{align*}
\]

Show that `[]`, `addl`, `dell`, `isin`, `set_of` and `invar` correctly implement the ADT `Set` by proving all propositions in the specification, suitably renamed, e.g. `invar ivs \rightarrow set_of (addl i ivs) = set_of ivs \cup \{i\}`.

In a second step, define two functions

\[
\begin{align*}
add & :: \text{intervals} \Rightarrow \text{intervals} \Rightarrow \text{intervals} \\
del & :: \text{intervals} \Rightarrow \text{intervals} \Rightarrow \text{intervals}
\end{align*}
\]

for union and difference and prove

\[
\begin{align*}
invar xs \land invar ys & \rightarrow set_of (\text{add } xs \text{ ys}) = set_of xs \cup set_of ys \\
invar xs \land invar ys & \rightarrow set_of (\text{del } xs \text{ ys}) = set_of ys - set_of xs
\end{align*}
\]

and that they preserve the invariant.

Make sure all functions in your implementation terminate as soon as possible. Both `add` and `del` should take time linear in the sum of the lengths of their arguments. They should not simply iterate `addl` and `dell`.

6.4 Maps

An even more versatile type than sets are maps from `\text{\{a\}}` to `\text{\{b\}}`. In fact, sets can be viewed as maps from `\text{\{a\}}` to `\text{\{bool\}}`. Conversely, many data structures for sets also support maps, e.g. BSTs. Although, for simplicity, we mostly focus on sets in this book, maps are used in a few places too.

Just as with sets, there is both an HOL type of maps and an ADT of maps. We start with the former, where \(\rightarrow\) is just nice syntax:

\[
\text{type\_synonym} \quad \text{'a} \rightarrow \text{'b} = \text{'a} \Rightarrow \text{'b option}
\]

These maps can also be viewed as partial functions. We define the following abbreviation:

\[
m(a \mapsto b) \equiv m(a := \text{Some } b)
\]
The ADT \textit{Map} is shown in Figure 6.2. Type \( 'm \) represents the type of maps from \( 'a \) to \( 'b \). The ADT \textit{Map} is very similar to the ADT \textit{Set} except that the abstraction function \textit{lookup} is also part of the interface: it abstracts a map to a function of type \( 'a \rightarrow 'b \). This implies that the equations are between functions of that type. We use the function update notation (Section 1.3) to explain \textit{update} and \textit{delete}: \textit{update} is modeled by \( \lambda m \ a \ b. \ m(a \mapsto b) \) and \textit{delete} by \( \lambda m \ a. \ m(a := \text{None}) \).

\begin{verbatim}
ADT Map = interface
empty :: 'm
update :: 'a \rightarrow 'b \Rightarrow 'm \Rightarrow 'm
delete :: 'a \Rightarrow 'm \Rightarrow 'm
lookup :: 'm \Rightarrow 'a \rightarrow 'b

abstraction lookup

invariant invar :: 'm \Rightarrow bool

specification
lookup empty = (\_ \_ \text{None}) \hspace{1cm} \text{(empty)}

invar empty

invar m \Rightarrow lookup (update a b m) = (lookup m)(a \mapsto b) \hspace{1cm} \text{(update)}

invar m \Rightarrow invar (update a b m) \hspace{1cm} \text{(update-inv)}

invar m \Rightarrow lookup (delete a m) = (lookup m)(a := \text{None}) \hspace{1cm} \text{(delete)}

invar m \Rightarrow invar (delete a m) \hspace{1cm} \text{(delete-inv)}
\end{verbatim}

Figure 6.2 ADT \textit{Map}

\section*{6.5 Implementing Maps by BSTs}

We implement maps as BSTs of type \(( 'a \times 'b \) \) \textit{tree}. The interface functions have the following straightforward implementations, ignoring the trivial \textit{empty}:

\begin{verbatim}
lookup :: ( 'a \times 'b \) \textit{tree} \Rightarrow 'a \rightarrow 'b
lookup () _ = None
lookup (l, (a, b), r) x = (case cmp x a of
LT \Rightarrow lookup l x |
EQ \Rightarrow Some b |
GT \Rightarrow lookup r x)
\end{verbatim}
update :: 'a ⇒ 'b ⇒ ('a × 'b) tree ⇒ ('a × 'b) tree
update x y () = ()
update x y (l, (a, b), r) = (case cmp x a of
    LT ⇒ (update x y l, (a, b), r) |
    EQ ⇒ (l, (x, y), r) |
    GT ⇒ (l, (a, b), update x y r))

delete :: 'a ⇒ ('a × 'b) tree ⇒ ('a × 'b) tree
delete _ () = ()
delete x (l, (a, b), r) = (case cmp x a of
    LT ⇒ (delete x l, (a, b), r) |
    EQ ⇒ if r = () then l |
        else let (ab', r') = split_min r in (l, (x, y), r') |
    GT ⇒ (l, (a, b), delete x r))

Function split_min is the one defined in Section 5.6.4.

The correctness proof proceeds as in Section 5.4. The intermediate level is the type
('a × 'b) list of association lists sorted w.r.t. the fst component:

\[
\text{sorted} \ ps \equiv \text{sorted} \ (\text{map} \ \text{fst} \ ps)
\]

Functions update, delete and lookup are easily implemented:

upd_list :: 'a ⇒ 'b ⇒ ('a × 'b) list ⇒ ('a × 'b) list
upd_list x y [] = [(x, y)]
upd_list x y ((a, b) # ps)
    = (if x < a then (x, y) # (a, b) # ps |
        else if x = a then (x, y) # ps else (a, b) # upd_list x y ps)

del_list :: 'a ⇒ ('a × 'b) list ⇒ ('a × 'b) list
del_list _ [] = []
del_list x ((a, b) # ps) = (if x = a then ps else (a, b) # del_list x ps)

map_of :: ('a × 'b) list ⇒ 'a ⇒ 'b
6.5 Implementing Maps by BSTs

\[
\begin{align*}
\text{map}_{\text{of}} \; [[]] &= (\lambda x.\; \text{None}) \\
\text{map}_{\text{of}} \; ([a, b] \neq ps) &= (\text{map}_{\text{of}} \; ps)(a \mapsto b)
\end{align*}
\]

It is easy to prove that association lists implement maps of type \( \forall a \to b \) via the abstraction function \( \text{map}_{\text{of}} \):

\[
\begin{align*}
\text{map}_{\text{of}} \; (\text{upd\_list} \; x \; y \; ps) &= (\text{map}_{\text{of}} \; ps)(x \mapsto y) \\
\text{sorted} \; ps \rightarrow \text{map}_{\text{of}} \; (\text{del\_list} \; x \; ps) &= (\text{map}_{\text{of}} \; ps)(x \leftarrow \text{None}) \\
\text{sorted} \; ps \rightarrow \text{sorted} \; (\text{upd\_list} \; x \; y \; ps) \\
\text{sorted} \; ps \rightarrow \text{sorted} \; (\text{del\_list} \; x \; ps)
\end{align*}
\]

The correctness of \( \text{map\_of} \) (as an operation on association lists) is trivial because \( \text{map\_of} \) is also the abstraction function and thus the requirement becomes \( \text{map\_of} \; ps \; a = \text{map\_of} \; ps \; a \).

We can also prove that \((\forall a \times b)\) trees implement association lists:

\[
\begin{align*}
\text{sorted} \; (\text{inorder} \; t) \rightarrow \\
\text{inorder} \; (\text{update} \; a \; b \; t) &= \text{upd\_list} \; a \; b \; (\text{inorder} \; t) \\
\text{sorted} \; (\text{inorder} \; t) \rightarrow \text{inorder} \; (\text{delete} \; x \; t) &= \text{del\_list} \; x \; (\text{inorder} \; t) \\
\text{sorted} \; (\text{inorder} \; t) \rightarrow \text{lookup} \; t \; x &= \text{map\_of} \; (\text{inorder} \; t) \; x
\end{align*}
\]

The \textit{Map} specification properties follow by composing the above two sets of implementation properties.

\textbf{Exercise 6.2.} Modify the ADT \textit{Map} as follows. Replace \textit{update} and \textit{delete} by a single function \textit{modify} :: \( \forall a \Rightarrow (\forall b \Rightarrow \text{option}) \Rightarrow (\forall m \Rightarrow \text{option}) \) with the specification that \textit{invar} \( m \) implies

\[
\begin{align*}
\text{lookup} \; (\text{modify} \; a \; f \; m) &= (\text{lookup} \; m)(a := f \; (\text{lookup} \; m \; a)) \\
\text{invar} \; (\text{modify} \; a \; f \; m)
\end{align*}
\]

Define \textit{update} and \textit{delete} with the help of \textit{modify} and prove the \textit{update} and \textit{delete} properties from the original ADT \textit{Map} from these definitions and the specification of \textit{modify}. Conversely, in the context of the original ADT \textit{Map}, define \textit{modify} in terms of \textit{update} and \textit{delete} and prove the above properties.
This is the first in a series of chapters examining balanced search trees where the height of the tree is logarithmic in its size and which can therefore be searched in logarithmic time.

The most popular first example of balanced search trees are red-black trees. We start with 2-3 trees, where nodes can have 2 or 3 children, because red-black trees are best understood as an implementation of (a variant of) 2-3 trees. We introduce red-black trees in the next chapter. The type of 2-3 trees is similar to binary trees but with an additional constructor \texttt{Node3}:

\begin{verbatim}
datatype 'a tree23 =
  Leaf |
  Node2 ('a tree23) 'a ('a tree23) |
  Node3 ('a tree23) 'a ('a tree23) 'a ('a tree23)
\end{verbatim}

The familiar syntactic sugar is sprinkled on top:

\begin{verbatim}
⟨⟩ ≡ Leaf 
⟨l, a, r⟩ ≡ Node2 l a r 
⟨l, a, m, b, r⟩ ≡ Node3 l a m b r
\end{verbatim}

The size, height and the completeness of a 2-3 tree are defined by adding an equation for \texttt{Node3} to the corresponding definitions on binary trees:

\begin{verbatim}
|⟨l, _, m, _, r⟩| = |l| + |m| + |r| + 1

h ⟨l, _, m, _, r⟩ = max (h l) (max (h m) (h r)) + 1

complete ⟨l, _, m, _, r⟩ = (h l = h m ∧ h m = h r ∧ complete l ∧ complete m ∧ complete r)
\end{verbatim}
A trivial induction yields complete \( t \longrightarrow 2^h t \leq |t| + 1 \): thus all operations on complete 2-3 trees have logarithmic complexity if they descend along a single branch and take constant time per node. This is the case and we will not discuss complexity in any more detail.

A nice property of 2-3 trees is that for every \( n \) there is a complete 2-3 tree of size \( n \). As we will see below, completeness can be maintained under insertion and deletion in logarithmic time.

**Exercise 7.1.** Define a function \( \text{maxt} :: \text{nat} \Rightarrow \text{unit} \Rightarrow \text{tree23} \) that creates the tree with the largest number of nodes given the height of the tree. We use type \( \text{unit} \) because we are not interested in the elements in the tree. Prove \( |\text{maxt } n| = (3^n - 1) \text{ div } 2 \) and that no tree of the given height can be larger: \( |t| \leq (3^h t - 1) \text{ div } 2 \). Note that both subtraction and division on type \( \text{nat} \) can be tedious to work with. You may want to prove the two properties as corollaries of subtraction- and division-free properties. Alternatively, work with \( \text{real} \) instead of \( \text{nat} \) by replacing \( \text{div} \) by \( \text{/} \).

### 7.1 Implementation of ADT Set

The implementation will maintain the usual ordering invariant and additionally completeness. When we speak of a 2-3 tree we will implicitly assume these two invariants now.

Searching a 2-3 tree is like searching a binary tree (see Section 5.2) but with one more defining equation:

\[
\text{isin } (l, a, m, b, r) \ x = \begin{cases} 
\text{isin } l \ x & \text{if } \text{cmp } x \ a = \text{LT} \\
\text{isin } m \ x & \text{if } \text{cmp } x \ b = \text{EQ} \\
\text{isin } r \ x & \text{if } \text{cmp } x \ b = \text{GT}
\end{cases}
\]

Insertion into a 2-3 tree must preserve the completeness invariant. Thus recursive calls must report back to the caller if the child has “overflown”, i.e. increased in height. Therefore insertion returns a result of type \('a upI\):

\[
\text{datatype } 'a upI = \begin{cases} 
\text{TI } ('a \text{ tree23}) & \text{if } t \text{ and } t' \text{ should have the same height,} \\
\text{OF } l \ x \ r & \text{if } t \text{ and } l \text{ and } r \text{ should have the same height.}
\end{cases}
\]

This is the idea: If insertion into \( t \) returns \( \text{TI } t' \) then \( t \) and \( t' \) should have the same height,

\( \text{OF } l \ x \ r \) then \( t \) and \( l \) and \( r \) should have the same height.

The insertion functions are shown in Figure 7.1. The actual work is performed by the recursive function \( \text{ins} \). The element to be inserted is propagated down to a leaf, which causes an overflow of the leaf. If an overflow is returned from a recursive call it
7.1 Implementation of ADT Set

\[ \text{insert } x \ t = \text{treeI} (\text{ins } x \ t) \]

\[
\text{ins} :: 'a \Rightarrow 'a \text{tree23} \Rightarrow 'a \text{upI}
\]

\[
\text{ins } \langle \rangle = \text{OF} \langle \rangle \ x \langle \rangle
\]

\[
\text{ins } \langle l, a, r \rangle = (\text{case } \text{cmp } x \ a \ if
\]

\[
\begin{align*}
LT \Rightarrow & \quad \text{case } \text{ins } x \ l \ of \\
& \quad TI \ l' \Rightarrow TI \langle l', a; r \rangle \\
& \quad \text{OF } l_1 \ b \ l_2 \Rightarrow TI \langle l_1, b; l_2, a; r \rangle \\
EQ \Rightarrow & \quad TI \langle l, a; r \rangle \\
GT \Rightarrow & \quad \text{case } \text{ins } x \ r \ of \\
& \quad TI \ r' \Rightarrow TI \langle l, a; r' \rangle \\
& \quad \text{OF } l_1 \ b \ l_2 \Rightarrow TI \langle l, a; l_1, b; l_2, r \rangle
\end{align*}
\]

\[
\text{ins } \langle l, a, m, b, r \rangle
\]

\[
= (\text{case } \text{cmp } x \ a \ of
\]

\[
\begin{align*}
LT \Rightarrow & \quad \text{case } \text{ins } x \ l \ of \\
& \quad TI \ l' \Rightarrow TI \langle l', a; m, b; r \rangle \\
& \quad \text{OF } l_1 \ c \ l_2 \Rightarrow \text{OF } \langle l_1, c; l_2 \rangle \ a \langle m, b; r \rangle \\
EQ \Rightarrow & \quad TI \langle l, a; m, b; r \rangle \\
GT \Rightarrow & \quad \text{case } \text{cmp } x \ b \ of \\
& \quad LT \Rightarrow \text{case } \text{ins } x \ m \ of \\
& \quad TI \ m' \Rightarrow TI \langle l, a; m', b; r \rangle \\
& \quad \text{OF } m_1 \ c \ m_2 \Rightarrow \text{OF } \langle l, a; m_1 \rangle \ c \langle m_2, b; r \rangle \\
EQ \Rightarrow & \quad TI \langle l, a; m, b; r \rangle \\
GT \Rightarrow & \quad \text{case } \text{ins } x \ r \ of \\
& \quad TI \ r' \Rightarrow TI \langle l, a; m, b; r' \rangle \\
& \quad \text{OF } l_1 \ c \ l_2 \Rightarrow \text{OF } \langle l, a; m \rangle \ b \langle l_1, c; l_2 \rangle
\end{align*}
\]

**Figure 7.1** Insertion into 2-3 tree

can be absorbed into a Node2 but in a Node3 it causes another overflow. At the root of the tree, function `treeI` converts values of type `\text{upI}` back into trees:

\[
\text{treeI} :: \text{'a upI} \Rightarrow 'a \text{tree23}
\]

\[
\text{treeI} (\text{TI } t) = t
\]

\[
\text{treeI} (\text{OF } l \ a \ r) = \langle l, a, r \rangle
\]
Deletion is dual. Recursive calls must report back to the caller if the child has "underflown", i.e. decreased in height. Therefore deletion returns a result of type \( \text{upD} \):

\[
\text{datatype } 'a \text{ upD} = TD (\ 'a \text{ tree23} ) \mid UF (\ 'a \text{ tree23} )
\]

This is the idea: If deletion from \( t \) returns

\( TD t' \) then \( t \) and \( t' \) should have the same height,

\( UF t' \) then \( t \) should be one level higher than \( t' \).

The main deletion functions are shown in Figure 7.2. The actual work is performed by the recursive function \( \text{del} \). If the element to be deleted is in a child, the result of a recursive call is reintegrated into the node via the auxiliary functions \( \text{nodeij} \) from Figure 7.3: \( \text{nodeij} \) creates a node with \( i \) children, where child \( j \) is given as an \( \text{upD} \) value, and wraps the node up in \( UF \) or \( TD \), depending on whether an underflow occurred or not. If the element to be deleted is in the node itself, a replacement is obtained and deleted from a child via \( \text{split_min} \). At the root of the tree, \( \text{upD} \) values are converted back into trees:

\[
\text{treeD} :: 'a \text{ upD} \Rightarrow 'a \text{ tree23}
\]

\[
\text{treeD} (\ TD t) = t
\]

\[
\text{treeD} (\ UF t) = t
\]

### 7.2 Preservation of Completeness

As explained in Section 5.4, we do not go into the automatic functional correctness proofs but concentrate on invariant preservation. To express the relationship between the height of a tree before and after insertion we define a height function \( hI \):

\[
hI :: 'a \text{ upI} \Rightarrow \text{nat}
\]

\[
hI (\ TI t) = h t
\]

\[
hI (\ OF l \_\_ ) = h l
\]

Intuitively, \( hI \) is the height of the tree before insertion. A routine induction proves

\[
\text{complete } t \rightarrow \text{complete } (\text{treeI} (\text{ins } a \ t)) \land hI (\text{ins } a \ t) = h t
\]

which implies by definition that

\[
\text{complete } t \rightarrow \text{complete } (\text{insert } a \ t)
\]
7.2 Preservation of Completeness

**Figure 7.2** Deletion from 2-3 tree: main functions
Figure 7.3  Deletion from 2-3 tree: auxiliary functions

node21 :: 'a upD ⇒ 'a ⇒ 'a tree23 ⇒ 'a upD
node21 (TD t₁) a t₂ = TD ⟨t₁, a, t₂⟩
node21 (UF t₁) a ⟨t₂, b, t₃⟩ = UF ⟨t₁, a, t₂, b, t₃⟩
node21 (UF t₁) a ⟨t₂, b, t₃, c, t₄⟩ = TD ⟨⟨t₁, a, t₂⟩, b, ⟨t₃, c, t₄⟩⟩

dnode22 :: 'a tree23 ⇒ 'a ⇒ 'a upD ⇒ 'a upD
node22 t₁ a (TD t₂) = TD ⟨t₁, a, t₂⟩
node22 ⟨t₁, b, t₂⟩ a (UF t₃) = UF ⟨t₁, b, t₂, a, t₃⟩
node22 ⟨t₁, b, t₂, c, t₃⟩ a (UF t₄) = TD ⟨⟨t₁, b, t₂⟩, c, ⟨t₃, a, t₄⟩⟩

dnode31 :: 'a upD ⇒ 'a ⇒ 'a tree23 ⇒ 'a ⇒ 'a tree23 ⇒ 'a upD
node31 (TD t₁) a t₂ b t₃ = TD ⟨t₁, a, t₂, b, t₃⟩
node31 (UF t₁) a ⟨t₂, b, t₃⟩ c t₄ = TD ⟨⟨t₁, a, t₂, b, t₃⟩, c, t₄⟩
node31 (UF t₁) a ⟨t₂, b, t₃, c, t₄⟩ d t₅
= TD ⟨⟨t₁, a, t₂⟩, b, ⟨t₃, c, t₄⟩, d, t₅⟩

dnode32 :: 'a tree23 ⇒ 'a ⇒ 'a upD ⇒ 'a ⇒ 'a tree23 ⇒ 'a upD
node32 t₁ a (TD t₂) b t₃ = TD ⟨t₁, a, t₂, b, t₃⟩
node32 t₁ a (UF t₂) b ⟨t₃, c, t₄⟩ = TD ⟨t₁, a, ⟨t₂, b, t₃⟩, c, t₄⟩
node32 t₁ a (UF t₂) b ⟨t₃, c, t₄⟩ d t₅
= TD ⟨t₁, a, ⟨t₂, b, t₃⟩, c, ⟨t₄, d, t₅⟩⟩

dnode33 :: 'a tree23 ⇒ 'a ⇒ 'a tree23 ⇒ 'a ⇒ 'a upD ⇒ 'a upD
node33 t₁ a t₂ b (TD t₃) = TD ⟨t₁, a, t₂, b, t₃⟩
node33 t₁ a ⟨t₂, b, t₃⟩ c (UF t₄) = TD ⟨t₁, a, ⟨t₂, b, t₃⟩, c, t₄⟩
node33 t₁ a ⟨t₂, b, t₃, c, t₄⟩ d (UF t₅)
= TD ⟨t₁, a, ⟨t₂, b, t₃⟩, c, ⟨t₄, d, t₅⟩⟩
To express the relationship between the height of a tree before and after deletion, we define:

\[
\begin{align*}
    h_D &: \text{a upD } \Rightarrow \text{nat} \\
    h_D (TD t) &= h t \\
    h_D (UF t) &= h t + 1
\end{align*}
\]

The intuition is that \( h_D \) is the height of the tree before deletion.

We now list a sequence of simple inductive properties that build on each other and culminate in completeness preservation of \textit{delete}:

- \textit{complete} \( r \land \text{complete (treeD l')} \land h r = hD l' \rightarrow \text{complete (treeD (node21 l' a r))} \)
- \( 0 < h r \rightarrow hD (node21 l' a r) = \max (hD l') (h r) + 1 \)
- \textit{split_min} \( t = (x, t') \land 0 < h t \land \text{complete t} \rightarrow hD t' = h t \)
- \textit{split_min} \( t = (x, t') \land \text{complete t} \land 0 < h t \rightarrow \text{complete (treeD t')} \)
- \textit{complete} \( t \rightarrow hD (del x t) = h t \)
- \textit{complete} \( t \rightarrow \text{complete (treeD (del x t))} \)
- \textit{complete} \( t \rightarrow \text{complete (delete x t)} \)

For each property of \textit{node21} there are analogous properties for the other \textit{nodeij} functions which we omit.

### 7.3 Converting a List into a 2-3 Tree

We consider the problem of converting a list of elements into a 2-3 tree. If the resulting tree should be a search tree, there is the obvious approach: insert the elements one by one starting from the empty tree. This takes time \( \Theta(n \lg n) \). This holds for any data structure where insertion takes time proportional to \( \lg n \). In that case inserting \( n \) elements one by one takes time proportional to \( \lg 1 + \cdots + \lg n = \lg(n!) \). Now \( n! \leq n^n \) implies \( \lg(n!) \leq n \lg n \). On the other hand, \( n^n \leq (n \cdot 1) \cdot (n \cdot 1 - 2) \cdot \cdots \cdot (1 \cdot n) = (n!)^2 \) implies \( \frac{1}{2} n \lg n \leq \lg(n!) \). Thus \( \lg(n!) \in \Theta(n \lg n) \) (which also follows from Stirling’s formula). We have intentionally proved a \( \Theta \) property because the \( O \) property is obvious but one might hope that \( \lg 1 + \cdots + \lg n \) has a lower order of growth than \( n \lg n \). However, since a search tree can be converted into a sorted list in linear time, the conversion into the search tree cannot be faster than sorting.

Now we turn to the actual topic of this section: converting a list \( xs \) into a 2-3 tree \( t \) such that \( inorder t = xs \) — in linear time. Thus we can take advantage of situations where we already know that \( xs \) is sorted. The bottom-up conversion algorithm is
particularly intuitive. It repeatedly passes over an alternating list $t_1, e_1, t_2, e_2, \ldots, t_k$ of trees and elements, combining trees and elements into new trees. Given elements $a_1, \ldots, a_n$ we start with the alternating list $\langle \rangle, a_1, \langle \rangle, a_2, \ldots, a_n, \langle \rangle$. On every pass over this list, we replace adjacent triples $t, a, t'$ by $(t, a, t')$, possibly creating a 3-node instead of a 2-node at the end of the list. Once a single tree is left over, we terminate.

We define this type of alternating (and non-empty) list as a new data type:

```plaintext
datatype 'a tree23s = T ('a tree23) | TTs ('a tree23) 'a ('a tree23s)
```

The following examples demonstrate the encoding of alternating lists as terms of type `'a tree23s`:

- Alternating list: $t_1 \ t_1, e_1 ; t_2; t_1, e_1; t_2; e_2; ts$
- Encoding: $T \ t_1 \ TTs \ t_1 \ e_1 (T \ t_2) \ TTs \ t_1 \ e_1 (TTs \ t_2 \ e_2 \ ts)$

We also need the following auxiliary functions:

```plaintext
len :: 'a tree23s ⇒ nat
len (T _) = 1
len (TTs _ _ ts) = len ts + 1

trees :: 'a tree23s ⇒ 'a tree23 set
trees (T t) = {t}
trees (TTs t _ ts) = {t} ∪ trees ts

inorder2 :: 'a tree23s ⇒ 'a list
inorder2 (T t) = inorder t
inorder2 (TTs t a ts) = inorder t @ a ≠ inorder2 ts

join_all :: 'a tree23s ⇒ 'a tree23
join_all (T t) = t
join_all ts = join_all (join_adj ts)

join_adj :: 'a tree23s ⇒ 'a tree23s
join_adj (TTs t1 a (T t2)) = T (t1, a, t2)
```

Repeatedly passing over the alternating list until only a single tree remains is expressed by the following functions:
7.3 Converting a List into a 2-3 Tree

join_adj (TTs t₁ a (TTs t₂ b (T₃))) = T₁ (a, t₂, b, t₃)
join_adj (TTs t₁ a (TTs t₂ b ts)) = TTs (t₁, a, t₂) b (join_adj ts)

Note that join_adj is not and does not need to be defined on single trees. We express this precondition with an abbreviation:
not_T ts ≡ ∀t. ts ≠ T t

Also note that join_all terminates only because join_adj shortens the list:
not_T ts → len (join_adj ts) < len ts

In fact, it reduces the length at least by a factor of 2:
not_T ts → len (join_adj ts) ≤ len ts div 2 \hspace{1cm} (7.1)

The whole process starts with a list of alternating leaves and elements:

```
tree23_of_list :: 'a list ⇒ 'a tree23
tree23_of_list as = join_all (leaves as)

leaves :: 'a list ⇒ 'a tree23s
leaves [] = T ()
leaves (a # as) = TTs () a (leaves as)
```

7.3.1 Functional Correctness

Functional correctness is easily established. The inorder and the completeness properties are proved independently by the following inductive lemmas:

not_T ts → inorder2 (join_adj ts) = inorder2 ts
inorder (join_all ts) = inorder2 ts
inorder (tree23_of_list as) = as

(∀t ∈ trees ts. complete t ∧ h t = n) ∧ not_T ts →
(∀t ∈ trees (join_adj ts). complete t ∧ h t = n + 1)

(∀t ∈ trees ts. complete t ∧ h t = n) → complete (join_all ts)
t ∈ trees (leaves as) → complete t ∧ h t = 0
complete (tree23_of_list as)
7.3.2 Running Time Analysis

Why does the conversion take linear time? Because the first pass over an alternating list of length \( n \) takes \( n \) steps, the next pass \( n/2 \) steps, the next pass \( n/4 \) steps, etc., and this sums up to \( 2n \). The time functions for the formal proof are shown in Appendix B.3. The following upper bound is easily proved by induction on the computation of \( \text{join}\_\text{adj} \):\

\[
\text{not}\_T\ ts \rightarrow T_{\text{join}\_\text{adj}}\ ts \leq \text{len}\ ts\ \text{div} 2
\]  

(7.2)

An upper bound \( T_{\text{join}\_\text{all}}\ ts \leq 2 \cdot \text{len}\ ts \) follows by induction on the computation of \( \text{join}\_\text{adj} \). We focus on the induction step:

\[
T_{\text{join}\_\text{all}}\ ts \\
= T_{\text{join}\_\text{adj}}\ ts + T_{\text{join}\_\text{all}}\ (\text{join}\_\text{adj}\ ts) + 1 \\
\leq \text{len}\ ts\ \text{div} 2 + 2 \cdot \text{len}\ (\text{join}\_\text{adj}\ ts) + 1 \\
\leq \text{len}\ ts\ \text{div} 2 + 2 \cdot (\text{len}\ ts\ \text{div} 2) + 1 \\
\leq 2 \cdot \text{len}\ ts
\] 

using (7.2) and IH

by (7.1)

because \( 1 \leq \text{len}\ ts \)

Now it is routine to derive

\[
T_{\text{tree}23\_\text{of}\_\text{list}}\ as \leq 3 \cdot |\text{as}| + 3
\]

7.4 Chapter Notes

The invention of 2-3 trees is credited to Hopcroft in 1970 by Cormen et al. [2009, p. 337]. Equational definitions were given by Hoffmann and O’Donnell [1982] (only insertion) and Reade [1992]. Our formalisation is based on teaching material by Franklyn Turbak and the article by Hinze [2018].
Red-Black Trees

Tobias Nipkow

Red-black trees are a popular implementation technique for BSTs: they guarantee logarithmic height just like 2-3 trees but the code is arguably simpler. The nodes are colored either red or black. Abstractly, red-black trees encode 2-3-4 trees where nodes have between 2 and 4 children. Each 2-3-4 node is encoded by a group of 2, 3 or 4 colored binary nodes as follows:

\[
\langle \rangle \approx \langle \rangle \\
\langle A, a, B \rangle \approx \langle A, a, B \rangle \\
\langle A, a, B, b, C \rangle \approx \langle A, a, B, b, C \rangle \text{ or } \langle A, a, B, b, C \rangle \\
\langle A, a, B, b, C, c, D \rangle \approx \langle A, a, B, b, C, c, D \rangle
\]

Color expresses grouping: a black node is the root of a 2-3-4 node, a red node is part of a bigger 2-3-4 node. Thus a red-black tree needs to satisfy the following properties or invariants:

1. The root is black.
2. Every \( \langle \rangle \) is considered black.
3. If a node is red, its children are black.
4. All paths from a node to a leaf have the same number of black nodes.

The final property expresses that the corresponding 2-3-4 tree is complete. The last two properties imply that the tree has logarithmic height (see below).

We implement red-black trees as binary trees augmented (see Section 4.4) with a color tag:

```haskell
datatype color = Red | Black

type_synonym 'a rbt = ('a × color) tree
```

Some new syntactic sugar is sprinkled on top:
Chapter 8  Red-Black Trees

\[ R \ l \ a \ r \equiv (l, (a, \text{Red}), r) \]
\[ B \ l \ a \ r \equiv (l, (a, \text{Black}), r) \]

The following functions get and set the color of a node:

\[
\begin{align*}
\text{color} \colon & \text{`a rbt} \Rightarrow \text{color} \\
\text{color} & (.()) = \text{Black} \\
\text{color} & (.(), (_, c), _) = c
\end{align*}
\]
\[
\begin{align*}
\text{paint} \colon & \text{color} \Rightarrow \text{`a rbt} \Rightarrow \text{`a rbt} \\
\text{paint} & (.()) = () \\
\text{paint} & c (.(), (a, _), r) = (.(), (a, c), r)
\end{align*}
\]

Note that the \text{color} of a leaf is by definition black.

8.1 Invariants

The above informal description of the red-black tree invariants is formalized as the predicate \text{rbt} which (for reasons of modularity) is split into a color and a height invariant \text{invc} and \text{invh}:

\[
\begin{align*}
\text{rbt} \colon & \text{`a rbt} \Rightarrow \text{bool} \\
\text{rbt} & t = \left( \text{invc} \ t \land \text{invh} \ t \land \text{color} \ t = \text{Black} \right)
\end{align*}
\]

The color invariant expresses that red nodes must have black children:

\[
\begin{align*}
\text{invc} \colon & \text{`a rbt} \Rightarrow \text{bool} \\
\text{invc} & (.()) = \text{True} \\
\text{invc} & (.(), (_, c), r) \\
& = ((c = \text{Red} \rightarrow \text{color} \ l = \text{Black} \land \text{color} \ r = \text{Black}) \land \\
& \text{invc} \ l \land \text{invc} \ r)
\end{align*}
\]

The height invariant expresses (via the \text{black height} \text{bh}) that all paths from the root to a leaf have the same number of black nodes:
8.2 Implementation of ADT Set

\[ \text{invh} :: 'a \text{ rbt} \Rightarrow \text{bool} \]
\[ \text{invh} \emptyset = \text{True} \]
\[ \text{invh} \langle l, (_ , _), r \rangle = (\text{bh} \ l = \text{bh} \ r \land \text{invh} \ l \land \text{invh} \ r) \]

\[ \text{bh} :: 'a \text{ rbt} \Rightarrow \text{nat} \]
\[ \text{bh} \emptyset = 0 \]
\[ \text{bh} \langle l, (_ , c), _ \rangle = (\text{if} \ c = \text{Black} \text{ then} \ \text{bh} \ l + 1 \text{ else} \ \text{bh} \ l) \]

Note that although \( \text{bh} \) traverses only the left spine of the tree, the fact that \( \text{invh} \) traverses the complete tree ensures that all paths from the root to a leaf are considered. (See Exercise 8.2)

The split of the invariant into \( \text{invc} \) and \( \text{invh} \) improves modularity: frequently one can prove preservation of \( \text{invc} \) and \( \text{invh} \) separately, which facilitates proof search. For compactness we will mostly present the combined invariance properties.

8.1.1 Logarithmic Height
In a red-black tree, i.e. \( \text{rbt} \ t \), every path from the root to a leaf has the same number of black nodes, and no such path has two red nodes in a row. Thus each leaf is at most twice as deep as any other leaf, and therefore \( h \ t \leq 2 \cdot \lg |t|_1 \). The detailed proof starts with the key inductive relationship between height and black height

\[ \text{invc} \ t \land \text{invh} \ t \rightarrow h \ t \leq 2 \cdot \text{bh} \ t + (\text{if} \ \text{color} \ t = \text{Black} \text{ then} \ 0 \text{ else} \ 1) \]

which has the easy corollary \( \text{rbt} \ t \rightarrow h \ t / 2 \leq \text{bh} \ t \). Together with the easy inductive fact

\[ \text{invc} \ t \land \text{invh} \ t \rightarrow 2^{\text{bh} \ t} \leq |t|_1 \]

this implies \( 2^{h \ t / 2} \leq 2^{\text{bh} \ t} \leq |t|_1 \) and thus \( h \ t \leq 2 \cdot \lg |t|_1 \) if \( \text{rbt} \ t \).

8.2 Implementation of ADT Set
We implement sets by red-black trees that are also BSTs. As usual, we only discuss the proofs of preservation of the \( \text{rbt} \) invariant.

Function \( \text{isin} \) is implemented as for all augmented BSTs (see Section 5.6.1).

8.2.1 Insertion
Insertion is shown in Figure 8.1. The workhorse is function \( \text{ins} \). It descends to the leaf where the element is inserted and it adjusts the colors on the way back up. The adjustment is performed by \( \text{baliL/baliR} \). They combine arguments \( l \ a \ r \) into a tree. If
there is a red-red conflict in l/r, they rebalance and replace it by red-black. Inserting into a red node needs no immediate balancing because that will happen at the black node above it:

\[
\text{ins } 1 \ (B \ (R \ 0) \ 2 \ (R \ 3)) = baliL \ (\text{ins} \ 1 \ (R \ 0) \ 2 \ (R \ 3)) = baliL \ (R \ 0 \ (\text{ins} \ 1) \ 2 \ (R \ 3)) = baliL \ (R \ 0 \ (R \ 1) \ 2 \ (R \ 3)) = R \ (B \ 0 \ 1 \ (B \ 2 \ (R \ 3)))
\]

Passing a red node up means an overflow occurred (as in 2-3 trees) that needs to be dealt with further up. At the latest, \text{insert} turns red into black at the very top.
Function \texttt{ins} preserves \texttt{invh} but not \texttt{invc}: it may return a tree with a red-red conflict at the root, as in the example above: \texttt{ins} 1 (\texttt{R} \texttt{L} 0 \texttt{L}) = \texttt{R} \texttt{L} 0 (\texttt{R} \texttt{L} 1 \texttt{L}). However, once the root node is colored black, everything is fine again. Thus we introduce the weaker invariant \texttt{invc2}:

\[
invc2 \equiv \texttt{invc} (\texttt{paint Black} \ t)
\]

It is easy to prove that \texttt{baliL} and \texttt{baliR} preserve \texttt{invh} and upgrade from \texttt{invc2} to \texttt{invc}:

\[
\begin{align*}
\texttt{invc} \land \texttt{invc} \land \texttt{invc2} \land \texttt{invc} \land \texttt{bh} \ l = \texttt{bh} \ r & \rightarrow \\
\texttt{invc} (\texttt{baliL} \ l \ a \ r) \land \texttt{invc} (\texttt{baliL} \ l \ a \ r) \land \texttt{bh} (\texttt{baliL} \ l \ a \ r) & = \texttt{bh} \ l + 1 \\
\texttt{invc} \land \texttt{invc} \land \texttt{invc2} \land \texttt{invc} \land \texttt{bh} \ l = \texttt{bh} \ r & \rightarrow \\
\texttt{invc} (\texttt{baliR} \ l \ a \ r) \land \texttt{invc} (\texttt{baliR} \ l \ a \ r) \land \texttt{bh} (\texttt{baliR} \ l \ a \ r) & = \texttt{bh} \ l + 1
\end{align*}
\]

Another easy induction yields

\[
\begin{align*}
\texttt{invc} \land \texttt{invc} & \rightarrow \\
\texttt{invc2} (\texttt{ins} \ x \ t) \land (\texttt{color} \ t = \texttt{Black} \rightarrow \texttt{invc} (\texttt{ins} \ x \ t)) & \land \\
\texttt{invc} (\texttt{ins} \ x \ t) \land \texttt{bh} (\texttt{ins} \ x \ t) & = \texttt{bh} \ t
\end{align*}
\]

The corollary \texttt{rbt} \rightarrow \texttt{rbt} (\texttt{insert} \ x \ t) is immediate.

### 8.2.2 Deletion

Deletion from a red-black tree is shown in Figure 8.2. It follows the deletion-by-replacing approach (Section 5.2.1). The tricky bit is how to maintain the invariants. As before, intermediate trees may only satisfy the weaker invariant \texttt{invc2}. Functions \texttt{del} and \texttt{split_min} decrease the black height of a tree with a black root node and leave the black height unchanged otherwise. To see that this makes sense, consider deletion from a singleton black or red node. The case that the element to be removed is not in the black tree can be dealt with by coloring the root node red. These are the precise input/output relations:

**Lemma 8.1.** \texttt{split_min} \( t = (x, t') \land t \neq \bot \land \texttt{invc} \land \texttt{invc} \rightarrow \\
\texttt{invc} t' \land (\texttt{color} t = \texttt{Red} \rightarrow \texttt{bh} t' = \texttt{bh} t \land \texttt{invc} t') \land \\
(\texttt{color} t = \texttt{Black} \rightarrow \texttt{bh} t' = \texttt{bh} t - 1 \land \texttt{invc2} t')

**Lemma 8.2.** \texttt{invc} \land \texttt{invc} \land t' = \texttt{del} \ x \ t \rightarrow \\
\texttt{invc} t' \land (\texttt{color} t = \texttt{Red} \rightarrow \texttt{bh} t' = \texttt{bh} t \land \texttt{invc} t') \land \\
(\texttt{color} t = \texttt{Black} \rightarrow \texttt{bh} t' = \texttt{bh} t - 1 \land \texttt{invc2} t')

It is easy to see that the \texttt{del}-Lemma implies correctness of \texttt{delete}:

**Corollary 8.3.** \texttt{rbt} \rightarrow \texttt{rbt} (\texttt{delete} \ x \ t)
delete x t = paint Black (del x t)

del :: 'a ⇒ 'a rbt ⇒ 'a rbt
del _ () = ()
del x (l, (a, _), r)
= (case cmp x a of
  LT ⇒ let l' = del x l
      in if l /= () ∧ color l = Black then baldL l' a r else R l' a r |
  EQ ⇒ let r = () then l
      else let (a', r') = split_min r
      in if color r = Black then baldR l a' r' else R l a' r' |
  GT ⇒ let r' = del x r
      in if r /= () ∧ color r = Black then baldR l a r' else R l a r')

split_min :: 'a rbt ⇒ 'a × 'a rbt
split_min (l, (a, _), r)
= (if l = () then (a, r)
  else let (x, l') = split_min l
  in (x, if color l = Black then baldL l' a r else R l' a r))

baldL :: 'a rbt ⇒ 'a ⇒ 'a rbt ⇒ 'a rbt
baldL (R t1 a t2) b t3 = R (B t1 a t2) b t3
baldL t1 a (B t2 b t3) = baliR t1 a (R t2 b t3)
baldL t1 a (R (B t2 b t3) c t4) = R (B t1 a t2) b (baliR t3 c (paint Red t4))
baldL t1 a t2 = R t1 a t2

baldR :: 'a rbt ⇒ 'a ⇒ 'a rbt ⇒ 'a rbt
baldR t1 a (R t2 b t3) = R t1 a (B t2 b t3)
baldR (B t1 a t2) b t3 = baliL (R t1 a t2) b t3
baldR (R t1 a t2) c t4 = R (baliL (paint Red t1) a t2) b (B t3 c t4)
baldR t1 a t2 = R t1 a t2

Figure 8.2  Deletion from red-black tree
The proofs of the two preceding lemmas need the following precise characterizations of $\text{baldL}$ and $\text{baldR}$, the counterparts of $\text{baliL}$ and $\text{baliR}$:

**Lemma 8.4.** $\text{invh l} \land \text{invh r} \land \text{bh l} + 1 = \text{bh r} \land \text{invc l} \land \text{invc r} \land \text{t} = \text{baldL l a r} \implies \text{invh t'} \land \text{bh t'} = \text{bh r} \land \text{invc2 t'} \land (\text{color r} = \text{Black} \implies \text{invc t'})$

**Lemma 8.5.** $\text{invh l} \land \text{invh r} \land \text{bh l} = \text{bh r} + 1 \land \text{invc l} \land \text{invc2 r} \land \text{t} = \text{baldR l a r} \implies \text{invh t'} \land \text{bh t'} = \text{bh l} \land \text{invc2 t'} \land (\text{color l} = \text{Black} \implies \text{invc t'})$

The proofs of the two preceding lemmas are by case analyses over the defining equations using the characteristic properties of $\text{baliL}$ and $\text{baliR}$ given above.

**Proof.** Lemma 8.2 is proved by induction on the computation of $\text{del x t}$. The base case is trivial. In the induction step $t = \langle l, (a, c), r \rangle$. If $x < a$ then we distinguish three subcases. If $l = \langle \rangle$ the claim is trivial. Otherwise the claim follows from the IH: if $\text{color l} = \text{Red}$ then the claim follows directly, if $\text{color l} = \text{Black}$ then it follows with the help of Lemma 8.4 (with $l = \text{del x l}$). The case $a < x$ is dual and the case $x = a$ is similar (using Lemma 8.1). We do not show the details because they are tedious but routine.

The proof of Lemma 8.1 is similar but simpler.

### 8.2.3 Deletion by Joining

As an alternative to deletion by replacement we also consider deletion by joining (see Section 5.2.1). The code for red-black trees is shown in Figure 8.3: compared to Figure 8.2, the $\text{EQ}$ case of $\text{del}$ has changed and $\text{join}$ is new.

Invariant preservation is proved much like before except that instead of $\text{split_min}$ we now have $\text{join}$ to take care of. The characteristic lemma is proved by induction on the computation of $\text{join}$:

**Lemma 8.6.** $\text{invh l} \land \text{invh r} \land \text{bh l} = \text{bh r} \land \text{invc l} \land \text{invc r} \land \text{t} = \text{join l r} \implies \text{invh t'} \land \text{bh t'} = \text{bh l} \land \text{invc2 t'} \land (\text{color l} = \text{Black} \land \text{color r} = \text{Black} \implies \text{invc t'})$

### 8.3 Implementation of ADT Map

Maps based on red-black trees are of course very similar to the above sets. In particular we can reuse the balancing and other auxiliary functions because they do not examine the contents of the nodes but only the color. We follow the general approach in Section 6.5. The representing type is $\langle 'a \times 'b \rangle \text{rbt}$.
\begin{verbatim}
del :: 'a ⇒ 'a rbt ⇒ 'a rbt
del _ () = ()
del x (l, (a, _), r)
  = (case cmp x a of
      LT ⇒ if l ≠ () ∧ color l = Black then baldL (del x l) a r
        else R (del x l) a r |
      EQ ⇒ join l r |
      GT ⇒ if r ≠ () ∧ color r = Black then baldR l a (del x r)
        else R l a (del x r))

join :: 'a rbt ⇒ 'a rbt ⇒ 'a rbt
join (l) t = t
join t (l) = t
join (R t1 a t2) (R t3 c t4)
  = (case join t2 t3 of
      R u2 b u3 ⇒ R (R t1 a u2) b (R u3 c t4) |
      t23 ⇒ R t1 a (R t23 c t4)
    )
join (B t1 a t2) (B t3 c t4)
  = (case join t2 t3 of
      R u2 b u3 ⇒ R (B t1 a u2) b (B u3 c t4) |
      t23 ⇒ baldL t1 a (R t23 c t4)
    )
join t1 (R t2 a t3) = R (join t1 t2) a t3 |
join (R t1 a t2) t3 = R t1 a (join t2 t3)
\end{verbatim}

Figure 8.3 Deletion from red-black tree by joining

Function \textit{lookup} is almost identical to its precursor in Section 6.5 except that the \textit{lhs} of the recursive case is \textit{lookup} \((l, ((a, b), _), r)\ x\) because of the (irrelevant) color field. There is no need to show the code.

Function \textit{update} is shown in Figure 8.4. It is a minor variation of insertion shown in Figure 8.1.

Deletion can be implemented by replacing and by joining. (In the source files we have chosen the second option.) In both cases, all we need is to adapt \textit{del} for sets by replacing \textit{cmp} \textit{x} \textit{a} by \textit{cmp} \textit{x} \((\textit{fst} \textit{a})\) (where the second \textit{a} is of type \textit{a} × \textit{b} and should be renamed, e.g. to \textit{ab}). Again, there is no need to show the code.
8.4 Exercises

Exercise 8.1. Show that the logarithmic height of red-black trees is already guaranteed by the color and height invariants:

\[ \text{invc} t \land \text{invh} t \longrightarrow h t \leq 2 \cdot \lg |t| + 2 \]

Exercise 8.2. We already discussed informally why the definition of \( \text{invh} \) captures "all paths from the root to a leaf have the same number of black nodes" although \( \text{bh} \) only traverses the left spine. This exercises formalizes that discussion. The following function computes the set of black heights (number of black nodes) of all paths:

\[ \text{bhs} :: 'a \ rbt \Rightarrow \text{nat set} \]
\[ \text{bhs} \ () = \{0\} \]
\[ \text{bhs} \ (l, (_, c), r) = (\text{let} \ H = \text{bhs} \ l \cup \text{bhs} \ r \in \text{if} \ c = \text{Black} \ then \ Suc \ 'H \ else \ H) \]

where the infix operator \( \cup \) is predefined as \( f \ 'A = \{y \mid \exists x \in A. \ y = f x\} \). Prove \( \text{invh} t \longleftrightarrow \text{bhs} t = \{\text{bh} t\} \). The \( \longleftrightarrow \) direction should be easy, the other direction should need some lemmas.

Exercise 8.3. Following Section 7.3, define a linear time function \( \text{rbt}_\text{of}_\text{list} :: 'a \ \text{list} \Rightarrow 'a \ \text{rbt} \) and prove \( \text{inorder} \ (\text{rbt}_\text{of}_\text{list} \ as) = as \) and \( \text{rbt} \ (\text{rbt}_\text{of}_\text{list} \ as) \).
8.5 Chapter Notes

Red-Black trees were invented by Bayer [1972] who called them “symmetric binary B-trees”. The red-black color convention was introduced by Guibas and Sedgewick [1978] who studied their properties in greater depth. The first functional version of red-black trees (without deletion) is due to Okasaki [1998] and everybody follows his code. A functional version of deletion was first given by Kahrs [2001]\(^1\) and Section 8.2.3 is based on it. Germane and Might [2014] presents a function for deletion by replacement that is quite different from the one in Section 8.2.2. Our starting point was an Isabelle proof by Reiter and Krauss (based on Kahrs). Other verifications of red-black trees are reported by Filliâtre and Letouzey [2004] (using their own deletion function) and Appel [2011] (based on Kahrs).

\(^1\)The code for deletion is not in the article but can be retrieved from this URL: [http://www.cs.ukc.ac.uk/people/staff/smk/redblack/rb.html](http://www.cs.ukc.ac.uk/people/staff/smk/redblack/rb.html)
The AVL tree [Adel'son-Vel'skiï and Landis 1962] (named after its inventors) is the granddaddy of efficient binary search trees. Its logarithmic height is maintained by rotating subtrees based on their height. For efficiency reasons the height of each subtree is stored in its root node. That is, the underlying data structure is a height-augmented tree (see Section 4.4):

\[ \text{type synonym} \ 'a \text{ tree}_{\text{ht}} = ('a \times \text{nat}) \text{ tree} \]

Function \( ht \) extracts the height field and \( \text{node} \) is a smart constructor that sets the height field:

\[
\begin{align*}
ht :: 'a \text{ tree}_{\text{ht}} & \Rightarrow \text{nat} \\
ht \langle \rangle & = 0 \\
ht \langle \_ , \_ , n , \_ \rangle & = n
\end{align*}
\]

\[
\begin{align*}
\text{node} :: 'a \text{ tree}_{\text{ht}} & \Rightarrow 'a \Rightarrow 'a \text{ tree}_{\text{ht}} \Rightarrow 'a \text{ tree}_{\text{ht}} \\
\text{node} \ l \ a \ r & = \langle l , \langle a , \text{max} \ (ht \ l) \ (ht \ r) + 1 \rangle , r \rangle
\end{align*}
\]

An AVL tree is a tree that satisfies the AVL invariant: the height of the left and right child of any node differ by at most 1

\[
\begin{align*}
\text{avl} :: 'a \text{ tree}_{\text{ht}} & \Rightarrow \text{bool} \\
\text{avl} \langle \rangle & = \text{True} \\
\text{avl} \ l , \langle \_ , n , r \rangle & = \left( \lvert \text{int} \ (ht \ l) - \text{int} \ (ht \ r) \rvert \leq 1 \land \\
& \quad n = \text{max} \ (ht \ l) \ (ht \ r) + 1 \land \text{avl} \ l \land \text{avl} \ r \right)
\end{align*}
\]

and the height field contains the correct value. The conversion function \( \text{int} :: \text{nat} \Rightarrow \text{int} \) is required because on natural numbers \( 0 - n = 0 \).


9.1 Logarithmic Height

AVL trees have logarithmic height. The key insight for the proof is that $M \, n$, the minimal number of leaves of an AVL tree of height $n$, satisfies the recurrence relation $M \, (n + 2) = M \, (n + 1) + M \, n$. Instead of formalizing this function $M$ we prove directly that an AVL tree of height $n$ has at least $\text{fib} \, (n + 2)$ leaves where $\text{fib}$ is the Fibonacci function:

\[
\text{fib} :: \text{nat} \Rightarrow \text{nat}
\]

\[
\text{fib} \, 0 = 0
\]

\[
\text{fib} \, 1 = 1
\]

\[
\text{fib} \, (n + 2) = \text{fib} \, (n + 1) + \text{fib} \, n
\]

**Lemma 9.1.** $\text{avl} \, t \rightarrow \text{fib} \, (h \, t + 2) \leq |t|_1$

*Proof. The proof is by induction on $t$. We focus on the induction step $t = (l, (a, n), r)$ and assume $\text{avl} \, t$. Thus the IHs reduce to $\text{fib} \, (h \, l + 2) \leq |l|_1$ and $\text{fib} \, (h \, r + 2) \leq |r|_1$. We prove $\text{fib} \, (\text{max} \, (h \, l) \, (h \, r) + 3) \leq |l|_1 + |r|_1$, from which $\text{avl} \, t \rightarrow \text{fib} \, (h \, t + 2) \leq |t|_1$ follows directly. There are two cases. We focus on $h \, l \geq h \, r$, $h \, l < h \, r$ is dual.

\[
\begin{align*}
\text{fib} \, (\text{max} \, (h \, l) \, (h \, r) + 3) &= \text{fib} \, (h \, l + 3) \\
&= \text{fib} \, (h \, l + 2) + \text{fib} \, (h \, l + 1) \\
&\leq |l|_1 + \text{fib} \, (h \, l + 1) \quad \text{by $\text{fib} \, (h \, l + 2) \leq |l|_1$} \\
&\leq |l|_1 + |r|_1 \quad \text{by $\text{fib} \, (h \, r + 2) \leq |r|_1$}
\end{align*}
\]

The last step is justified because $h \, l + 1 \leq h \, r + 2$ (which follows from $\text{avl} \, t$) and $\text{fib}$ is monotone. \qed

Now we prove a well-known exponential lower bound for $\text{fib}$ where $\varphi \equiv (1 + \sqrt{5}) / 2$:

**Lemma 9.2.** $\varphi^n \leq \text{fib} \, (n + 2)$

*Proof. The proof is by induction on $n$ by $\text{fib}$ computation induction. The case $n = 0$ is trivial and the case $n = 1$ is easy. Now consider the induction step:

\[
\begin{align*}
\text{fib} \, (n + 2 + 2) &= \text{fib} \, (n + 2 + 1) + \text{fib} \, (n + 2) \\
&\geq \varphi^n + \varphi^n \quad \text{by IHs} \\
&= (\varphi + 1) \cdot \varphi^n \\\n&= \varphi^{n + 2} \quad \text{because $\varphi + 1 = \varphi^2$} \quad \square
\end{align*}
\]

Combining the two lemmas yields $\text{avl} \, t \rightarrow \varphi^t \leq |t|_1$ and thus

**Corollary 9.3.** $\text{avl} \, t \rightarrow h \, t \leq 1 / \lg \, \varphi \cdot \lg \, |t|_1$
9.2 Implementation of ADT Set

9.2.1 Insertion

Insertion follows the standard approach: insert the element as usual and reestablish the AVL invariant on the way back up.

\[
\begin{align*}
\text{insert} :: 'a => 'a tree ht => 'a tree ht \\
\text{insert} x () = () \\
\text{insert} x (l, (a, n), r) = \begin{cases}
\text{case cmp x a of} \\
LT & \Rightarrow \text{balL (insert x l) a r} \\
EQ & \Rightarrow (l, (a, n), r) \\
GT & \Rightarrow \text{balR l a (insert x r)}
\end{cases}
\end{align*}
\]

Functions \text{balL}/\text{balR} readjust the tree after an insertion into the left/right child. The AVL invariant has been lost if the difference in height has become 2 — it cannot become more because the height can only increase by 1. Consider the definition of \text{balL} in Figure 9.1 (\text{balR} in Figure 9.2 is dual). If the AVL invariant has not been lost, i.e., if \(ht AB \neq ht C + 2\), then we can just return the AVL tree \(\text{node AB c C}\). But if \(ht AB = ht C + 2\), we need to “rotate” the subtrees suitably. Clearly \(AB\) must be of the form \(h A; (a, _)\); \(B\). There are two cases, which are illustrated in Figure 9.1. Triangles of the same height denote trees of the same height. A +1 at the bottom denotes an additional level due to insertion of the new element.

If \(ht B \leq ht A\) then \text{balL} performs what is known as a single rotation.

If \(ht A < ht B\) then \(B\) must be of the form \((B_1, (b, _), B_2)\) (where either \(B_1\) or \(B_2\) has increased in height) and \text{balL} performs what is known as a double rotation.

It is easy to check that in both cases the tree on the right satisfies the AVL invariant.

Preservation of \(avl\) by \text{insert} cannot be proved in isolation but needs to be proved simultaneously with how \text{insert} changes the height (because \(avl\) depends on the height and \text{insert} requires \(avl\) for correct behaviour):

**Theorem 9.4.** \(avl t \rightarrow avl (insert x t) \land h (insert x t) \in \{h t, h t + 1\}\)

The proof is by induction on \(t\) followed by a complete case analysis (which Isabelle automates).

9.2.2 Deletion

Figure 9.3 shows deletion-by-replacing (see 5.2.1). The recursive calls are dual to insertion: in terms of the difference in height, deletion of some element from one child...
\[ \text{balL} :: 'a \text{ tree}_\text{ht} \Rightarrow 'a \Rightarrow 'a \text{ tree}_\text{ht} \Rightarrow 'a \text{ tree}_\text{ht} \]

\[
\text{balL} \ AB \ c \ C \\
= (\text{if} \ \text{ht} \ AB = \text{ht} \ C + 2 \\
\quad \text{then case} \ AB \ \text{of} \\
\quad \quad \langle A, (a, x), B \rangle \Rightarrow \\
\quad \quad \quad \text{if} \ \text{ht} \ B \leq \text{ht} \ A \ \text{then} \ \text{node} \ A \ a \ (\text{node} \ B \ c \ C) \\
\quad \quad \quad \text{else case} \ B \ \text{of} \\
\quad \quad \quad \quad \langle B_1, (b, \_), B_2 \rangle \Rightarrow \ \text{node} \ (\text{node} \ A \ a \ B_1) \ b \ (\text{node} \ B_2 \ c \ C) \\
\quad \quad \quad \text{else} \ \text{node} \ AB \ c \ C)
\]

---

**Figure 9.1** Function \(\text{balL}\)
is the same as insertion of some element into the other child. Thus functions \(balR/balL\) can again be employed to restore the invariant.

An element is deleted from a node by replacing it with the maximal element of the left child (the minimal element of the right child would work just as well). Function \(split\_max\) performs that extraction and uses \(balL\) to restore the invariant after splitting an element off the right child.
The fact that balR/balL can be reused for deletion can be illustrated by drawing the corresponding rotation diagrams. We look at how the code for balL behaves when an element has been deleted from C. Dashed rectangles at the bottom indicate a single additional level that may or may not be there. A -1 indicates that the level has disappeared due to deletion.

Single rotation in balL after deletion in C:

Double rotation in balL after deletion in C:

At least one of B_1 and B_2 must have the same height as A.

Preservation ofavl by delete can be proved in the same manner as for insert but we provide more of the details (partly because our Isabelle proof is less automatic). The following lemmas express that the auxiliary functions preserve avl:

\[
\begin{align*}
\text{avl } l & \land \text{avl } r \land h \ r - 1 \leq h \ l \land h \ l \leq h \ r + 2 \rightarrow \text{avl } (\text{balL } l \ a \ r) \\
\text{avl } l & \land \text{avl } r \land h \ l - 1 \leq h \ r \land h \ r \leq h \ l + 2 \rightarrow \text{avl } (\text{balR } l \ a \ r) \\
\text{avl } t & \land t \neq \{} \rightarrow \\
\text{avl } (\text{fst } (\text{split_max } t)) & \land \\
\text{h } t \in \{h (\text{fst } (\text{split_max } t)), h (\text{fst } (\text{split_max } t)) + 1\}
\end{align*}
\]

The first two are proved by the obvious cases analyses, the last one also requires induction.
As for insert, preservation of \textit{avl} by delete needs to be proved simultaneously with how delete changes the height:

\textbf{Theorem 9.5.} \textit{avl} \( t \land t' = \text{delete} \ x \ t \longrightarrow \text{avl} \ t' \land h \ t \in \{ h \ t', h \ t' + 1 \} \)

\textit{Proof.} The proof is by induction on \( t \) followed by the case analyses dictated by the code for delete. We sketch the induction step. Let \( t = (l, (a, n), r) \) and \( t' = \text{delete} \ x \ t \) and assume the IHs and \textit{avl} \( t \). The claim \textit{avl} \( t' \) follows from the preservation of \textit{avl} by balL, balR and split\_max as shown above. The claim \( h \ t \in \{ h \ t', h \ t' + 1 \} \) follows directly from the definitions of balL and balR. \( \square \)

\section*{9.3 Exercises}

\textbf{Exercise 9.1.} The logarithmic height of AVL trees can be proved directly. Prove \( \text{avl} \ t \land h \ t = n \longrightarrow 2^\text{n div 2} \leq |t|_1 \) by \textit{fib} computation induction on \( n \). This implies \( \text{avl} \ t \longrightarrow h \ t \leq 2 \cdot \lg |t|_1 \).

\textbf{Exercise 9.2.} Fibonacci trees are defined in analogy to Fibonacci numbers:

\begin{verbatim}
  fibt : nat \rightarrow unit tree
  fibt 0 = ()
  fibt 1 = ((), ()
  fibt (n + 2) = (fibt (n + 1), (), fibt n)
\end{verbatim}

We are only interested in the shape of these trees. Therefore the nodes just contain dummy unit values (). Hence we need to define the AVL invariant again for trees without annotations:

\begin{verbatim}
  avl0 : 'a tree \Rightarrow bool
  avl0 () = True
  avl0 (l, _, r) = (|int (h l) - int (h r)| \leq 1 \land avl0 l \land avl0 r)
\end{verbatim}

Prove the following properties of Fibonacci trees:

\( \text{avl0} (fibt n) \quad |\text{fibt} \ n|_1 = \text{fib} (n + 2) \)

Conclude that the Fibonacci trees are minimal (w.r.t. their size) among all AVL trees of a given height:

\( \text{avl} \ t \longrightarrow |\text{fibt} \ (h \ t)|_1 \leq |t|_1 \)

\textbf{Exercise 9.3.} Show that every almost complete tree is an AVL tree:

\( \text{acomplete} \ t \longrightarrow \text{avl0} \ t \)

As in the previous exercise we consider trees without height annotations.

\textbf{Exercise 9.4.} Generalize AVL trees to height-balanced trees where the condition
in the invariant is replaced by
\[ |\text{int} (h_l) - \text{int} (h_r)| \leq m \]
where \( m \geq 1 \) is some fixed integer. Modify the invariant and the insertion and deletion functions and prove that the latter fulfill the same correctness theorems as before. You do not need to prove the logarithmic height of height-balanced trees.

Exercise 9.5. Following Section 7.3, define a linear-time function `avl_of_list :: 'a list => 'a tree_ht` and prove both `inorder (avl_of_list as) = as` and `avl (avl_of_list as)`.

### 9.4 An Optimization

Instead of recording the height of the tree in each node, it suffices to record the balance factor, i.e. the difference in height of its two children. Rather than the three integers -1, 0 and 1 we utilize a new data type:

```plaintext
datatype bal = Lh | Bal | Rh

type_synonym 'a tree_bal = ('a * bal) tree
```

The names `Lh` and `Rh` stand for “left-heavy” and “right-heavy”. The AVL invariant for these trees reflect these names:

```plaintext
avl :: 'a tree_bal => bool
avl () = True
avl (l, (_ , b), r) = ((case b of
    Lh => h_l = h_r + 1 |
    Bal => h_r = h_l |
    Rh => h_r = h_l + 1) \and
    avl l \and avl r)
```

The code for insertion (and deletion) is similar to the height-based version. The key difference is that the test if the AVL invariant as been lost cannot be based on the height anymore. We need to detect if the tree has increased in height upon insertion based on the balance factors. The key insight is that a height increase is coupled with a change from `Bal` to `Lh` or `Rh`. Except when we transition from `()` to `()`, `(a, Bal)`, `(a)`. This insight is encoded in the test `incr`:
9.4 An Optimization

The test for a height increase compares the trees before and after insertion. Therefore it has been pulled out of the balance functions into insertion:

\[
\text{is\_bal} :: \text{a} \rightarrow \text{b} \rightarrow \text{bool} \\
\text{is\_bal} (\text{a}, \text{b}) = (\text{b} = \text{Bal})
\]

\[
\text{incr} :: \text{a} \rightarrow \text{b} \rightarrow \text{bool} \\
\text{incr} t t' = (t \neq \text{a} \lor t' \neq \text{a} \lor \text{is\_bal} t' \land \neg \text{is\_bal} t)
\]

The balance functions are shown in Figure 9.4. Function rot2 implements double rotations. Function balL is called if the left child AB has increased in height. If the tree was LH then single or double rotations are necessary to restore balance. Otherwise we simply need to adjust the balance factors. Function balR is dual to balL.

For deletion we need to test if the height has decreased and decr implements this test:

\[
\text{decr} :: \text{a} \rightarrow \text{b} \rightarrow \text{bool} \\
\text{decr} t t' = (t \neq \text{a} \lor t' \neq \text{a} \lor \text{is\_bal} t' \land \neg \text{is\_bal} t')
\]

The functions incr and decr are almost dual except that incr implicitly assumes \( t' \neq \text{a} \lor \text{t}' \neq \text{a} \lor \text{is\_bal} t' \land \neg \text{is\_bal} t \) because insertion is guaranteed to return a Node. Thus we could use decr instead of incr but not the other way around.
Figure 9.4 Functions balL and balR
Deletion and split_max change in the same manner as insertion:

\[
delete :: 'a \Rightarrow 'a \text{ tree_bal} \Rightarrow 'a \text{ tree_bal}
delete \_ () = ()
delete x (l, (a, ba), r)
= (\text{case} \ cmp \ x \ a \ of
  LT \Rightarrow \text{let} \ l' = \text{delete} \ x \ l
    \text{in} \ if \ \text{decr} \ l \ l' \ \text{then} \ \text{balR} \ l' \ a \ ba \ \text{else} \ (l', (a, ba), r)
  | EQ \Rightarrow \text{if} \ l = () \ \text{then} \ r
    \text{else} \ \text{let} \ (l', a') = \text{split_max} \ l
      \text{in} \ if \ \text{decr} \ l \ l' \ \text{then} \ \text{balR} \ l' \ a' \ ba \ r
      \text{else} \ (l', (a', ba), r)
  | GT \Rightarrow \text{let} \ r' = \text{delete} \ x \ r
    \text{in} \ if \ \text{decr} \ r \ r' \ \text{then} \ \text{balL} \ l \ a \ ba \ r' \ \text{else} \ (l, (a, ba), r'))
\]

split_max :: 'a \text{ tree_bal} \Rightarrow 'a \text{ tree_bal} \times 'a

split_max (l, (a, ba), r)
= (if r = () \ then \ (l, a)
  \text{else} \ \text{let} \ (r', a') = \text{split_max} \ r;
    t' = if \ \text{decr} \ r \ r' \ \text{then} \ \text{balL} \ l \ a \ ba \ r' \ \text{else} \ (l, (a, ba), r')
  \text{in} \ (t', a'))

In the end we have the following correctness theorems:

**Theorem 9.6.** \( \text{avl} \ t \land t' = \text{insert} \ x \ t \longrightarrow \text{avl} \ t' \land h \ t' = h \ t + (\text{if} \ \text{incr} \ t \ t' \ \text{then} \ 1 \ \text{else} \ 0) \)

This theorem tells us not only that \( \text{avl} \) is preserved but also that \( \text{incr} \) indicates correctly if the height has increased or not. Similarly for deletion and \( \text{decr} \):

**Theorem 9.7.** \( \text{avl} \ t \land t' = \text{delete} \ x \ t \longrightarrow \text{avl} \ t' \land h \ t' = h \ t + (\text{if} \ \text{decr} \ t \ t' \ \text{then} \ 1 \ \text{else} \ 0) \)

The proofs of both theorems follow the standard pattern of induction followed by an exhaustive (automatic) cases analysis. The proof for \text{delete} requires an analogous lemma for \text{split_max}:

\[
\text{split_max} \ t = (t', a) \land \text{avl} \ t \land t \neq () \longrightarrow \text{avl} \ t' \land h \ t' = h \ t + (\text{if} \ \text{decr} \ t \ t' \ \text{then} \ 1 \ \text{else} \ 0)
\]
9.5 Exercises

Exercise 9.6. We map type \('a\ tree\_bal\) back to type \('a\ \times nat\) tree called \('a\ tree\_ht\) in the beginning of the chapter:

\[
debal :: \text{type}\,(\text{a}\ tree\_bal) \Rightarrow (\text{type}\,(\text{a}\ 
\times nat)\ tree)
\]

\[
debal\;\langle\rangle = \langle\rangle \\
\text{debal}\;\langle l, (a,\;\_),\;r\rangle = \langle\text{debal}\;l, (a,\;\max\;\text{h}\;l\;\text{h}\;r)\;+\;1),\;\text{debal}\;r\rangle
\]

Prove that the AVL property is preserved: \(avl\;t \rightarrow avl\_ht\; \text{debal}\;t\) where \(avl\_ht\) is the \(avl\) predicate on type \('a\ tree\_ht\) from the beginning of the chapter.

Define a function \(\text{debal2}\) of the same type that traverses the tree only once and in particular does not use function \(h\). Prove \(avl\;t \rightarrow \text{debal2}\;t = \text{debal}\;t\).

Exercise 9.7. To realize the full space savings potential of balance factors we encode them directly into the node constructors and work with the following special tree type:

\[
datatype\;\text{a}\;\text{tree4} = \text{Leaf} \\
| \text{Lh}\;\text{a}\;\text{tree4}\;\text{a}\;\text{a}\;\text{tree4} \\
| \text{Bal}\;\text{a}\;\text{tree4}\;\text{a}\;\text{a}\;\text{tree4} \\
| \text{Rh}\;\text{a}\;\text{tree4}\;\text{a}\;\text{a}\;\text{tree4}
\]

On this type define the AVL invariant, insertion, deletion and all necessary auxiliary functions. Prove theorems 9.6 and 9.7. Hint: modify the theory underlying Section 9.4.
10 Beyond Insert and Delete: $\cup$, $\cap$ and $\setminus$

Tobias Nipkow

So far we looked almost exclusively at insertion and deletion of single elements, with the exception of the conversion of whole lists of elements into search trees (see Section 7.3 and Exercises 8.3 and 9.5). This chapter is dedicated to operations that combine two sets (implemented by search trees) by union, intersection and difference. We denote set difference by $\setminus$ rather than $\setminus$.

Let us focus on set union for a moment and assume that insertion into a set of size $s$ takes time proportional to $\lg s$. Consider two sets $A$ and $B$ of size $m$ and $n$ where $m \leq n$. The naive approach is to insert the elements from one set one by one into the other set. This takes time proportional to $\lg n + \cdots + \lg(n + m - 1)$ or $\lg m + \cdots + \lg(m + n - 1)$ depending on whether the smaller set is inserted into the larger one or the other way around. Of course the former sum is less than or equal to the latter sum. To estimate the growth of $\lg n + \cdots + \lg(n + m - 1) = \lg(n \cdots (n + m - 1))$ we can easily generalize the derivation of $\lg(n!) \in \Theta(n \lg n)$ in the initial paragraph of Section 7.3. The result is $\lg(n \cdots (n + m - 1)) \in \Theta(m \lg n)$. That is, inserting $m$ elements into an $n$ element set one by one takes time $\Theta(m \lg n)$.

There is a second, possibly naive sounding algorithm for computing the union: flatten both trees to ordered lists (using function $\text{inorder}$ from Exercise 4.1), merge both lists and convert the resulting list back into a suitably balanced search tree. All three steps take linear time. The last step is the only slightly nontrivial one but has been dealt with before (see Section 7.3 and Exercises 8.3 and 9.5). This algorithm takes time $O(m + n)$ which is significantly better than $O(m \lg n)$ if $m \approx n$ but significantly worse if $m << n$.

This chapter presents a third approach which has the following salient features:

- Union, intersection and difference take time $O(m \lg(n + 1))$
- It works for a whole class of balanced trees, including AVL, red-black and weight-balanced trees.
- It is based on a single function for joining two balanced trees to form a new balanced tree.
ADT $\text{Set2} = \text{Set} +$

interface
union :: 's ⇒ 's ⇒ 's
inter :: 's ⇒ 's ⇒ 's
diff :: 's ⇒ 's ⇒ 's

specification
\[
\begin{align*}
invar s_1 ∧ invar s_2 & \rightarrow \text{set}(\text{union } s_1 s_2) = \text{set } s_1 ∪ \text{set } s_2 \quad \text{(union)} \\
invar s_1 ∧ invar s_2 & \rightarrow \text{invar}(\text{union } s_1 s_2) \quad \text{(union-inv)} \\
invar s_1 ∧ invar s_2 & \rightarrow \text{set}(\text{inter } s_1 s_2) = \text{set } s_1 ∩ \text{set } s_2 \quad \text{(inter)} \\
invar s_1 ∧ invar s_2 & \rightarrow \text{invar}(\text{inter } s_1 s_2) \quad \text{(inter-inv)} \\
invar s_1 ∧ invar s_2 & \rightarrow \text{set}(\text{diff } s_1 s_2) = \text{set } s_1 − \text{set } s_2 \quad \text{(diff)} \\
invar s_1 ∧ invar s_2 & \rightarrow \text{invar}(\text{diff } s_1 s_2) \quad \text{(diff-inv)}
\end{align*}
\]

Figure 10.1 ADT Set2

We call it the join approach. It is easily and efficiently parallelizable, a property we will not explore here.

The join approach is at least as fast as the one-by-one approach: from $m + n ≤ mn$ it follows that $\frac{n}{m} + 1 ≤ n$ (if $m, n ≥ 2$). The join approach is also at least as fast as the tree-to-list-to-tree approach because $m + n = m(\frac{n}{m} + 1)$ (if $m ≥ 1$).

10.1 Specification of Union, Intersection and Difference

Before explaining the join approach we extend the ADT Set by three new functions union, inter and diff. The specification in Figure 10.1 is self-explanatory.

10.2 Just Join

Now we come to the heart of the matter, the definition of union, intersection and difference in terms of a single function join. We promised that the algorithms would be generic across a range of balanced trees. Thus we assume that we operate on augmented trees of type ($'a × 'b$) tree where '$a$ is the type of the elements and '$b$ is the balancing information (which we can ignore here). This enables us to formulate the algorithms via pattern-matching. A more generic approach is the subject of Exercise 10.2.

The whole section is parameterized by the join function and an invariant:
\[
\begin{align*}
\text{join} :: ('a × 'b) \text{ tree} & ⇒ 'a ⇒ ('a × 'b) \text{ tree} ⇒ ('a × 'b) \text{ tree} \\
\text{inv} :: ('a × 'b) \text{ tree} & ⇒ \text{bool}
\end{align*}
\]
**Figure 10.2** Specification of \( \text{join} \) and \( \text{inv} \)

Function \( \text{inv} \) is meant to take care of the balancedness property only, not the BST property. Functions \( \text{join} \) and \( \text{inv} \) are specified with the help of the standard tree functions \( \text{set\_tree} \) and \( \text{bst} \) in Figure 10.2. With respect to the set of elements, \( \text{join} \) must behave like union. But it need only return a BST if both trees are BSTs and the element \( a \) lies in between the elements of the two trees, i.e. if \( \text{bst} \langle l, (a, \_), r \rangle \). The structural invariant \( \text{inv} \) must be preserved by formation and destruction of trees. Thus we can see \( \text{join} \) as a smart constructor that builds a balanced tree.

To define union and friends we need a number of simple auxiliary functions. Function \( \text{split\_min} \) decomposes a tree into its leftmost (minimal) element and the remaining tree; the remaining tree is reassembled via \( \text{join} \), thus preserving \( \text{inv} \):

\[
\text{split\_min} :: \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle
\]

\[\text{split\_min} \langle l, (a, \_), r \rangle = (\text{if } l = \langle \rangle \text{ then } (a, r) \text{ else let } (m, l') = \text{split\_min} l \text{ in } (m, \text{join} l' a r))\]

Function \( \text{join2} \) is reduced to \( \text{join} \) with the help of \( \text{split\_min} \):

\[
\text{join2} :: \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle
\]

\[\text{join2} l r = (\text{if } r = \langle \rangle \text{ then } l \text{ else let } (m, r') = \text{split\_min} r \text{ in } \text{join} l m r')\]

Function \( \text{split} \) splits a BST w.r.t. a given element \( a \) into a triple \( (l, b, r) \) such that \( l \) contains the elements less than \( a \), \( r \) contains the elements greater than \( a \), and \( b \) is true iff \( a \) was in the input tree:

\[
\text{split} :: 'a \Rightarrow \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle \times \text{bool} \times \langle 'a \times 'b \rangle \Rightarrow \langle 'a \times 'b \rangle
\]

\[\text{split} \langle \rangle = (\langle \rangle, \text{False}, \langle \rangle)\]
Chapter 10  Beyond Insert and Delete: \( \cup, \cap \) and \(-\)

\[
split x \langle l, (a, \_), r \rangle \\
= (\text{case } \text{cmp } x a \text{ of} \\
\quad LT \Rightarrow (l_1, b, l_2) = \text{split } x \text{ in } l_1, b, \text{join } l_2 \text{ a } r \mid \\
\quad EQ \Rightarrow (l, \text{True}, r) \mid \\
\quad GT \Rightarrow (r_1, b, r_2) = \text{split } x \text{ in } \text{join } l a r_1, b, r_2)
\]

The following example demonstrates the workings of \(\text{split}\):

\[
\begin{array}{c}
\text{Assume } a < b < c < d < e. \text{ The call } \text{split } c \text{ descends the input BST along the path } a, e, b, d, \text{ splits the tree into two parts on each level and reassembles the parts into the two separate output trees on the way back up using } \text{join}. \text{ For simplicity the example assumes that } \text{join} \text{ just puts the subtrees together but no rebalancing is needed.} \\
\text{Insertion and deletion can be define in terms of the } \text{split} \text{ and } \text{join}:
\end{array}
\]

\[
\begin{align*}
\text{insert ::} & \quad 'a \Rightarrow ('a \times 'b) \text{ tree } \Rightarrow ('a \times 'b) \text{ tree} \\
& \quad \text{insert } x t = (\text{let } (l, b, r) = \text{split } x \text{ in } \text{join } l x r)
\end{align*}
\]

\[
\begin{align*}
\text{delete ::} & \quad 'a \Rightarrow ('a \times 'b) \text{ tree } \Rightarrow ('a \times 'b) \text{ tree} \\
& \quad \text{delete } x t = (\text{let } (l, b, r) = \text{split } x \text{ in } \text{join2 } l r)
\end{align*}
\]

The efficiency can be improved a little by taking the returned \(b\) into account (how?). Alternatively, insertion and deletion can be defined by means of union and difference (Exercise 10.1).

But we have bigger functions to fry: union, intersection and difference. They are shown in Figure 10.3. All three are divide-and-conquer algorithms that follow the same schema: both input trees are split at an element \(a\) (by construction or explicitly), the algorithm is applied recursively to the two trees of the elements below \(a\) and to the two trees of the elements above \(a\), and the two results are suitably joined.

The following diagram demonstrates the behaviour of \(\text{union}\):
10.2 Just Join

union :: ('a × 'b) tree ⇒ ('a × 'b) tree ⇒ ('a × 'b) tree
union () t = t
union t () = t
union (l₁, (a, _), r₁) t₂
= (let (l₂, b₂, r₂) = split a t₂
    in join (union l₁ l₂) a (union r₁ r₂))

inter :: ('a × 'b) tree ⇒ ('a × 'b) tree ⇒ ('a × 'b) tree
inter () t = ()
inter t () = ()
inter (l₁, (a, _), r₁) t₂
= (let (l₂, b₂, r₂) = split a t₂;
    l' = inter l₁ l₂; r' = inter r₁ r₂
    in if b₂ then join l' a r' else join2 l' r')

diff :: ('a × 'b) tree ⇒ ('a × 'b) tree ⇒ ('a × 'b) tree
diff () t = ()
diff t () = t
diff t₁ (l₂, (a, _), r₂)
= (let (l₁, b₁, r₁) = split a t₁
    in join2 (diff l₁ l₂) (diff r₁ r₂))

Figure 10.3  Union, intersection and difference

10.2.1 Correctness
We need to prove that union, inter and diff satisfy the specification in Figure 10.1 where set = set_tree and invar t = inv t ∧ bst t. That is, for each function we show its set-theoretic property and preservation of inv and bst using the assumptions in
Figure 10.2. Most of the proofs in this section are obvious and automatic inductions and we do not discuss them.

First we need to prove suitable properties of the auxiliary functions $\text{split}_\text{min}$, $\text{join}_2$ and $\text{split}$:

\[ \text{split}_\text{min} \; t = (m, t') \land t \neq \langle \rangle \longrightarrow \]
\[ m \in \text{set}_\text{tree} \; t \land \text{set}_\text{tree} \; t = \{m\} \cup \text{set}_\text{tree} \; t' \]

\[ \text{split}_\text{min} \; t = (m, t') \land \text{bst} \; t \land t \neq \langle \rangle \longrightarrow \]
\[ \text{bst} \; t' \land (\forall x \in \text{set}_\text{tree} \; t'. \; m < x) \]

\[ \text{split}_\text{min} \; t = (m, t') \land \text{inv} \; t \land t \neq \langle \rangle \longrightarrow \text{inv} \; t' \]

\[ \text{set}_\text{tree} \; (\text{join}_2 \; l \; r) = \text{set}_\text{tree} \; l \cup \text{set}_\text{tree} \; r \quad (10.5) \]

\[ \text{bst} \; l \land \text{bst} \; r \land (\forall x \in \text{set}_\text{tree} \; l. \; \forall y \in \text{set}_\text{tree} \; r. \; x < y) \longrightarrow \]
\[ \text{bst} \; (\text{join}_2 \; l \; r) \]

\[ \text{inv} \; l \land \text{inv} \; r \longrightarrow \text{inv} \; (\text{join}_2 \; l \; r) \]

\[ \text{split} \; x \; t = (l, b, r) \land \text{bst} \; t \longrightarrow \]
\[ \text{set}_\text{tree} \; l = \{a \in \text{set}_\text{tree} \; t \mid a < x\} \land \]
\[ \text{set}_\text{tree} \; r = \{a \in \text{set}_\text{tree} \; t \mid x < a\} \land \]
\[ b = (x \in \text{set}_\text{tree} \; t) \land \text{bst} \; l \land \text{bst} \; r \quad (10.6) \]

\[ \text{split} \; x \; t = (l, b, r) \land \text{inv} \; t \longrightarrow \text{inv} \; l \land \text{inv} \; r \]

The correctness properties of $\text{insert}$ and $\text{delete}$ are trivial consequences and are not shown. We move on to $\text{union}$. Its correctness properties are concretizations of the properties ($\text{union}$) and ($\text{union-inv}$) in Figure 10.1:

\[ \text{bst} \; t_2 \longrightarrow \text{set}_\text{tree} \; (\text{union} \; t_1 \; t_2) = \text{set}_\text{tree} \; t_1 \cup \text{set}_\text{tree} \; t_2 \]

\[ \text{bst} \; t_1 \land \text{bst} \; t_2 \longrightarrow \text{bst} \; (\text{union} \; t_1 \; t_2) \]

\[ \text{inv} \; t_1 \land \text{inv} \; t_2 \longrightarrow \text{inv} \; (\text{union} \; t_1 \; t_2) \]

All three $\text{union}$ properties are proved by computation induction. The first property follows easily from assumption (10.1) and (10.6). The assumption $\text{bst} \; t_2$ (but not $\text{bst} \; t_1$) is required because $t_2$ is split and (10.6) requires $\text{bst}$. Preservation of $\text{bst}$ follows from assumption (10.2) with the help of the first $\text{union}$ property and the preservation of $\text{bst}$ by $\text{split}$. Preservation of $\text{inv}$ follows from assumptions (10.3) and (10.4) with the help of the preservation of $\text{inv}$ by $\text{split}$.

The correctness properties of $\text{inter}$ look similar:

\[ \text{bst} \; t_1 \land \text{bst} \; t_2 \longrightarrow \text{set}_\text{tree} \; (\text{inter} \; t_1 \; t_2) = \text{set}_\text{tree} \; t_1 \cap \text{set}_\text{tree} \; t_2 \]

\[ \text{bst} \; t_1 \land \text{bst} \; t_2 \longrightarrow \text{bst} \; (\text{inter} \; t_1 \; t_2) \]

\[ \text{inv} \; t_1 \land \text{inv} \; t_2 \longrightarrow \text{inv} \; (\text{inter} \; t_1 \; t_2) \]
The proof of the preservation properties are automatic but the proof of the set\_tree property is more involved than the corresponding proof for union and we take a closer look at the induction. We focus on the case $t_1 = \langle l_1, (a, \_), r_1 \rangle$ and $t_2 \neq \emptyset$. Let $L_1 = set\_tree l_1$ and $R_1 = set\_tree r_1$. Let $(l_2, b, r_2) = split t_2 a, L_2 = set\_tree l_2, R_2 = set\_tree r_2$ and $A = (if b then \{a\} else \{\})$. The separation properties

$$a \notin L_1 \cup R_1 \quad a \notin L_2 \cup R_2$$

$$L_2 \cap R_2 = \{\} \quad L_1 \cap R_2 = \{\} \quad L_2 \cap R_1 = \{\}$$

follow from bst $t_1$, bst $t_2$ and (10.6). Now for the main proof:

$$set\_tree t_1 \cap set\_tree t_2 = (L_1 \cup R_1 \cup \{a\}) \cap (L_2 \cup R_2 \cup A)$$

by (10.6), bst $t_2$

$$= L_1 \cap L_2 \cap R_1 \cap R_2 \cup A$$

by the separation properties

$$= set\_tree (inter t_1 t_2)$$

by (10.1), (10.5), IHs, bst $t_1$, bst $t_2$, (10.6)

The correctness properties of diff follow the same pattern and their proofs are similar to the proofs of the inter properties. This concludes the generic join approach.

### 10.3 Joining Red-Black Trees

This section shows how to implement join efficiently on red-black trees. The basic idea is simple: descend along the spine of the higher of the two trees until reaching a subtree whose height is the same as the height of the lower tree. With suitable changes this works for other balanced trees as well [Blelloch et al. 2022]. The function definitions are shown in Figure 10.4. Function join calls joinR (ascending along the right spine of l) if l is the higher tree, or calls joinL (descending along the left spine of r) if r is the higher tree, or returns $B l x r$ otherwise. The running time is linear in the black height (and thus logarithmic in the size) if we assume that the black height is stored in each node; our implementation of red-black trees would have to be augmented accordingly. Note that in joinR (and similarly in joinL) the comparison is not $bh l = bh r$ but $bh l \leq bh r$ to simplify the proofs.

#### 10.3.1 Correctness

We need to prove that join on red-black trees (and a suitable inv) satisfies its specification in Figure 10.2. We start with properties of joinL; the properties of function joinR are completely symmetric. These are the three automatically provable inductive propositions:

$$invc \ l \land invc \ r \land invh \ l \land invh \ r \land bh \ l \leq bh \ r \rightarrow$$

$$invc2 \ (joinL \ l x r) \land$$

$$(bh \ l \neq bh \ r \land color \ r = \text{Black} \rightarrow invc \ (joinL \ l x r)) \land$$

$$invh \ (joinL \ l x r) \land bh \ (joinL \ l x r) = bh \ r$$
joinL :: 'a rbt ⇒ 'a ⇒ 'a rbt ⇒ 'a rbt
joinL l x r = (if bh r ≤ bh l then R l x r
  else case r of
    ⟨l', (x', Red), r'⟩ ⇒ R (joinL l x l') x' r'
    ⟨l', (x', Black), r'⟩ ⇒ baliL (joinL l x l') x' r')

joinR :: 'a rbt ⇒ 'a ⇒ 'a rbt ⇒ 'a rbt
joinR l x r = (if bh l ≤ bh r then R l x r
  else case l of
    ⟨l', (x', Red), r'⟩ ⇒ R l x (joinR r' x r)
    ⟨l', (x', Black), r'⟩ ⇒ baliR l x (joinR r' x r))

join :: 'a rbt ⇒ 'a ⇒ 'a rbt ⇒ 'a rbt
join l x r = (if bh r < bh l then paint Black (joinR l x r)
  else if bh l < bh r then paint Black (joinL l x r)
  else B l x r)

Figure 10.4 Function join on red-black trees

bh l ≤ bh r → set_tree (joinL l x r) = set_tree l ∪ \{x\} ∪ set_tree r
bst ⟨l, (a, n), r⟩ ∧ bh l ≤ bh r → bst (joinL l a r)

Because joinL employs baliL from the chapter on red-black trees, the proof of the first proposition makes use of the property of baliL displayed in Section 8.2.1.

We define the invariant inv required for the specification in Figure 10.2 as follows:

inv t = (invc t ∧ inv h t)

Although weaker than rbt, it still guarantees logarithmic height (see Exercise 8.1). Note that rbt itself does not work because it does not satisfy property (10.4). The properties of join and inv are now easy consequences of the joinL (and joinR) properties shown above.
10.4 Exercises

Exercise 10.1. Define alternative versions \( \text{insert}' \) and \( \text{delete}' \) of \( \text{insert} \) and \( \text{delete} \) using \( \text{union} \) and \( \text{diff} \) (and \( \text{join} \) and \( \langle \rangle \)). Prove their correctness as in Section 10.2.1: \( \text{set_tree} \) yields the right result and \( \text{bst} \) is preserved.

Exercise 10.2. Define an alternative version \( \text{diff1} \) of \( \text{diff} \) where in the third equation pattern matching is on \( t_1 \) and \( t_2 \) is \( \text{split} \). Prove that \( \text{bst} t_1 \wedge \text{bst} t_2 \) implies both \( \text{set_tree} (\text{diff1} t_1 t_2) = \text{set_tree} t_1 - \text{set_tree} t_2 \) and \( \text{bst} (\text{diff1} t_1 t_2) \).

Exercise 10.3. Following the general idea of the join function for red-black trees, define a join function for 2-3-trees. Start with two functions \( \text{joinL, joinR} :: 'a \text{ tree23} \Rightarrow 'a \Rightarrow 'a \text{ tree23} \Rightarrow 'a \text{ tree23} \) and combine them into the overall join function:

\[
\text{join} :: 'a \text{ tree23} \Rightarrow 'a \Rightarrow 'a \text{ tree23} \Rightarrow 'a \text{ tree23}
\]

Prove the following correctness properties:

\[
\text{complete } l \wedge \text{complete } r \rightarrow \text{complete } (\text{join } l x r)
\]
\[
\text{complete } l \wedge \text{complete } r \rightarrow \text{inorder } (\text{join } l x r) = \text{inorder } l \odot x \# \text{inorder } r
\]

The corresponding (and needed) properties of \( \text{joinL} \) and \( \text{joinR} \) are slightly more involved.

10.5 Chapter Notes

The join approach goes back to Adams [1993]. Blelloch et al. [2022] generalized the approach from weight-balanced trees to AVL trees, red-black trees and treaps. In particular they proved the \( O(m \log (\frac{n}{m} + 1)) \) bound for the work (and an \( O(\log m \log n) \) bound for the span).
Arrays via Braun Trees

Tobias Nipkow

Braun trees are a subclass of almost complete trees. In this chapter we explore their use as arrays and in Chapter 15 as priority queues.

11.1 Array

So far we have discussed sets (or maps) over some arbitrary linearly ordered type. Now we specialize that linearly ordered type to \textit{nat} to model arrays. In principle we could model arrays as maps from a subset of natural numbers to the array elements. Because arrays are contiguous, it is more appropriate to model them as lists. The type \texttt{\textquotesingle a list} comes with two array-like operations (see Appendix A):

- **Indexing:** \texttt{xs!n} is the \textit{n}th element of the list \texttt{xs}.
- **Updating:** \texttt{xs[n := x]} is \texttt{xs} with the \textit{n}th element replaced by \texttt{x}.

By convention, indexing starts with \texttt{n = 0}. If \texttt{n \geq length xs} then \texttt{xs!n} and \texttt{xs[n := x]} are underdefined: they are defined terms but we do not know what their value is.

Note that operationally, indexing and updating take time linear in the index, which may appear inappropriate for arrays. However, the type of lists is only an abstract model that specifies the desired functional behaviour of arrays but not their running time complexity.

The ADT of arrays is shown in Figure 11.1. Type \texttt{'a ar} is the type of arrays, type \texttt{'a} the type of elements in the arrays. The abstraction function \texttt{list} abstracts arrays to lists. It would make perfect sense to include \texttt{list} in the interface as well. In fact, our implementation below comes with a (reasonably efficiently) executable definition of \texttt{list}.

The behaviour of \texttt{lookup}, \texttt{update}, \texttt{size} and \texttt{array} is specified in terms of their counterparts on lists and requires that the invariant is preserved. What distinguishes the specifications of \texttt{lookup} and \texttt{update} from the standard schema (see Chapter 6) is that they carry a size precondition because the result of \texttt{lookup} and \texttt{update} is only specified if the index is less than the size of the array.
ADT Array =

interface
lookup :: 'ar ⇒ nat ⇒ 'a
update :: nat ⇒ 'a ⇒ 'ar ⇒ 'ar
len :: 'ar ⇒ nat
array :: 'a list ⇒ 'ar

abstraction list :: 'ar ⇒ 'a list
invariant invar :: 'ar ⇒ bool

specification
invar ar ∧ n < len ar → lookup ar n = list ar ! n (lookup)
invar ar ∧ n < len ar → list (update n x ar) = (list ar)[n := x] (update-inv)
invar ar ∧ n < len ar → invar (update n x ar) (update-inv)
invar ar ⇒ len ar = |list ar| (len)
list (array xs) = xs (array)
invar (array xs) (array-inv)

Figure 11.1 ADT Array

11.2 Braun Trees

One can implement arrays by any one of the many search trees presented in this book. Instead we take advantage of the fact that the keys are natural numbers and implement arrays by so-called Braun trees which are almost complete and thus have minimal height.

The basic idea is to index a node in a binary tree by the non-zero bit string that leads from the root to that node in the following fashion. Starting from the least significant bit and while we have not reached the leading 1 (which is ignored), we examine the bits one by one. If the current bit is 0, descend into the left child, otherwise into the right child. Instead of bit strings we use the natural numbers 1–15 that they represent. The Braun tree with nodes indexed by 1–15 is shown in Figure 11.2. The numbers are the indexes and not the elements stored in the nodes. For example, the index 14 is 0111 in binary (least significant bit first). If you follow the path left-right-right corresponding to 011 in Figure 11.2 you reach node 14.

A tree t is suitable for representing an array if the set of indexes of all its nodes is the interval \{1..|t|\}. The following tree is unsuitable because the node indexed by 2 is missing:
It turns out that the following invariant guarantees that a tree $t$ contains exactly the nodes indexed by $1, \ldots, j_t$:

$$\text{braun} :: \forall a \text{tree } \Rightarrow \text{bool}$$

$$\text{braun} () = \text{True}$$

$$\text{braun} (l, _, r) = (|l| = |r| \lor |l| = |r| + 1) \land \text{braun } l \land \text{braun } r$$

The disjunction can alternatively be expressed as $|r| \leq |l| \leq |r| + 1$. We call a tree a **Braun tree** if it satisfies predicate $\text{braun}$.

Although we do not need or prove this here, it is interesting to note that a tree that contains exactly the nodes indexed by $1, \ldots, |t|$ is a Braun tree.

Let us now prove the earlier claim that Braun trees are almost complete. First, a lemma about the composition of almost complete trees:

**Lemma 11.1.** $\text{almost-complete } l \land \text{almost-complete } r \land |l| = |r| + 1 \rightarrow \text{almost-complete } (l, x, r)$

**Proof.** Using Lemmas 4.7 and 4.8 and the assumptions we obtain

$$h (l, x, r) = \lceil \log (|r| + 1) \rceil + 1 \quad (\ast)$$

$$mh (l, x, r) = \lceil \log |r| \rceil + 1 \quad (\ast\ast)$$

Because $1 \leq |r|$, there is an $i$ such that $2^i \leq |r| < 2^i + 1$ and thus $2^i < |r| + 1 \leq 2^i + 1$. This implies $i = \lceil \log |r| \rceil$ and $i + 1 = \lceil \log (|r| + 1) \rceil$. Together with $(\ast)$ and $(\ast\ast)$ this implies $\text{almost-complete } (l, x, r)$. 

Now we can show that all Braun trees are almost complete. Thus we know that they have optimal height (Lemma 4.6) and can even quantify it (Lemma 4.7).
Lemma 11.2. \(\text{braun } t \rightarrow \text{acomplete } t\)

Proof by induction. We focus on the induction step where \(t = \langle l, x, r \rangle\). By assumption we have \(\text{acomplete } l\) and \(\text{acomplete } r\). Because \(t = \text{braun } t\) we can distinguish two cases. First assume \(|l| = |r| + 1\). The claim \(\text{acomplete } t\) follows immediately from the previous lemma. Now assume \(|l| = |r|\). By definition, there are four cases to consider when proving \(\text{acomplete } t\). By symmetry it suffices to consider only two of them. If \(h \leq h_r\) and \(mh_r < mh_l\) then \(\text{acomplete } t\) reduces to \(\text{acomplete } r\), which is true by assumption. Now assume \(h \leq h_r\) and \(mh_l \leq mh_r\). Because \(|l| = |r|\), the fact that the height of an almost complete tree is determined uniquely by its size (Lemma 4.7) implies \(h_l = h_r\) and thus \(\text{acomplete } t\) reduces to \(\text{acomplete } l\), which is again true by assumption.

Note that the proof does not rely on the fact that it is the left child that is potentially one bigger than the right one; it merely requires that the difference in size between two siblings is at most 1.

11.3 Arrays via Braun Trees

In this section we implement arrays by means of Braun trees and verify correctness and complexity. We start by defining array-like functions on Braun trees. After the above explanation of Braun trees the following lookup function will not come as a surprise:

\[
\text{lookup} :: 'a \text{ tree} \Rightarrow \text{nat} \Rightarrow 'a
\]

\[
\text{lookup} \; \langle l, x, r \rangle \; n = (\text{if } n = 1 \text{ then } x \text{ else } \text{lookup} \; (\text{if } \text{even } n \text{ then } l \text{ else } r) \; (n \text{ div } 2))
\]

The least significant bit is the parity of the index and we advance to the next bit by \text{div} 2. The function is called \text{lookup1} rather than \text{lookup} to emphasize that it expects the index to be at least 1. This simplifies the implementation via Braun trees but is in contrast to the \text{Array} interface where by convention indexing starts with 0.

Function \text{update1} descends in the very same manner but also performs an update when reaching 1:

\[
\text{update} :: 'a \text{ nat} \Rightarrow 'a \text{ tree} \Rightarrow 'a \text{ tree}
\]

\[
\text{update} \; x \; \langle \rangle = \langle \rangle, \; x, \; \langle \rangle, \; \text{update} \; n \; x \; \langle l, a, r \rangle = (\text{if } n = 1 \text{ then } \langle l, x, r \rangle
\]
11.3 Arrays via Braun Trees

Figure 11.3 Array implementation via Braun trees

\[
\begin{align*}
\text{lookup} \ (t, \ _) \ n &= \ \text{lookup1} \ t \ (n + 1) \\
\text{update} \ n \ x \ (t, \ m) &= \ (\text{update1} \ (n + 1) \ x \ t, \ m) \\
\text{len} \ (t, \ m) &= \ m \\
\text{array} \ xs &= \ (\text{adds} \ xs \ 0 \ (\), \ |xs|) \\
\end{align*}
\]

else if even \( n \) then \( (\text{update1} \ (n \div 2) \ x \ l, \ a, \ r) \)
else \( (l, \ a, \ \text{update1} \ (n \div 2) \ x \ r) \)

The second equation updates existing entries in case \( n = 1 \). The first equation, however, creates a new entry and thus supports extending the tree. That is, \( \text{update1} \ (|t| + 1) \ x \ t \) extends the tree with a new node \( x \) at index \( |t| + 1 \). Function \( \text{adds} \) iterates this process (again expecting \( |t| + 1 \) as the index) and thus adds a whole list of elements:

\[
\text{adds} :: 'a \:\text{list} \Rightarrow 'a \text{tree} \Rightarrow 'a \text{tree} \\
\text{adds} \ [\] \ _ \ t = t \\
\text{adds} \ (x \# \ xs) \ n \ t = \text{adds} \ xs \ (n + 1) \ (\text{update1} \ (n + 1) \ x \ t)
\]

The implementation of the Array interface in Figure 11.3 is just a thin wrapper around the corresponding functions on Braun trees. An array is represented as a pair of a Braun tree and its size. Note that although \( \text{update1} \) can extend the tree, the specification and implementation of the array \( \text{update} \) function does not support that: \( n \) is expected to be below the length of the array. Flexible arrays are specified and implemented in Section 11.4.

11.3.1 Functional Correctness

The invariant on arrays is obvious:

\[
\text{invar} \ (t, \ l) = (\text{braun} \ t \land l = |t|)
\]

The abstraction function \( \text{list} \) could be defined in the following intuitive way, where \([m..<n]\) is the list of natural numbers from \( m \) up to but excluding \( n \) (see Appendix A):

\[
\text{list} \ t = \text{map} \ (\text{lookup1} \ t) \ [1..<|t| + 1]
\]
Instead we define `list` recursively and derive the above equation later on

\[
list \colon 'a \text{ tree} \Rightarrow 'a \text{ list} \\
list \langle \rangle = [] \\
list \langle l, x, r \rangle = x \# \text{splice} (\text{list} l) (\text{list} r)
\]

This definition is best explained by looking at Figure 11.2. The subtrees with root 2 and 3 will be mapped to the lists \([2, 4, 6, 8, 10, 12, 14]\) and \([1, 3, 5, 7, 9, 11, 13, 15]\). The obvious way to combine these two lists into \([1, 2, 3, \ldots, 15]\) is to splice them:

\[
\text{splice} \colon 'a \text{ list} \Rightarrow 'a \text{ list} \Rightarrow 'a \text{ list} \\
\text{splice} \langle \rangle \ ys = ys \\
\text{splice} (x \# xs) ys = x \# \text{splice} ys xs
\]

Note that because of this reasonably efficient \(O(n \log n)\), see Section 11.3.2) implementation of `list` we can also regard `list` as part of the interface of arrays.

Before we embark on the actual proofs we state a helpful arithmetic truth that is frequently used implicitly below:

\[
\text{braun} \langle l, x, r \rangle \land n \in \{1..|l, x, r|\} \land 1 < n \rightarrow \\
(odd n \rightarrow n \div 2 \in \{1..|r|\}) \land (even n \rightarrow n \div 2 \in \{1..|l|\})
\]

where \(\{m..n\} = \{k \mid m \leq k \land k \leq m\}\).

We will now verify that the implementation in Figure 11.3 of the `Array` interface in Figure 11.1 satisfies the given specification.

We start with proposition \((\text{len})\), the correctness of function `len`. Because of the invariant, \((\text{len})\) follows directly from

\[|\text{list} t| = |t|\]

which is proved by induction. This fact is used implicitly in many proofs below.

The following proposition implies the correctness property \((\text{lookup})\):

\[
\text{braun} t \land i < |t| \rightarrow \text{list} t ! i = \text{lookup}1 t (i + 1) \tag{11.1}
\]

The proof is by induction and uses the following proposition that is also proved by induction:

\[
n < |xs| + |ys| \land |ys| \leq |xs| \land |xs| \leq |ys| + 1 \rightarrow \\
\text{splice} xs ys \land n = (\text{if} \ even n \text{ then} xs \text{ else} ys) ! (n \div 2)
\]

As a corollary to \((11.1)\) we obtain that function `list` can indeed be expressed via `lookup1`:
11.3 Arrays via Braun Trees

11.3 Arrays via Braun Trees

[Some text]

It follows by list extensionality:

\[\text{list} = \text{map} (\text{lookup} (1 \ldots |t| + 1))\]

Let us now verify update as implemented via update1. The following two preservation properties (proved by induction) prove (update-inv):

\[\text{update1} (n \times t) m = (\text{if } n = m \text{ then } x \text{ else } \text{lookup1 } t m)\]

The following property relating lookup1 and update1 is again proved by induction:

\[\text{list} (\text{update1} (n \times t)) = (\text{list } t)[n - 1 := x]\]

Finally we turn to the constructor array. It is implemented in terms of adds and update1. Their correctness is captured by the following properties whose inductive proofs build on each other:

\[\text{list} (\text{add} x \times t) = \text{list } t @ x\]

The last two properties imply the remaining proof obligations (array) and (array-inv). The proof of (11.5) requires the following two properties of splice which are proved by simultaneous induction:

\[|ys| \leq |xs| \rightarrow \text{splice } (xs @ [x]) \text{ys} = \text{splice } xs \text{ys} @ [x]\]

11.3.2 Running Time Analysis

The running time of lookup and update is obviously logarithmic because of the logarithmic height of Braun trees. We sketch why list and array both have running time \(O(n \log n)\). Linear time versions are presented in Section 11.5.

Function list is similar to bottom-up merge sort and splice is similar to merge. We focus on splice because it performs almost all the work. Consider calling list on a complete tree of height \(h\). At each level \(k\) (starting with 0 for the root) of the tree.
ADT Array_Flex = Array +

interface
add_lo :: 'a ⇒ 'ar ⇒ 'ar
del_lo :: 'ar ⇒ 'ar
add_hi :: 'a ⇒ 'ar ⇒ 'ar
del_hi :: 'ar ⇒ 'ar

specification
invar ar → invar (add_lo a ar)  (add_lo-inv)
invar ar → list (add_lo a ar) = a ≠ list ar  (add_lo)
invar ar → invar (del_lo ar)  (del_lo-inv)
invar ar → list (del_lo ar) = tl (list ar)  (del_lo)
invar ar → invar (add_hi a ar)  (add_hi-inv)
invar ar → list (add_hi a ar) = list ar @ [a]  (add_hi)
invar ar → invar (del_hi ar)  (del_hi-inv)
invar ar → list (del_hi ar) = butlast (list ar)  (del_hi)

splice is called \(2^h\) times with lists of size (almost) \(2^{h-k-1}\). The running time of splice with lists of the same length is proportional to the size of the lists. Thus the running time at each level is \(O(2^k 2^{h-k-1}) = O(2^{h-1}) = O(2^h)\). Thus all the splices together require time \(O(h 2^h)\). Because complete trees have size \(n = 2^h\), the bound \(O(n \lg n)\) follows.

Function array is implemented via adds and thus via repeated calls of update1. At the beginning of Section 7.3 we show that because update1 has logarithmic complexity, calling it \(n\) times on a growing tree starting with a leaf takes time \(\Theta(n \lg n)\).

Flexible Arrays
Flexible arrays can be grown and shrunk at either end. Figure 11.4 shows the specification of all four operations. (For tl and butlast see Appendix A.) Array_Flex extends the basis specification Array in Figure 11.1.

Below we first implement the Array_Flex functions on Braun trees. In a final step an implementation of Array_Flex on (tree, size) pairs is derived.

We have already seen that update1 adds an element at the high end. The inverse operation del_hi removes the high end, assuming that the given index is the size of the tree:
11.4 Flexible Arrays

\[
del_hi :: \text{nat} \Rightarrow \text{tree} \\
del_hi \_ \_ = \text{'}\text{tree}' \_ \\
del_hi n (l, x, r) \\
= (\text{if } n = 1 \text{ then } \_ \\
\text{else if } \text{even } n \text{ then } (del_hi \ (n \text{ div } 2) \ l, x, r) \text{ else } (l, x, del_hi \ (n \text{ div } 2) \ r))
\]

This was easy but extending an array at the low end seems hard because one has to shift the existing entries. However, Braun trees support a logarithmic implementation:

\[
add_lo :: \text{a} \Rightarrow \text{tree} \Rightarrow \text{tree} \\
add_lo x \_ = (\_ \_ , x, \_ ) \\
add_lo x (l, a, r) = (\text{add_lo } a \ r, x, l)
\]

The intended functionality is \text{list } (\text{add_lo } x \ t) = x \ # \ \text{list } t. \text{ Function add_lo installs the new element } x \text{ at the root of the tree. Because add_lo needs to shift the indices of the elements already in the tree, the left child (indices 2, 4, \ldots ) becomes the new right child (indices 3, 5, \ldots ). The old right child becomes the new left child with the old root } a \text{ added in at index 2 and the remaining elements at indices 4, 6, \ldots . In the following example, add_lo 0 transforms the left tree into the right one. The numbers in the nodes are the actual elements, not their indices.}

\[
\begin{array}{c}
1 \\
2 3 \\
4 5 6
\end{array}
\quad add_lo 0, \\
\begin{array}{c}
0 \\
1 2 \\
3 5 4 6
\end{array}
\]

Function \text{del_lo} simply reverses \text{add_lo} by removing the root and merging the children:

\[
del_lo :: \text{tree} \Rightarrow \text{tree} \\
del_lo \_ = \_ \\
del_lo (l, \_ , r) = \text{merge } l \ r
\]

\[
merge :: \text{tree} \Rightarrow \text{tree} \Rightarrow \text{tree}
\]
Chapter 11  Arrays via Braun Trees

Figure 11.5  Flexible array implementation via Braun trees

\[
\begin{align*}
  \text{add}_\text{lo} \; x \; (t, l) &= \; (\text{add}_\text{lo} \; x \; t, l + 1) \\
  \text{del}_\text{lo} \; (t, l) &= \; (\text{del}_\text{lo} \; t, l - 1) \\
  \text{add}_\text{hi} \; x \; (t, l) &= \; (\text{update} \; (l + 1) \; x \; t, l + 1) \\
  \text{del}_\text{hi} \; (t, l) &= \; (\text{del}_\text{hi} \; t, l - 1)
\end{align*}
\]

Figure 11.5 shows the obvious implementation of the functions in the \textit{Array\_Flex} specification from Figure 11.4 (on the left-hand side) with the help of the corresponding Braun tree operations (on the right-hand side). It is an extension of the basic array implementation from Figure 11.3. All \textit{Array\_Flex} functions have logarithmic time complexity because the corresponding Braun tree functions do because they descend along one branch of the tree.

11.4.1 Functional Correctness

We now have to prove the properties in Figure 11.4. We have already dealt with \textit{update} and thus \textit{add}_\text{hi} above. Properties (\textit{add}_\text{hi}-inv) and (\textit{add}_\text{hi}) follow from (11.3), (11.4) and (11.5) stated earlier.

Correctness of \textit{del}_\text{hi} on Braun trees is captured by the following two properties proved by induction:

\[
\begin{align*}
  \text{braun} \; t &\rightarrow \text{braun} \; (\text{del}_\text{hi} \; \mid t \; t) \\
  \text{braun} \; t &\rightarrow \text{list} \; (\text{del}_\text{hi} \; \mid t \; t) = \text{butlast} \; (\text{list} \; t)
\end{align*}
\]

They imply (\textit{del}_\text{hi}) and (\textit{del}_\text{hi}-inv). The proof of (11.6) requires the following property of \textit{splice}, which is proved by induction:

\[
\begin{align*}
  \text{butlast} \; (\text{splice} \; xs \; ys) \\
  = (\text{if} \; |ys| < |xs| \; \text{then} \; \text{splice} \; (\text{butlast} \; xs) \; ys \; \text{else} \; \text{splice} \; xs \; (\text{butlast} \; ys))
\end{align*}
\]

Correctness of \textit{add}_\text{lo} on Braun trees (properties (\textit{add}_\text{lo}) and (\textit{add}_\text{lo}-inv)) follows directly from the following two inductive properties:

\[
\begin{align*}
  \text{braun} \; t &\rightarrow \text{list} \; (\text{add}_\text{lo} \; a \; t) = a \; \# \; \text{list} \; t \\
  \text{braun} \; t &\rightarrow \text{braun} \; (\text{add}_\text{lo} \; x \; t)
\end{align*}
\]
Finally we turn to del_lo. Inductions (for merge) and case analyses (for del_lo) yield the following properties:

\[
\begin{align*}
\text{braun } (l, x, r) & \to \text{list } (\text{merge } l \ r) = \text{splice } (\text{list } l) (\text{list } r) \\
\text{braun } (l, x, r) & \to \text{braun } (\text{merge } l \ r) \\
\text{braun } t & \to \text{list } (\text{del_lo } t) = \text{tl } (\text{list } t) \\
\text{braun } t & \to \text{braun } (\text{del_lo } t)
\end{align*}
\]

The last two properties imply (del_lo) and (del_lo-inv).

11.5  
Bigger, Better, Faster, More!

In this section we meet efficient versions of some old and new functions on Braun trees. The implementation of the corresponding array operations is trivial and is not discussed.

11.5.1 Fast Size of Braun Trees

The size of a Braun tree can be computed without having to traverse the entire tree:

\[
\begin{align*}
\text{size_fast } :: & \ 'a \text{ tree } \Rightarrow \text{nat} \\
\text{size_fast } \emptyset & = 0 \\
\text{size_fast } (l, \_, r) & = (\text{let } n = \text{size_fast } r \text{ in } 1 + 2 \cdot n + \text{diff } l \ n)
\end{align*}
\]

\[
\begin{align*}
\text{diff } :: & \ 'a \text{ tree } \Rightarrow \text{nat } \Rightarrow \text{nat} \\
\text{diff } \emptyset & = 0 \\
\text{diff } (l, \_, r) \ n & = (\text{if } n = 0 \text{ then } 1 \\
& \quad \text{else if even } n \text{ then } \text{diff } r \ (n \div 2 - 1) \quad \text{else } \text{diff } l \ (n \div 2))
\end{align*}
\]

Function size_fast descends down the right spine, computes the size of a Node as if both children were the same size \((1 + 2 \cdot n)\), but adds diff \(l \ n\) to compensate for bigger left children. Correctness of size_fast

Lemma 11.3. \(\text{braun } t \to \text{size_fast } t = |t|\)

follows from this property of diff:

\[
\text{braun } t \land |t| \in \{n, n + 1\} \to \text{diff } t \ n = |t| - n
\]

The running time of size_fast is quadratic in the height of the tree (Exercise 11.3).
11.5.2 Initializing a Braun Tree with a Fixed Value

Above we only considered the construction of a Braun tree from a list. Alternatively one may want to create a tree (array) where all elements are initialized to the same value. Of course one can call `update1 n` times, but one can also build the tree directly:

```haskell
braun_of_naive x n
= (if n = 0 then ()
  else let m = (n - 1) div 2
       in if odd n
         then (braun_of_naive x m, x, braun_of_naive x m)
         else (braun_of_naive x (m + 1), x,
               braun_of_naive x m))
```

This solution also has time complexity $O(n \log n)$ but it can clearly be improved by sharing identical recursive calls. Function `braun2_of` shares as much as possible by producing trees of size $n$ and $n + 1$ in parallel:

```haskell
braun2_of :: 'a ⇒ nat ⇒ 'a tree × 'a tree
braun2_of x n
= (if n = 0 then ((()), (()), x, ()))
  else let (s, t) = braun2_of x ((n - 1) div 2)
        in if odd n then ((s, x, s), (t, x, s)) else ((t, x, s), (t, x, t))

braun_of :: 'a ⇒ nat ⇒ 'a tree
braun_of x n = fst (braun2_of x n)
```

The running time is clearly logarithmic.

The correctness properties (see Appendix A for replicate)

```haskell
list (braun_of x n) = replicate n x
braun (braun_of x n)
```

are corollaries of the more general statements

```haskell
braun2_of x n = (s, t) →
list s = replicate n x ∧ list t = replicate (n + 1) x
braun2_of x n = (s, t) →
|s| = n ∧ |t| = n + 1 ∧ braun s ∧ braun t
```

which can both be proved by induction.
11.5 Bigger, Better, Faster, More!

11.5.3 Converting a List into a Braun Tree

We improve on function `adds` from Section 11.3 that has running time $\Theta(n \lg n)$ by developing a linear-time function. Given a list of elements $[1, 2, \ldots]$, we can subdivide it into sublists $[1], [2, 3], [4, \ldots, 7], \ldots$ such that the $k$th sublist contains the elements of level $k$ of the corresponding Braun tree. This is simply because on each level we have the entries whose index has $k + 1$ bits. Thus we need to process the input list in chunks of size $2^k$ to produce the trees on level $k$. But we also need to get the order right. To understand how that works, consider the last two levels of the tree in Figure 11.2:

```
  4     6     5     7
 / \   / \   / \   / \
8   12 10  14  9  13 11 15
```

If we rearrange them in increasing order of the root labels

```
  4     5     6     7
 / \   / \   / \   / \
8   12  9  13 10  14 11 15
```

the following pattern emerges: the left subtrees are labeled $[8, \ldots, 11]$, the right subtrees $[12, \ldots, 15]$. Call $t_i$ the tree with root label $i$. The correct order of subtrees, i.e. $t_4, t_6, t_5, t_7$, is restored when the three lists $[t_4, t_5], [2, 3]$ (the labels above) and $[t_6, t_7]$ are combined into new trees by going through them simultaneously from left to right, yielding $[\langle t_4, 2 \rangle, \langle t_5, 3, t_7 \rangle]$, the level above.

Abstracting from this example we arrive at the following code. Loosely speaking, `brauns k xs` produces the Braun trees on level $k$.

```haskell
brauns :: nat ⇒ 'a list ⇒ 'a tree list
brauns k xs
= (if xs = [] then []
  else let ys = take 2^k xs;
       zs = drop 2^k xs;
       ts = brauns (k + 1) zs
       in nodes ts ys (drop 2^k ts))
```

Function `brauns` chops off a chunk $ys$ of size $2^k$ from the input list and recursively converts the remainder of the list into a list $ts$ of (at most) $2^{k+1}$ trees. This list is (conceptually) split into `take 2^k ts` and `drop 2^k ts` which are combined with $ys$. 
by function nodes that traverses its three argument lists simultaneously. As a local optimization, we pass all of ts rather than just take $2^k$ ts to nodes.

$$
nodes :: 'a tree list \Rightarrow 'a list \Rightarrow 'a tree list \Rightarrow 'a tree list
$$

Because the input list may not have exactly $2^n - 1$ elements, some of the chunks of elements and trees may be shorter than $2^k$. To compensate for that, function nodes implicitly pads lists of trees at the end with leaves. This padding is the purpose of equations two to four.

The top-level function for turning a list into a tree simply extracts the first (and only) element from the list computed by brauns 0:

$$
brauns1 :: 'a list \Rightarrow 'a tree
brauns1 xs = (if xs = [] then () else brauns 0 xs ! 0)
$$

### 11.5.3.1 Functional Correctness

The key correctness lemma below expresses a property of Braun trees: the subtrees on level $k$ consist of all elements of the input list $xs$ that are $2^k$ elements apart, starting from some offset. To state this concisely we define

$$
take_nths :: nat \Rightarrow nat \Rightarrow 'a list \Rightarrow 'a list
take_nths _ _ [] = []
$$

The result of $take_nths i k xs$ is every $2^k$-th element in $drop i xs$.

A number of simple properties follow by easy inductions:

$$
take_nths i k (drop j xs) = take_nths (i + j) k xs \quad (11.7)
take_nths 0 0 xs = xs \quad (11.8)
splice (take_nths 0 1 xs) (take_nths 1 1 xs) = xs \quad (11.9)
$$
take\_nths \(i\) \(m\) (take\_nths \(j\) \(n\) \(xs\))
\(=\) take\_nths \((i \cdot 2^n + j)\) \((m + n)\) \(xs\)  
(11.10)
\(\)take\_nths \(i\) \(k\) \(xs\) = \([] \leftarrow |xs| \leq i\)  
(11.11)
i < |xs| \(\rightarrow\) hd (take\_nths \(i\) \(k\) \(xs\)) = \(xs \downarrow i\)  
(11.12)
\(|xs| = |ys| \lor |xs| = |ys| + 1 \rightarrow\)
take\_nths \(0\) \(1\) (splice \(xs\) \(ys\)) = \(xs \land\)  
take\_nths \(1\) \(1\) (splice \(xs\) \(ys\)) = \(ys\)  
(11.13)
\(|take\_nths \(0\) \(1\) \(xs\)| = |take\_nths \(1\) \(1\) \(xs\)| \(\lor\)  
|take\_nths \(0\) \(1\) \(xs\)| = |take\_nths \(1\) \(1\) \(xs\)| + 1  
(11.14)

We also introduce a predicate relating a tree to a list:

\texttt{braun\_list :: 'a tree \(\Rightarrow\) 'a list \(\Rightarrow\) bool}

\texttt{braun\_list \(\langle\rangle\) \(xs\) = (\(xs = []\))}
\texttt{braun\_list \(\langle l, x, r \rangle\) \(xs\)
\(=\) (\(xs \neq []\) \(\land x = hd \, xs\) \(\land\)
\texttt{braun\_list \(l\) (take\_nths \(1\) \(1\) \(xs\)) \(\land\)
\texttt{braun\_list \(r\) (take\_nths \(2\) \(1\) \(xs\))})}

This definition may look a bit mysterious at first but it satisfies a simple specification:
\texttt{braun\_list \(t\) \(xs\) \(\leftarrow\) braun \(t\) \(\land\) \(xs = list\, t\) (see below). The idea of the above definition}
\texttt{is that instead of relating \(\langle l, x, r \rangle\) to \(xs\) via splice we invert the process and relate}
l and \(r\) to the even and odd numbered elements of \texttt{drop \(1\) \(xs\)}.

\textbf{Lemma 11.4.} \(\texttt{braun\_list \(t\) \(xs\) \(\leftarrow\) braun \(t\) \(\land\) \(xs = list\, t\)\)

\textit{Proof} by induction on \(t\). The base case is trivial. In the induction step the key
properties are (11.14) to prove \(braun\, t\) and (11.9) and (11.13) to prove \(xs = list\, t\). \(\square\)

The correctness proof of \texttt{brauns} rests on a few simple inductive properties:
\(|\texttt{nodes\, \(ls\)\, \(xs\)\, \(rs\)}| = |\(xs|\)  
(11.15)
i < |\(xs|\) \(\rightarrow\)
\texttt{nodes\, \(ls\)\, \(xs\)\, \(rs\) \(\downarrow\) \(i\)
\(=\) \texttt{(if \(i < |\(ls|\) then \(ls\) \(\downarrow\) \(i\) else \(\langle\rangle\), \(xs\) \(\downarrow\) \(i),
\texttt{if \(i < |\(rs|\) then \(rs\) \(\downarrow\) \(i\) else \(\langle\rangle\)\})}  
(11.16)
\(|\texttt{brauns\, \(k\)\, \(xs\)| = \texttt{min\, |\(xs|\, 2^k}\)\)
(11.17)

The main theorem expresses the following correctness property of the elements of
\texttt{brauns \(k\)\, \(xs\): every tree \texttt{brauns \(k\)\, \(xs\) \(\downarrow\) \(i\) is a Braun tree and its list of elements is}
take\_nths \(i\) \(k\) \(xs\):
Chapter 11  Arrays via Braun Trees

Theorem 11.5. \( i < \min |xs| 2^k \mapsto \)
\( \texttt{braun\_list (brauns } k \ x \ ! \ i \ (\texttt{take\_nths } i \ k \ x \) s) \)

Proof by induction on the length of \( xs \). Assume \( i < \min |xs| 2^k \), which implies \( xs \neq [] \). Let \( zs = \text{drop } 2^k \ x \) s. Thus \( |zs| < |xs| \) and therefore the IH applies to \( zs \) and yields
\[
\forall i. \ j. \ j = i + 2^k \land i < \min |zs| 2^k + 1 \mapsto
\texttt{braun\_list (ts } i \ (\texttt{take\_nths } j (k + 1) \ x s))
\]
where \( ts = \texttt{brauns } (k + 1) \ x s \). Let \( ts' = \text{drop } 2^k \ ts \). Below we examine \( \texttt{nodes } ts \_ \ ts' \ ! \ i \) with the help of (11.16). Thus there are four similar cases of which we only discuss one representative one: assume \( i < |ts| \) and \( i \geq |ts'| \).

\[
\texttt{braun\_list (brauns } k \ x s \ ! \ i \ (\texttt{take\_nths } i \ k \ x s) \)
\]
\[
\texttt{braun\_list (ts } i \ (\texttt{take\_nths } (2^k + i) (k + 1) \ x s) \land
\texttt{braun\_list () (\texttt{take\_nths } (2^k + 1 + i) (k + 1) \ x s)
by (11.16), (11.10), (11.11), (11.12) and assumptions}
\]
\[
\texttt{True}
by (*), (11.11), (11.17) and assumptions}
\]

Setting \( i = k = 0 \) in this theorem we obtain the correctness of \( \texttt{brauns1} \) using Lemma 11.4 and (11.8):

Corollary 11.6. \( \texttt{braun (brauns1 x s) \land list (brauns1 x s)} = x s \)

11.5.3.2 Running Time Analysis
Function \( T_{\text{nodes}} \) is shown in Appendix B.4. It is obviously linear:
\[
T_{\text{nodes}} \text{ is x s r s} = |x s| + 1 \quad (11.18)
\]
Function \( T_{\text{brauns}} \) assumes that \( 2^k \) can be computed in constant (i.e. 0) time like all basic arithmetic operations. This is justified if \( k \) is bounded, in which case \( 2^k \) can be implemented as a table lookup.

\[
T_{\text{brauns}} :: \texttt{nat} \Rightarrow \texttt{a list} \Rightarrow \texttt{nat}
\]
\[
T_{\text{brauns}} \ k \ x s
= (\texttt{if } x s = [] \texttt{ then } 0 \texttt{ else let y s = take } 2^k \ x s; \ z s = \texttt{drop } 2^k \ x s; \ t s = \texttt{brauns } (k + 1) \ x s \in T_{\text{take}} 2^k \ x s + T_{\text{drop}} 2^k \ x s + T_{\text{brauns}} (k + 1) \ z s + T_{\text{drop}} 2^k \ t s +
T_{\text{nodes}} \ t s \ y s (\texttt{drop } 2^k \ t s)) +
1
\]
Lemma 11.7. $\text{brauns}_k xs \leq 9 \cdot (|xs| + 1)$

Proof. The proof is by induction on the length of $xs$. If $xs = []$ the claim is trivial. Now assume $xs \neq []$ and let $zs = \text{drop} \ 2^k \ xs$. In the first step we simplify the body using (11.17), (11.18) and simple properties of $\text{take}$, $\text{drop}$, $\text{take}$ and $\text{drop}$ and $\text{min}$:

\[
\begin{align*}
\text{brauns}_k xs &= 3 \cdot (\min 2^k |zs| + 1) + (\min 2^k (|zs| - 2^k) + 1) + \text{brauns}_k (k + 1) zs + 1 \\
&\leq 4 \cdot \min 2^k |zs| + \text{brauns}_k (k + 1) zs + 5 \\
&= 4 \cdot \min 2^k |zs| + 9 \cdot (|zs| + 1) + 5 \\
&= 4 \cdot \min 2^k |zs| + 9 \cdot (|zs| - 2^k + 1) + 5 \\
&= 4 \cdot \min 2^k |zs| + 4 \cdot (|zs| - 2^k) + 5 \cdot (|zs| - 2^k + 1) + 9 \\
&= 4 \cdot |zs| + 5 \cdot (|zs| - 2^k + 1) + 9 \\
&\leq 4 \cdot |zs| + 5 \cdot |zs| + 9 \\
&= 9 \cdot (|zs| + 1)
\end{align*}
\]

because $|zs| - 2^k + 1 \leq |zs|$

11.5.4 Converting a Braun Tree into a List

We improve on function $\text{list}$ that has running time $O(n \lg n)$ by developing a linear-time version. Imagine that we want to invert the computation of $\text{brauns}$ and thus of $\text{brauns}_1$. Thus it is natural to convert not merely a single tree but a list of trees. Looking once more at the reordered list of subtrees

\[
\begin{array}{cccccc}
4 & 5 & 6 & 7 \\
8 & 12 & 9 & 13 & 10 & 14 & 11 & 15
\end{array}
\]

the following strategy strongly suggests itself: first the roots, then the left children, then the right children. The recursive application of this strategy also takes care of the required reordering of the subtrees. Of course we have to ignore any leaves we encounter. This is the resulting function:

\[
\begin{align*}
\text{list\_fast\_rec} :: \ ('a \ \text{tree list}) \Rightarrow \ ('a \ \text{list}) \\
\text{list\_fast\_rec} \ ts \\
= \ (\text{let} \ us = \ \text{filter} \ (\lambda t. \ t \neq ()) \ ts \\
\text{in} \ \text{if} \ us = [] \text{ then } [] \\
\text{else} \ \text{map \ value} \ us \ @ \ \text{list\_fast\_rec} \ (\text{map \ left} \ us \ @ \ \text{map \ right} \ us))
\end{align*}
\]
value \langle l, x, r \rangle = x
left \langle l, x, r \rangle = l
right \langle l, x, r \rangle = r

Function list_fast_rec terminates because left and right remove the top node of a non-\langle \rangle tree. Thus the sum of the sizes of all trees in ts decreases with each recursive call because us is a non-empty list of non-\langle \rangle trees.

This is the top level function to extract a list from a single tree:

\[
\text{list_fast :: 'a tree} \Rightarrow \text{'a list}
\]
\[
\text{list_fast } t = \text{list_fast_rec } [t]
\]

From list_fast one can easily derive an efficient fold function on Braun trees that processes the elements in the tree in the order of their indexes.

11.5.4.1 Functional Correctness

We want to prove correctness of list_fast: list_fast \( t = \text{list } t \) if braun \( t \). A direct proof of list_fast_rec \([t]\) = list \( t \) will fail and we need to generalize this statement to all lists of length \( 2^k \). Reusing the infrastructure from the previous subsection this can be expressed as follows:

**Theorem 11.8.** \(|ts| = 2^k \land (\forall i < 2^k. \text{braun_list } (ts ! i) (\text{take_nths } i \ k \ xs)) \rightarrow \text{list_fast_rec } ts = xs\)

**Proof** by induction on the length of xs. Assume the two premises. There are two cases.

First assume \(|zs| < 2^k\). Then

\[
ts = \text{map } (\lambda x: \langle \rangle, x, \langle \rangle) \ x @ \text{replicate } n \ \langle \rangle
\]

where \( n = |ts| - |zs| \). This can be proved pointwise. Take some \( i < 2^k \). If \( i < |zs| \) then \( \text{take_nths } i \ k \ xs = \text{take } 1 \ (\text{drop } i \ xs) \) (which can be proved by induction on xs).

By definition of braun_list it follows that \( t ! i = \langle l, xs ! i, r \rangle \) for some \( l \) and \( r \) such that \( \text{braun_list } l [] \) and \( \text{braun_list } l [] \) and thus \( l = r = \langle \rangle \), i.e. \( t ! i = \langle \rangle, xs ! i, \langle \rangle \). If \( i < |xs| \) then \( \text{take_nths } i \ k \ xs = [] \) by (11.11) and thus \( \text{braun_list } (ts ! i) [] \) by the second premise and thus \( ts ! i = \langle \rangle \) by definition of braun_list. This concludes the proof of (\(*\)). The desired list_fast_rec \( ts = xs \) follows easily by definition of list_fast_rec.

Now assume \( \neg |zs| < 2^k \). Then for all \( i < 2^k \)
11.5 Bigger, Better, Faster, More!

\( ts \mid i \neq \langle \rangle \land \text{value}(ts \mid i) = xs \mid i \land \\
\text{braun_list}(\text{left}(ts \mid i)) (\text{take_nth}(i + 2^k) (k + 1) xs) \land \\
\text{braun_list}(\text{right}(ts \mid i)) (\text{take_nth}(i + 2 \cdot 2^k) (k + 1) xs) \)

follows from the second premise with the help of (11.10), (11.11) and (11.12). We obtain two consequences:

\[
\text{map value} \ ts = \text{take} \ 2^k \ xs
\]

\[
\text{list_fast_rec} (\text{map left} \ ts \ @ \ \text{map right} \ ts) = \text{drop} \ 2^k \ xs
\]

The first consequence follows by pointwise reasoning, the second consequence with the help of the IH and (11.7). From these two consequences the desired conclusion \( \text{list_fast_rec} \ ts = \ xs \) follows by definition of \( \text{list_fast_rec} \).

11.5.4.2 Running Time Analysis

We focus on \( \text{list_fast_rec} \). After a few simplifications with basic properties of \( \text{map} \) and \( \text{T_append} \) the definition of \( \text{T_list_fast_rec} \) looks like this:

\[
\text{T_list_fast_rec} :: 'a \ \text{tree list} \Rightarrow \text{nat}
\]

\[
\text{T_list_fast_rec} \ ts = \text{(let } \text{us} = \text{filter } (\lambda \ t. \ t \neq \langle \rangle )) \ \text{ts} \text{ in } |ts| + 1 + \\
\text{if } \text{us} = \langle \rangle \text{ then } 0 \text{ else } 5 \cdot (|us| + 1) + \text{T_list_fast_rec} (\text{map left} \ us \ @ \ \text{map right} \ us) \text{) } + 1
\]

The following inductive proposition is an abstraction of the core of the termination argument of \( \text{list_fast_rec} \) above.

\[
(\forall \ t \in \text{set ts}. \ t \neq \langle \rangle ) \rightarrow \sum_{t\in ts} k \cdot |t| = (\sum_{t\in \text{map left ts} \ @ \ \text{map right ts}} k \cdot |t|) + k \cdot |ts| \quad (11.19)
\]

The suggestive notation \( \sum_{x \leftarrow xs. \ f x} \) abbreviates \( \text{sum_list}(\text{map} \ f \ xs) \).

Now we can state and prove a linear upper bound of \( \text{T_list_fast_rec} \):

**Theorem 11.9.** \( \text{T_list_fast_rec} \ ts \leq (\sum_{t\in ts} 14 \cdot |t| + 1) + 2 \)

*Proof* by induction on the size of \( ts \) (which decreases with each recursive call as we argued above). If \( us = \langle \rangle \) the claim is easily seen to be true. Now assume \( us \neq \langle \rangle \) and let \( \text{children} = \text{map left us} \ @ \ \text{map right us} \).

\[
\text{T_list_fast_rec} \ ts = \text{T_list_fast_rec} \ \text{children} + 5 \cdot |us| + |ts| + 7
\]

\[
\leq (\sum_{t\in \text{children}} 14 \cdot |t| + 1) + 5 \cdot |us| + |ts| + 9 \quad \text{by IH}
\]

\[
= (\sum_{t\in \text{children}} 14 \cdot |t| + 7 \cdot |us| + |ts| + 9)
\]
Chapter 11  Arrays via Braun Trees

\[ (\sum_{t \leftarrow \text{children}} 14 \cdot |t|) + 14 \cdot |us| + |ts| + 2 \quad \text{because } us \neq [] \]

\[ = (\sum_{t \leftarrow us} 14 \cdot |t|) + |ts| + 2 \quad \text{by (11.19)} \]

\[ \leq (\sum_{t \leftarrow ts} 14 \cdot |t|) + |ts| + 2 \]

\[ = ((\sum_{t \leftarrow ts} 14 \cdot |t| + 1) + 2) \]

11.6 Experiences

Exercise 11.1. Instead of first showing that Braun trees are almost complete, give a direct proof of \( \text{braun } t \rightarrow h = \lfloor \log |t| \rfloor \) by first showing \( \text{braun } t \rightarrow 2^h t \leq 2 \cdot |t| + 1 \) by induction.

Exercise 11.2. Let \( lh \), the "left height", compute the length of the left spine of a tree. Prove that the left height of a Braun tree is equal to its height: \( \text{braun } t \rightarrow lh = h t \).

Exercise 11.3. Give a readable proof of the fact that Braun trees satisfy the same height as size property:

\[ \text{braun } (l, x, r) \rightarrow h = h \lor h = h + 1 \]

Hint: use the fact that Braun trees are almost complete (and thus height optimal).

Exercise 11.4. Show that function \( \text{bal} \) in Section 4.3.1 produces Braun trees: \( n \leq |zs| \land \text{bal } n zs = (t, zs) \rightarrow \text{braun } t \). (Isabelle hint: \( \text{bal} \) needs to be qualified as \( \text{Balance.bal} \).)

Exercise 11.5. One can view Braun trees as tries (see Chapter 12) by indexing them not with a \( \text{nat} \) but a \( \text{bool list} \) where each bit tells us whether to go left or right (as explained at the start of Section 11.2). Function \( \text{nat.of} \) specifies the intended correspondence:

\[ \text{nat.of} :: \text{bool list} \Rightarrow \text{nat} \]

\[ \text{nat.of} [] = 1 \]

\[ \text{nat.of} (b \# bs) = 2 \cdot \text{nat.of} bs + (\text{if } b \text{ then } 1 \text{ else } 0) \]

Define the counterparts of \( \text{lookup1} \) and \( \text{update1} \)

\[ \text{lookup_trie} :: 'a \Rightarrow \text{bool list} \Rightarrow 'a \]

\[ \text{update_trie} :: \text{bool list} \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a \text{ tree} \]

and prove their correctness:

\[ \text{braun } t \land \text{nat.of} bs \in \{1, |t|\} \rightarrow \text{lookup_trie} t bs = \text{lookup1 } t (\text{nat.of} bs) \]

\[ \text{update_trie} bs x t = \text{update1 } (\text{nat.of} bs) x t \]
Exercise 11.6. Function \texttt{del\_lo} is defined with the help of function \texttt{merge}. Define a recursive function \texttt{del\_lo2 :: tree \Rightarrow tree} without recourse to any auxiliary function and prove \( \texttt{del\_lo2} \ t = \texttt{del\_lo} \ t \).

Exercise 11.7. Prove correctness of function \texttt{braun\_of\_naive} defined in Section 11.5.2: \( \texttt{list} (\texttt{braun\_of\_naive} x n) = \texttt{replicate} n x \).

Exercise 11.8. Show that the running time of \texttt{size\_fast} is quadratic in the height of the tree: Define the running time functions \( T_{\text{diff}} \) and \( T_{\text{size\_fast}} \) (taking 0 time in the base cases) and prove \( T_{\text{size\_fast}} \ t \leq (h t)^2 \).

11.7 Chapter Notes

Braun trees were investigated by \textit{Rem and Braun} [1983] and later, in a functional setting, by \textit{Hoogerwoord} [1992] who coined the term “Braun tree”. Section 11.5 is partly based on work by \textit{Okasaki} [1997]. The whole chapter is based on work by \textit{Nipkow and Sewell} [2020].
A trie is a search tree where keys are strings, i.e. lists of some type of characters. A trie can be viewed as a tree-shaped finite automaton where the root is the start state. For example, the set of strings \{a, an, can, car, cat\} is encoded as the trie in Figure 12.1. The solid states are accepting, i.e. those nodes terminate the string leading to them.

What distinguishes tries from ordinary search trees is that the access time is not logarithmic in the size of the tree but linear in the length of the string, at least assuming that at each node the transition to the sub-trie takes constant time.

12.1 Abstract Tries via Functions

A nicely abstract model of tries is the following type:

```plaintext
datatype 'a trie = Nd bool ('a -> 'a trie)
```

Parameter ‘a is the type of ‘characters’. In a node Nd b f, b indicates if it is an accepting node and f maps characters to sub-tries. Remember (from Section 6.4) that \(\rightarrow\) is a type of maps with update notation \(f(a \mapsto b)\). There is no trie invariant, i.e. the invariant is simply True: there are no ordering, balance or other requirements. This is an abstract model that ignores efficiency considerations like fast access to sub-tries.

Figure 12.2 shows how the ADT Set is implemented by means of tries. The definitions are straightforward. For simplicity, delete does not try to shrink the trie. For example:
empty :: 'a trie
empty = Nd False (λ _ None)

isin :: 'a trie ⇒ 'a list ⇒ bool
isin (Nd b _) [] = b
isin (Nd _ m) (k # xs)
= (case m k of None ⇒ False | Some t ⇒ isin t xs)

insert :: 'a list ⇒ 'a trie ⇒ 'a trie
insert [] (Nd _ m) = Nd True m
insert (x # xs) (Nd b m)
= (let s = case m x of None ⇒ empty | Some t ⇒ t
    in Nd b (m(x ⇒ insert xs s)))

delete :: 'a list ⇒ 'a trie ⇒ 'a trie
delete [] (Nd _ m) = Nd False m
delete (x # xs) (Nd b m)
= Nd b (case m x of None ⇒ m | Some t ⇒ m(x ⇒ delete xs t))

Formally:

delete [a] (Nd False [a ⇒ Nd True (λ _ None)])
= Nd False [a ⇒ Nd False (λ _ None)]

where [x ⇒ t] ≡ (λ _ None)(x ⇒ t). The resulting trie is correct (it represents the empty set of strings) but could have been shrunk to Nd False (λ _ None). We will remedy this "defect" in later, more operational definitions of tries.

12.1.1 Functional Correctness

For the correctness proof we take a lazy approach and define the abstraction function in a trivial manner via isin:
12.2 Binary Tries

A binary trie is a trie over the alphabet `bool`. That is, binary tries represent sets of `bool list`. More concretely, every node has two children:

```
datatype trie = Lf | Nd bool (trie × trie)
```

A binary trie, for example

```
Nd False (Nd True (Nd False (Lf, Lf), Nd True (Lf, Lf)), Lf)
```

can be visualized like this:

```
    False
  /   \
False  True
```

Lfs are not shown at all. The edge labels indicated that `False` refers to the left and `True` to the right child. This convention is encoded in the following auxiliary functions selecting from and modifying pairs:

```
sel2 :: bool ⇒ 'a × 'a ⇒ 'a
sel2 b (a1, a2) = (if b then a2 else a1)

mod2 :: ('a ⇒ 'a) ⇒ bool ⇒ 'a × 'a ⇒ 'a × 'a
```

Correctness of `empty` and `isin (set empty = {} and isin t xs = (xs ∈ set t))` are trivial, correctness of insertion and deletion are easily proved by induction:

```
set (insert xs t) = set t ∪ {xs}
set (delete xs t) = set t − {xs}
```

This simple model of tries leads to simple correctness proofs but is inefficient because of the function space in `'a ⇒ 'a trie`. Now we investigate two efficient implementations: First binary tries where `a` is specialized to `bool`. Then ternary tries, where the maps `a ⇒ 'a trie` are represented by search trees.
empty :: trie
empty = Lf

isin :: trie ⇒ bool list ⇒ bool
isin Lf ks = False
isin (Nd b lr) ks = (case ks of [] ⇒ b | k ≠ ks' ⇒ isin (sel2 k lr) ks')

insert :: bool list ⇒ trie ⇒ trie
insert [] Lf = Nd True (Lf, Lf)
insert [] (Nd _ lr) = Nd True lr
insert (k ≠ ks) Lf = Nd False (mod2 (insert ks) k (Lf, Lf))
insert (k ≠ ks) (Nd b lr) = Nd b (mod2 (insert ks) k lr)

delete :: bool list ⇒ trie ⇒ trie
delete _ Lf = Lf
delete ks (Nd b lr)
  = (case ks of [] ⇒ node False lr
           | k ≠ ks' ⇒ node b (mod2 (delete ks') k lr))

node b lr = (if b ∧ lr = (Lf, Lf) then Lf else Nd b lr)

Figure 12.3 Implementation of Set by binary tries

mod2 f b (a1, a2) = (if b then (a1, f a2) else (f a1, a2))

The implementation of the Set interface is shown in Figure 12.3. In our abstract tries, deletion could generate non-empty sub-tries that do not contain an accepting Nd. In contrast, our binary delete employs a smart constructor node that shrinks a non-accepting Nd to a Lf if both children have become empty. For example delete [True] (Nd False (Lf, Nd True (Lf, Lf))) = Lf.

To ensure that tries are fully shrunk at all times, we make this constraint an invariant: if both sub-tries of a Nd are Lfs, the Nd must be accepting.
invar :: trie ⇒ bool
invar Lf = True
invar (Nd b (l, r)) = (invar l ∧ invar r ∧ (l = Lf ∧ r = Lf → b))

Of course we will need to prove that it is invariant.

12.2.1 Correctness

For the correctness proof we take the same lazy approach as above:

set_trie :: trie ⇒ bool list set
set_trie t = {xs | isin t xs}

The two non-trivial functional correctness properties

set_trie (insert xs t) = set_trie t ∪ {xs}  (12.1)
set_trie (delete xs t) = set_trie t - {xs}  (12.2)

are simple consequences of the following inductive properties:

isin (insert xs t) ys = (xs = ys ∨ isin t ys)
isin (delete xs t) ys = (xs ≠ ys ∧ isin t ys)

The invariant is not required because it only expresses a space optimality property. Preservation of the invariant is easily proved by induction:

invar t → invar (insert xs t)
invar t → invar (delete xs t)

12.2.2 Exercises

Exercise 12.1. Show that distinct tries (which satisfy invar) represent distinct sets:

invar t₁ ∧ invar t₂ → (set_trie t₁ = set_trie t₂) = (t₁ = t₂)

This is in contrast with most BST representations of sets.

Exercise 12.2. Define a union operation union :: trie ⇒ trie ⇒ trie on binary tries and prove set_trie (union t₁ t₂) = set_trie t₁ ∪ set_trie t₂ and invar t₁ ∧ invar t₂ ⇒ invar (union t₁ t₂). Similarly for intersection where you should be able to prove invar (inter t₁ t₂) outright.

Exercise 12.3. This exercise is about searching tries with wildcard patterns, i.e. strings that can contain a special symbol that matches any character. We model such
patterns with type `bool option list` where any Boolean value matches `None` but only `b` matches `Some b`. Define a function `matches :: 'a option list ⇒ 'a list ⇒ bool` that expresses when a wildcard pattern is matched by a `bool list`. Then define a function `isins :: trie ⇒ bool option list ⇒ bool list list` that searches a trie with a wildcard pattern and returns all the `bool list` in the trie that match the pattern. Prove its correctness: `(xs ∈ set (isins t ps)) = (isin t xs ∧ matches ps xs)`.

Exercise 12.4. This exercise is about nearest-neighbour search, namely finding all strings in a trie within a given Hamming distance of the search key. The Hamming distance of two lists of the same length is the number of positions where they differ. Define a function `Hdist :: 'a list ⇒ 'a list ⇒ nat` that computes the Hamming distance. Then define a function `near :: trie ⇒ bool list ⇒ nat ⇒ bool list list` such that `near t xs d` is a list of all `ys` in `t` of the same length as `xs` that have Hamming distance at most `d` from `xs`. Prove its correctness: `(ys ∈ set (near t xs d)) = (|zs| = |ys| ∧ isin t ys ∧ Hdist xs ys ≤ d)`.

12.3 Binary Patricia Tries

Tries can contain long branches without branching. These can be contracted by storing the branch directly in the start node. The result is called a Patricia trie. The following figure shows the contraction of a trie into a Patricia trie:

This is the data type of binary Patricia tries:

```
datatype trieP = LfP | NdP (bool list) bool (trieP × trieP)
```

The implementation of the Set ADT by binary Patricia tries is shown in Figure 12.4; function `nodeP` is displayed separately. The key auxiliary function is `lcp` where `lcp xs ys = (ps, xs', ys')` such that `ps` is the longest common prefix of `xs` and `ys` and `xs'/ys'` is what remains of `xs/ys` after dropping `ps`. Function `lcp` is used by both `insertP` and
deleteP to analyse how the given key and the prefix stored in the NdP overlap. For the detailed case analysis see the code.

Just as for simple binary tries, deletion may enable shrinking. For example, 

\[ NdP \; xs \; False \; (NdP \; ys \; b \; lr, \; LfP) \]

can be shrunk to 

\[ NdP \; (xs \; @ \; False \; \neq \; ys) \; b \; lr \]

because both tries represent the same set. Function deleteP performs shrinking with the help of the smart constructor nodeP that merges two nested NdP's if there is no branching:

\[
\text{nodeP} \; ps \; b \; lr \\
= (\text{if} \; b \; \text{then} \; NdP \; ps \; b \; lr \\
\text{else case} \; lr \; \text{of} \\
\text{LfP} \; \text{LfP} \Rightarrow \text{LfP} \; | \\
(LfP, \; NdP \; ks \; b \; lr) \Rightarrow NdP \; (ps \; @ \; True \; \neq \; ks) \; b \; lr \; | \\
(NdP \; ks \; b \; lr, \; LfP) \Rightarrow NdP \; (ps \; @ \; False \; \neq \; ks) \; b \; lr \; | \\
_ {\Rightarrow} \; NdP \; ps \; b \; lr)
\]

This shrinking property motivates the following invariant: any non-branching NdP must be accepting (because otherwise it could be merged with its children).

\[
\text{invarP} :: \text{trieP} \Rightarrow \text{bool} \\
\text{invarP} \; LfP = \text{True} \\
\text{invarP} \; (NdP \; _ \; b \; (l, \; r)) \\
= (\text{invarP} \; l \wedge \text{invarP} \; r \wedge (l = LfP \vee r = LfP \rightarrow b))
\]

It is tempting to think that \( \text{invarP} \; t = \text{invar} \; (\text{abs_trieP} \; t) \) but this is not the case. Find a \( t \) such that \( \neg \; \text{invarP} \; t \) but \( \text{invar} \; (\text{abs_trieP} \; t) \).

### 12.3.1 Correctness

This is an exercise in stepwise data refinement. We have already proved that trie implements Set via an abstraction function. Now we map trieP back to trie via another abstraction function. Afterwards the overall correctness follows trivially by composing the two abstraction functions.

The abstraction function abs_trieP is defined via the auxiliary function prefix_trie that prefixes a trie with a bit list:

\[
\text{abs_trieP} :: \text{trieP} \Rightarrow \text{trie} \\
\text{abs_trieP} \; LfP = \text{Lf}
\]
emptyP :: trieP
emptyP = LfP

isinP :: trieP ⇒ bool list ⇒ bool
isinP LfP _ = False
isinP (NdP ps b lr) ks
= (let n = |ps|
in if ps = take n ks then case drop n ks of
[[] ⇒ b |
  k # x ⇒ isinP (sel2 k lr) x
else False)

insertP :: bool list ⇒ trieP ⇒ trieP
insertP ks LfP = NdP ks True (LfP, LfP)
insertP ks (NdP ps b lr)
= (case lcp ks ps of
  (_, [], []) ⇒ NdP ps True lr |
  (qs, [], p # ps') ⇒
    let t = NdP ps' b lr
    in NdP qs True (if p then (LfP, t) else (t, LfP)) |
  (_, k # ks', []) ⇒ NdP ps b (mod2 (insertP ks') k lr) |
  (qs, k # ks', _ # ps') ⇒
    let tp = NdP ps' b lr; tk = NdP ks' True (LfP, LfP)
in NdP qs False (if k then (tp, tk) else (tk, tp)))

deleteP :: bool list ⇒ trieP ⇒ trieP
deleteP ks LfP = LfP
deleteP ks (NdP ps b lr)
= (case lcp ks ps of
  (_, [], []) ⇒ nodeP ps False lr |
  (_, _, _ # _) ⇒ NdP ps b lr |
  (_, k # ks', []) ⇒ nodeP ps b (mod2 (deleteP ks') k lr)

lcp :: 'a list ⇒ 'a list ⇒ 'a list × 'a list × 'a list
lcp [] ys = ([], [], ys)
lcp xs [] = ([], xs, [])
lcp (x # xs) (y # ys)
= (if x ≠ y then ([], x # xs, y # ys)
else let (ps, zs', ys') = lcp xs ys in (x # ps, zs', ys'))
abs_trieP \( (\text{NdP} \ ps \ b \ (l, \ r)) \)
\[= \prefix_trie \ ps \ (\text{Nd} \ b \ (abs_trieP \ l, \ abs_trieP \ r))\]

\(\prefix_trie :: \text{bool list} \Rightarrow \text{trie} \Rightarrow \text{trie}\)
\(\prefix_trie \ [] \ t = t\)
\(\prefix_trie \ (k \neq ks) \ t = (\text{let } t' = \prefix_trie \ ks \ t \text{ in Nd False (if } k \text{ then (Lf, } t') \text{ else (}t', \ Lf)))\)

Correctness of \(\text{emptyP}\) is trivial. Correctness of the remaining operations is proved by induction and requires a number of supporting inductive lemmas which we display before the corresponding correctness properties.

Correctness of \(\text{isinP}\):
\(\text{isin} \ (\prefix_trie \ ps \ t) \ ks = (ps = \text{take} \ |ps| \ ks \wedge \text{isin} \ t \ (\text{drop} \ |ps| \ ks))\)

\(\text{isinP} \ t \ ks = \text{isin} \ (abs_trieP \ t) \ ks\)

Correctness of \(\text{insertP}\):
\(\prefix_trie \ ks \ (\text{Nd True} \ (\text{Lf, Lf})) = \text{insert} \ ks \ Lf\)
\(\text{insert} \ ps \ (\prefix_trie \ ps \ (\text{Nd b lr})) = \prefix_trie \ ps \ (\text{Nd True lr})\)
\(\text{insert} \ (ks \oplus ks') \ (\prefix_trie \ ks \ t) = \prefix_trie \ ks \ (\text{insert} \ ks' \ t)\)
\(\prefix_trie \ (ps \oplus qs) \ t = \prefix_trie \ ps \ (\prefix_trie \ qs \ t)\)
\(lcp \ ks \ ps = (qs, ks', ps') \mapsto\)
\(ks = qs \oplus ks' \wedge ps = qs \oplus ps' \wedge (ks' \neq [] \wedge ps' \neq []) \rightarrow \text{hd} \ ks' \neq \text{hd} \ ps'\)
\(\text{abs_trieP} \ (\text{insertP} \ ks \ t) = \text{insert} \ ks \ (abs_trieP \ t)\) \tag{12.3}\)
\(\text{invarP} \ t \mapsto \text{invarP} \ (\text{insertP} \ xs \ t)\)

Correctness of \(\text{deleteP}\):
\(\text{delete} \ xs \ (\prefix_trie \ xs \ (\text{Nd b} \ (l, r))) \)
\[= (\text{if } (l, r) = (\text{Lf, Lf}) \text{ then Lf else prefix_trie} \ xs \ (\text{Nd False} \ (l, r)))\]
\(\text{delete} \ (xs \oplus ys) \ (\prefix_trie \ xs \ t) \)
\[= (\text{if delete} \ ys \ t = \text{Lf} \text{ then Lf else prefix_trie} \ xs \ (\text{delete} \ ys \ t))\]
\(\text{abs_trieP} \ (\text{deleteP} \ ks \ t) = \text{delete} \ ks \ (abs_trieP \ t)\) \tag{12.4}\)
\(\text{invarP} \ t \mapsto \text{invarP} \ (\text{deleteP} \ xs \ t)\)

It is now trivial to obtain the correctness of the \(\text{trieP}\) implementation of sets. The invariant is still \text{invarP} and has already been dealt with. The abstraction function is simply the composition of the two abstraction abstraction functions: \text{set_trieP}
Chapter 12  Tries

\[
\text{set_trie} \circ \text{abs_trieP}. The required functional correctness properties (ignoring emptyP and isinP) are trivial compositions of (12.1)/(12.2) and (12.3)/(12.4):
\]

\[
\text{set_trieP (insertP xs t)} = \text{set_trieP t} \cup \{xs\}
\]

\[
\text{set_trieP (deleteP xs t)} = \text{set_trieP t} - \{xs\}
\]

12.3.2 Exercises

The exercises for binary tries (Section 12.2.2) can be repeated for binary Patricia tries.

12.4 Ternary Tries

What if we want to implement our original abstract tries over type \(\text{\textquoteleft a}\) efficiently, not just binary tries? For example the following one:

Ternary tries implement the \(\text{\textquoteleft a} \rightarrow \text{\textquoteleft a}\) maps as BSTs. The above trie can be represented (non-uniquely) by the following ternary trie:

The ternary trie diagram should be interpreted as follows. The left and right children of a node form the BST. The middle child is the sub-trie that the character in the node maps to. Accepting nodes are gray. The name ternary trie derives from the fact that nodes have three children. However, conceptually they are BSTs that map elements of type \(\text{\textquoteleft a}\) to further such BSTs, i.e. the middle child isn’t really a child but part of the contents of the node.

Using the unbalanced tree implementation of maps from Section 6.5 (any other map implementation works just as well) we define ternary tries as follows:
12.4 Ternary Tries

As before, the bool field indicates if it is an accepting node. Note that trie3 differs from the above graphical display of a ternary trie: in the latter representation, nodes and not transitions are labeled and thus the empty string cannot be represented.

The invariant for ternary tries requires that in all nodes the invariant invar of the map implementation holds:

\[
\text{invar3} :: \text{trie3} \Rightarrow \text{bool} \\
\text{invar3} (\text{Nd}_3 m) = (\text{invar m} \land (\forall a t. \text{lookup m a} = \text{Some t} \rightarrow \text{invar3 t}))
\]

The self-explanatory implementation of the Set interface is shown in Figure 12.5. Function delete does not try to shrink the trie. Remember that lookup and update come from the Map interface.

12.4.1 Functional Correctness

This is another example of stepwise refinement, just like in the correctness proof for binary Patricia tries in Section 12.3. We show that ‘a trie3 implements ‘a trie (from Section 12.1) via this abstraction function:

\[
\text{abs3} :: \text{trie3} \Rightarrow \text{trie} \\
\text{abs3} (\text{Nd}_3 b t) = \text{Nd} b (\lambda a. \text{map_option abs3 (lookup t a)})
\]

\[
\text{map_option} :: (\text{a} \Rightarrow \text{b}) \Rightarrow \text{a option} \Rightarrow \text{b option} \\
\text{map_option f None} = \text{None} \\
\text{map_option f (Some x)} = \text{Some (f x)}
\]

The correctness properties (ignoring empty3) have easy inductive proofs:

\[
\text{isin3} t xs = \text{isin (abs3 t) xs} \\
\text{invar3 t} \rightarrow \text{abs3 (insert3 xs t)} = \text{insert xs (abs3 t)} \\
\text{invar3 t} \rightarrow \text{abs3 (delete3 xs t)} = \text{delete xs (abs3 t)} \\
\text{invar3 t} \rightarrow \text{invar3 (insert3 xs t)} \\
\text{invar3 t} \rightarrow \text{invar3 (delete3 xs t)}
\]
```haskell
empty3 :: 'a trie3
    empty3 = Nd3 False ()

isin3 :: 'a trie3 ⇒ 'a list ⇒ bool
    isin3 (Nd3 b _) [] = b
    isin3 (Nd3 _ m) (x # xs)
        = (case lookup m x of None ⇒ False | Some t ⇒ isin3 t xs)

insert3 :: 'a list ⇒ 'a trie3 ⇒ 'a trie3
    insert3 [] (Nd3 _ m) = Nd3 True m
    insert3 (x # xs) (Nd3 b m)
        = Nd3 b
            (update x
                (insert3 xs (case lookup m x of None ⇒ empty3 | Some t ⇒ t)) m)

delete3 :: 'a list ⇒ 'a trie3 ⇒ 'a trie3
    delete3 [] (Nd3 _ m) = Nd3 False m
    delete3 (x # xs) (Nd3 b m)
        = Nd3 b
            (case lookup m x of None ⇒ m | Some t ⇒ update x (delete3 xs t) m)
```

**Figure 12.5** Implementation of Set via ternary tries

We had already shown that 'a trie implements 'a set and composing the abstraction functions and correctness theorems to show that 'a trie3 implements 'a set is trivial.

### 12.5 Chapter Notes

Tries were first sketched by De La Briandais [1959] and described in more detail by Fredkin [1960] who coined their name based on the word reTRIEval. However, “trie” is usually pronounced like “try” rather than “tree” to avoid confusion. Patricia tries are due to Morrison [1968]. Ternary tries are due to Bentley and Sedgewick [1997].
Part III

Priority Queues
Priority Queues

Tobias Nipkow

A priority queue of linearly ordered elements is like a multiset where one can insert arbitrary elements and remove minimal elements. Its specification as an ADT is shown in Figure 13.1 where Min_mset m \equiv \text{Min}(\text{set_mset} m) and Min yields the minimal element of a finite and non-empty set of linearly ordered elements.

**ADT** Priority.Queue =

**interface**

- empty :: 'q
- insert :: 'a \Rightarrow 'q \Rightarrow 'q
- del_min :: 'q \Rightarrow 'q
- get_min :: 'q \Rightarrow 'a

**abstraction** mset :: 'q \Rightarrow 'a multiset

**invariant** invar :: 'q \Rightarrow bool

**specification**

\[
\begin{align*}
\text{mset empty} & = \{\}\quad \text{(empty)}
\end{align*}
\]

\[
\begin{align*}
\text{invar empty} & \quad \text{(empty-inv)}
\end{align*}
\]

\[
\begin{align*}
\text{invar q} & \rightarrow \text{mset (insert x q)} = \text{mset q} + \{x\} \quad \text{(insert)}
\end{align*}
\]

\[
\begin{align*}
\text{invar q} & \rightarrow \text{invar (insert x q)} \quad \text{(insert-inv)}
\end{align*}
\]

\[
\begin{align*}
\text{invar q} & \wedge \text{mset q} \neq \{\}
\end{align*}
\]

\[
\begin{align*}
\rightarrow \text{mset (del_min q)} & = \text{mset q} - \{\text{get_min q}\} \quad \text{(del_min)}
\end{align*}
\]

\[
\begin{align*}
\text{invar q} & \wedge \text{mset q} \neq \{\} \rightarrow \text{invar (del_min q)} \quad \text{(del_min-inv)}
\end{align*}
\]

\[
\begin{align*}
\text{invar q} & \wedge \text{mset q} \neq \{\} \rightarrow \text{get_min q} = \text{Min_mset (mset q)} \quad \text{(get_min)}
\end{align*}
\]

**Figure 13.1** ADT Priority.Queue

Mergeable priority queues (see Figure 13.2) provide an additional function merge (sometimes: meld or union) with the obvious functionality.

Our priority queues are simplified. The more general version contains elements that are pairs of some item and its priority.
Chapter 13  
Priority Queues

ADT Priority_Queue_Merge = Priority_Queue +

interface
merge :: 'q ⇒ 'q ⇒ 'q

specification
invar q1 ∧ invar q2 → mset (merge q1 q2) = mset q1 + mset q2
invar q1 ∧ invar q2 → invar (merge q1 q2)

Figure 13.2  ADT Priority_Queue_Merge

Exercise 13.1. Give a list-based implementation of mergeable priority queues with constant-time get_min and del_min. Verify the correctness of your implementation w.r.t. Priority_Queue_Merge.

13.1  Heaps

A popular implementation technique for priority queues are heaps, i.e. trees where the minimal element in each subtree is at the root:

heap :: 'a tree ⇒ bool
heap () = True
heap (l, m, r)
  = (∀x ∈ set_tree l ∪ set_tree r. m ≤ x) ∧ heap l ∧ heap r

Function mset_tree extracts the multiset of elements from a tree:

mset_tree :: 'a tree ⇒ 'a multiset
mset_tree () = {}
mset_tree (l, a, r) = {a} + mset_tree l + mset_tree r

When verifying a heap-based implementation of priority queues the invariant invar and the abstraction function mset in the ADT Priority_Queue are instantiated by heap and mset_tree. The correctness proofs need to talk about both multisets and (because of the heap invariant) sets of elements in a heap. We will only show the relevant multiset properties because the set properties follow easily via the fact set_mset (mset_tree t) = set_tree t.

Both empty and get_min have obvious implementations:
empty = ⟨⟩

get_min ⟨_, a, _⟩ = a

If a heap-based implementation provides a merge function (e.g. skew heaps in Chapter 21), then insert and del_min can be defined like this:

\[
\text{insert } x \langle \rangle = \text{merge } \langle \rangle, x, \langle \rangle \langle \rangle
\]

\[
\text{del_min } \langle \rangle = \langle \rangle
\]

\[
\text{del_min } \langle l, _, r \rangle = \text{merge } l \ r
\]

Note that the following tempting definition of merge is functionally correct but leads to very unbalanced heaps:

\[
\text{merge } \langle \rangle t = t
\]

\[
\text{merge } t \langle \rangle = t
\]

\[
\text{merge } \langle l_1, a_1, r_1 \rangle : : t_1 \langle l_2, a_2, r_2 \rangle : : t_2 = (\text{if } a_1 \leq a_2 \text{ then } \langle l_1, a_1, \text{merge } r_1 t_2 \rangle \text{ else } \langle l_2, a_2, \text{merge } t_1 r_2 \rangle)
\]

Many of the more advanced implementations of heaps focus on improving this merge function. We will see examples of this in the next chapter on leftist heaps, as well as in the chapters on skew heaps and pairing heaps.

Exercise 13.2. Show functional correctness of the above definition of merge (w.r.t. Priority_Queue.Merge) and prove functional correctness of the implementations of insert and del_min (w.r.t. Priority_Queue).

Exercise 13.3. Define a function list from a heap to a sorted list of its elements and prove mset (list t) = mset_tree t and heap t → sorted (list t).

13.2 Chapter Notes

The idea of the heap goes back to Williams [1964] who also coined the name. In imperative implementations, priority queues frequently also provide an operation decrease_key: given some direct reference to an element in the priority queue, decrease its element’s priority. This is not completely straightforward in a functional language. Lammich and Nipkow [2019] present an implementation, a Priority Search Tree.
Leftist Heaps

Tobias Nipkow

Leftist heaps are heaps in the sense of Section 13.1 and implement mergeable priority queues. The key idea is to maintain the invariant that at each node the minimal height of the right child is \( \leq \) that of the left child. We represent leftist heaps as augmented trees that store the minimal height in every node:

\[
\text{type synonym} \; 'a \text{ lheap} = ('a \times \text{nat}) \text{ tree}
\]

\[
mht :: 'a \text{ lheap} \Rightarrow \text{nat}
mht () = 0
mht (_, (_, n), _) = n
\]

There are two invariants: the standard heap invariant (on augmented trees)

\[
\text{heap} :: ('a \times 'b) \text{ tree} \Rightarrow \text{bool}
\text{heap} () = \text{True}
\text{heap} (l, (m, _), r)
= ((\forall x \in \text{set_tree} \, l \cup \text{set_tree} \, r. \, m \leq x) \land \text{heap} \, l \land \text{heap} \, r)
\]

and the structural invariant that requires that the minimal height of the right child is no bigger than that of the left child (and that the minimal height information in the node is correct):

\[
\text{ltree} :: 'a \text{ lheap} \Rightarrow \text{bool}
\text{ltree} () = \text{True}
\text{ltree} (l, (_, n), r) = (m \text{h} \, r \leq m \text{h} \, l \land n = m \text{h} \, r + 1 \land \text{ltree} \, l \land \text{ltree} \, r)
\]

Thus a tree is a leftist tree if for every subtree the right spine is a shortest path from the root to a leaf. Pictorially:
Now remember \(2^{mh} t \leq |t|_1\), i.e. \(mh t \leq \lg |t|_1\). Because the expensive operations on leftist heaps descend along the right spine, this means that their running time is logarithmic in the size of the heap.

Exercise 14.1. An alternative definition of leftist tree is via the length of the right spine of the tree:

\[
\begin{align*}
\text{rank} &:: \forall t. \text{tree} \Rightarrow \text{nat} \\
\text{rank} \langle \rangle &= 0 \\
\text{rank} \langle \_, \_, r \rangle &= \text{rank} r + 1
\end{align*}
\]

Prove that \(\text{ltree} t \rightarrow \text{rank} t = mh t\).

Exercise 14.2. Define \(\text{ltree0} :: \forall t. \text{tree} \Rightarrow \text{bool}\), a pared-down version of \(\text{ltree}\) that works on arbitrary trees without any height information stored in the nodes. Thus \(\text{ltree} t \rightarrow \text{ltree0} (\text{map_tree} \text{fst} t)\). Prove that any complete tree is a leftist tree: \(\text{complete} t \rightarrow \text{ltree0} t\).

14.1 Implementation of ADT Priority_Queue_Merge

The key operation is \(\text{merge}\):

\[
\begin{align*}
\text{merge} &:: \forall t. \text{lheap} \Rightarrow \forall t. \text{lheap} \Rightarrow \forall t. \text{lheap} \\
\text{merge} \langle \rangle t &= t \\
\text{merge} t \langle \rangle &= t \\
\text{merge} \langle l_1, (a_1, n_1), r_1 \rangle \langle l_2, (a_2, n_2), r_2 \rangle &= \begin{cases} t_1 & \text{if } a_1 \leq a_2 \\ t_2 & \text{else} \end{cases} \\
&= (\text{if } a_1 \leq a_2 \text{ then } \text{node} l_1 a_1 (\text{merge} r_1 t_2) \\
&\quad \text{else } \text{node} l_2 a_2 (\text{merge} t_1 r_2))
\end{align*}
\]

\[
\begin{align*}
\text{node} &:: \forall t. \text{lheap} \Rightarrow \forall t. \Rightarrow \forall t. \text{lheap} \\
\text{node} l a r &= \begin{cases} (\text{let } mhl = \text{mht} l; mhr = \text{mht} r \\ \text{in } \begin{cases} l_1, (a, mhr + 1), r & \text{if } mhr \leq mhl \\ r, (a, mhl + 1), l & \text{else} \end{cases} & \text{if } a_1 \leq a_2 \text{ then } \text{node} l_1 a_1 (\text{merge} r_1 t_2) \\
&\quad \text{else } \text{node} l_2 a_2 (\text{merge} t_1 r_2))
\end{cases}
\end{align*}
\]

Termination of \(\text{merge}\) can be proved either by the sum of the sizes of the two arguments (which goes down with every call) or by the lexicographic product of the
two size measures: either the first argument becomes smaller or it stays unchanged and the second argument becomes smaller.

As shown in Section 13.1, once we have merge, the other operations are easily definable. We repeat the definitions of those operations that change because this chapter employs augmented rather than ordinary trees:

\[
\begin{align*}
\text{get\_min} &:: \text{'a lheap} \Rightarrow \text{'a} \\
\text{get\_min} \ _\ _, \ (a, \ _), \ _) &= a \\
\text{insert} &:: \text{'a} \Rightarrow \text{'a lheap} \Rightarrow \text{'a lheap} \\
\text{insert} \ x \ t &= \text{merge} \ \langle\langle\rangle, \ (x, 1), \ \langle\rangle\rangle \ t
\end{align*}
\]

14.2 Correctness

The above implementation is proved correct with respect to the ADT Priority\_Queue\_Merge where

\[
\begin{align*}
\text{mset\_tree} &:: \ (\text{'a} \times \text{'b}) \ \text{tree} \Rightarrow \text{'a multiset} \\
\text{mset\_tree} \ () &= \{\} \\
\text{mset\_tree} \ l, \ (a, \ _), \ r) &= \{a\} + \text{mset\_tree} \ l + \text{mset\_tree} \ r \\
\text{invar} \ t &= \ (\text{heap} \ t \land \text{ltree} \ t)
\end{align*}
\]

Correctness of get\_min follows directly from the heap invariant:

\[
\text{heap} \ t \land \ t \neq () \longrightarrow \text{get\_min} \ t = \text{Min} \ (\text{set\_tree} \ t)
\]

From the following inductive lemmas about merge

\[
\begin{align*}
\text{mset\_tree} \ (\text{merge} \ t_1 \ t_2) &= \text{mset\_tree} \ t_1 + \text{mset\_tree} \ t_2 \\
\text{ltree} \ l \land \text{ltree} \ r &\longrightarrow \text{ltree} \ (\text{merge} \ l \ r) \\
\text{heap} \ l \land \text{heap} \ r &\longrightarrow \text{heap} \ (\text{merge} \ l \ r)
\end{align*}
\]

correctness of insert and del\_min follow easily:

\[
\begin{align*}
\text{mset\_tree} \ (\text{insert} \ x \ t) &= \text{mset\_tree} \ t + \{x\} \\
\text{mset\_tree} \ (\text{del\_min} \ t) &= \text{mset\_tree} \ t - \{\text{get\_min} \ t\} \\
\text{ltree} \ t &\longrightarrow \text{ltree} \ (\text{insert} \ x \ t) \\
\text{heap} \ t &\longrightarrow \text{heap} \ (\text{insert} \ x \ t) \\
\text{ltree} \ t &\longrightarrow \text{ltree} \ (\text{del\_min} \ t)
\end{align*}
\]
heap \( t \rightarrow \text{heap (del\_min } t) \)

Of course the above proof (ignoring the ltree part) works for any mergeable priority queue implemented as a heap.

### 14.3 Running Time Analysis

We simplify matters by counting only calls of the only recursive function, \textit{merge}. The running time functions are shown in Appendix B.5. By induction on the computation of \textit{merge} we obtain

\[
\text{ltree } l \land \text{ltree } r \rightarrow T_{\text{merge}} l r \leq mh l + mh r + 1
\]

With \(2^{mh} t \leq |t|_1\) it follows that

\[
\text{ltree } l \land \text{ltree } r \rightarrow T_{\text{merge}} l r \leq \lg |l|_1 + \lg |r|_1 + 1
\]  \hspace{1cm} \text{(14.1)}

which implies logarithmic bounds for insertion and deletion:

\[
\text{ltree } t \rightarrow T_{\text{insert}} x t \leq \lg |t|_1 + 2
\]

\[
\text{ltree } t \rightarrow T_{\text{del\_min}} t \leq 2 \cdot \lg |t|_1
\]

The derivation of the bound for insertion is trivial, as is the proof of the \(T_{\text{del\_min}}\) bound for \(t = \langle\rangle\). The case \(t = \langle l, \_, r\rangle\) and \text{ltree } \(t\) needs a little lemma:

\[
T_{\text{del\_min}} t = T_{\text{merge}} l r
\]

\[
\leq \lg |l|_1 + \lg |r|_1 + 1
\]

\[
\leq 2 \cdot \lg |t|_1
\]

using (14.1) because \(\lg x + \lg y + 1 < 2 \cdot \lg (x + y)\) if \(0 < x\) and \(0 < y\)

### 14.4 Converting a List into a Leftist Heap

We follow the pattern of bottom-up merge sort (Section 2.5) and of the conversions from lists to 2-3 trees (Section 7.3). In both cases we repeatedly pass over a list of objects, merging pairs of adjacent objects in each pass. However, the complexity differs: in merge sort, each merge takes linear time, which leads to the overall complexity of \(O(n \lg n)\); when converting a list into a 2-3 tree, each combination of two trees takes only constant time, which leads to a linear overall complexity. So what happens if the merge step takes logarithmic time, as in (14.1)? But first the algorithm, which is very similar to merge sort:

\[
\text{merge\_adj :: 'a heap list } \Rightarrow \text{'a heap list}
\]

\[
\text{merge\_adj } [] = []
\]

\[
\text{merge\_adj } [t] = [t]
\]
14.4 Converting a List into a Leftist Heap

merge_adj \( (t_1 \# t_2 \# ts) = \text{merge} t_1 t_2 \# \text{merge_adj} ts \)

merge_all :: 'a lheap list ⇒ 'a lheap
merge_all [] = ⟨⟩
merge_all [t] = t
merge_all ts = merge_all (merge_adj ts)

lheap_list :: 'a list ⇒ 'a lheap
lheap_list xs = merge_all (map (λx. ⟨⟩, (x, 1), ⟨⟩) xs)

Termination of merge_all follows because merge_adj decreases the length of the list if \( |ts| \geq 2 \):

\[ |\text{merge_adj} ts| = (|ts| + 1) \text{ div } 2 \]

Functional correctness is straightforward: from the inductive properties

\[ (\forall t \in \text{set ts. heap } t) \rightarrow (\forall t \in \text{set (merge_adj ts). heap } t) \]
\[ (\forall t \in \text{set ts. heap } t) \rightarrow \text{heap (merge_all ts)} \]
\[ (\forall t \in \text{set ts. ltree } t) \rightarrow (\forall t \in \text{set (merge_adj ts). ltree } t) \]
\[ (\forall t \in \text{set ts. ltree } t) \rightarrow \text{ltree (merge_all ts)} \]
\[ \sum_e (\text{image_mset mset_tree (mset (merge_adj ts))}) \]
\[ = \sum_e (\text{image_mset mset_tree (mset ts)}) \]
\[ \text{mset_tree (merge_all ts)} = \sum_e (\text{mset (map mset_tree ts)}) \]

it follows directly that lheap_list xs yields a leftist heap with the same multiset of elements as in xs:

\[ \text{heap (lheap_list ts)} \quad \text{ltree (lheap_list ts)} \]
\[ \text{mset_tree (lheap_list xs)} = \text{mset xs} \]

The running time analysis is more interesting. We only count the time for merge to keep things simple.

\[ T_{\text{merge_adj}} :: \text{'a lheap list ⇒ nat} \]
\[ T_{\text{merge_adj}} [] = 0 \]
\[ T_{\text{merge_adj}} [\_] = 0 \]
\[ T_{\text{merge_adj}} (t_1 \# t_2 \# ts) = T_{\text{merge}} t_1 t_2 + T_{\text{merge_adj}} ts \]

The remaining time functions are displayed in Appendix B.5.
To simplify things further we assume that the length of the initial list \(xs\) and thus the length of all intermediate lists of heaps are powers of 2 and in any of the intermediate lists all heaps have the same size.

Because the complexity of \(\text{merge}\) is logarithmic in the size of the two heaps (14.1), the following upper bound for \(\text{merge\_adj}\) follows by an easy computation induction:

\[
(\forall t \in \text{set ts. ltree } t) \land (\forall t \in \text{set ts. } |t| = n) \longrightarrow \\
T_{\text{merge\_adj}} ts \leq (|ts| \text{ div } 2) \cdot Tm n
\]

where \(Tm n \equiv 2 \cdot \lfloor \log_2 (n + 1) \rfloor + 1\).

The complexity of \(\text{merge\_all}\) can be expressed as a sum:

\[
(\forall t \in \text{set ts. ltree } t) \land (\forall t \in \text{set ts. } |t| = n) \land |ts| = 2^k \longrightarrow \\
T_{\text{merge\_all}} ts \leq (\sum_{i=1}^{k} 2^{k-i} \cdot Tm (2^i - 1 \cdot n))
\]

Each summand is the complexity of one \(\text{merge\_adj}\) call on heap lists whose lengths go down from \(2^k\) to 2 and whose heaps go up in size from \(n\) to \(2^k - 1 \cdot n\). The proof is by induction on the computation of \(\text{merge\_all}\).

The following lemma will permit us to find a closed upper bound for the sum in (14.2). The proof is a straightforward induction on \(k\).

**Lemma 14.1.** \((\sum_{i=1}^{k} 2^{k-i} \cdot (2 \cdot i + 1)) = 5 \cdot 2^k - 2 \cdot k - 5\)

Now we can upper-bound \(T_{\text{heap\_list}}\) as follows if \(|xs| = 2^k\):

\[
T_{\text{heap\_list}} xs = T_{\text{merge\_all}} (\text{map } (\lambda x. (\langle x, 1 \rangle, \langle x \rangle)) xs) \\
\leq \sum_{i=1}^{k} 2^{k-i} \cdot Tm (2^i - 1) \quad \text{(by (14.2)) (where } n = 1) \text{ and } |xs| = 2^k \\
\leq \sum_{i=1}^{k} 2^{k-i} \cdot (2 \cdot \log (2 \cdot 2^i - 1) + 1) \\
= \sum_{i=1}^{k} 2^{k-i} \cdot (2 \cdot i + 1) \\
= 5 \cdot 2^k - 2 \cdot k - 5 \quad \text{by Lemma 14.1}
\]

Thus (14.2) implies that \(T_{\text{heap\_list}} xs\) is upper-bounded by a function linear in \(|xs|\):

\[
|xs| = 2^k \longrightarrow T_{\text{heap\_list}} xs \leq 5 \cdot |xs|
\]

The assumption \(|xs| = 2^k\) merely simplifies technicalities. With more care one can show that \(T_{\text{heap\_list}} \in O(n)\) holds for all inputs of length \(n\).

Finally note that the above complexity analysis has nothing to do with leftist heaps or priority queues and works for any \(\text{merge}\) function of the given logarithmic complexity. Our proofs generalize easily. One can even go one step further and show that \(\text{merge\_all}\) has linear complexity as long as \(\text{merge}\) has sublinear complexity. This is a special case of the master theorem [Cormen et al. 2009] for divide-and-conquer algorithms, because \(\text{merge\_all}\) is just divide-and-conquer in reverse. However, proving
even this special case (let alone the full master theorem) is much harder than the proofs above.

14.5 Chapter Notes

Leftist heaps were invented by Crane [1972]. Another version of leftist trees, based on weight rather than height, was introduced by Cho and Sahni [1998].
15 Priority Queues via Braun Trees

Tobias Nipkow

In Chapter 11 we introduced Braun trees and showed how to implement arrays. In
the current chapter we show how to implement priority queues by means of Braun
trees. Because Braun trees have logarithmic height this guarantees logarithmic running
times for insertion and deletion. Remember that every node \( (l, x, r) \) in a Braun tree
satisfies \( |l| = |r| \lor |l| = |r| + 1 \) (*).

15.1 Implementation of ADT Priority Queue

We follow the heap approach in Section 13.1. Functions empty, get_min, heap and
mset_tree are defined as in that section.

Insertion and deletion maintain the Braun tree property (*) by inserting into the
right (and possibly smaller) child, deleting from the left (and possibly larger) child,
and swapping children to reestablish (*).

Insertion is straightforward and clearly maintains both the heap and the Braun
tree property:

\[
\begin{align*}
\text{insert} & : 'a \Rightarrow 'a \text{ tree} \Rightarrow 'a \text{ tree} \\
\text{insert} \ a \ \langle \rangle &= \langle \rangle, \ a, \langle \rangle \\
\text{insert} \ a \ \langle l, x, r \rangle &= \begin{cases} \\
n\text{insert} \ r \ a, \ l & \text{if } a < x \\
\text{insert} \ a \ x, \ l & \text{else}
\end{cases}
\end{align*}
\]

To delete the minimal (i.e. root) element from a tree, extract the leftmost element
from the tree and let it sift down to its correct position in the tree in the manner of
heapsort:

\[
\begin{align*}
\text{del_min} & : 'a \text{ tree} \Rightarrow 'a \text{ tree} \\
\text{del_min} \ \langle \rangle &= \langle \rangle \\
\text{del_min} \ \langle \rangle, \ x, \ r &= \langle \rangle
\end{align*}
\]
15.2 Correctness

We outline the correctness proofs for \texttt{insert} and \texttt{del\textunderscore min} by presenting the key lemmas. Correctness of \texttt{insert} is straightforward:

\[
|\text{insert } x \ t| = |t| + 1 \\
\text{mset\textunderscore tree } (\text{insert } x \ t) = \{x\} + \text{mset\textunderscore tree } t \\
\text{braun } t \rightarrow \text{braun } (\text{insert } x \ t) \\
\text{heap } t \rightarrow \text{heap } (\text{insert } x \ t)
\]

Correctness of \texttt{del\textunderscore min} builds on analogous correctness lemmas for the auxiliary functions:

\[
\begin{align*}
\text{del\textunderscore left } t &= (x, t') \land t \neq () \rightarrow \text{mset\textunderscore tree } t = \{x\} + \text{mset\textunderscore tree } t' \\
\text{del\textunderscore left } t &= (x, t') \land t \neq () \land \text{heap } t \rightarrow \text{heap } t' \\
\text{del\textunderscore left } t &= (x, t') \land t \neq () \rightarrow |t| = |t'| + 1
\end{align*}
\] (15.1)
15.3 Running Time Analysis

The running time functions are shown in Appendix B.6. Intuitively, all operations are linear in the height of the tree, which in turn is logarithmic in the number of elements (see Section 11.2).

Upper bounds for the running times of \textit{insert}, \textit{del_left} and \textit{sift_down} are proved by straightforward inductions:

\[ T_{\text{insert}} a t \leq h t + 1 \]

\[ t \neq \langle \rangle \rightarrow T_{\text{del_left}} t \leq h t \] (15.3)

\[ \text{braun} (l, a, r) \rightarrow T_{\text{sift_down}} l x r \leq \max (h l) (h r) + 1 \] (15.4)

The analysis of \textit{del_min} requires a bit more work, including another auxiliary inductive fact:

\[ \text{del_left} t = (x, t') \land t \neq \langle \rangle \rightarrow h t' \leq h t \] (15.5)

Lemma 15.1. \( \text{braun} t \rightarrow T_{\text{del_min}} t \leq 2 \cdot h t \)

Proof by induction on \( t \). The base case is trivial. If \( t = \langle l, x, r \rangle \), the case \( l = \langle \rangle \) is again trivial. Assume \( l \neq \langle \rangle \). The call of \textit{del_min} must yield a pair: \textit{del_left} \( l = (y, l') \). Now we are ready for the main derivation:

\[ T_{\text{del_min}} t = T_{\text{del_left}} l + T_{\text{sift_down}} r y l' \]

\[ \leq \text{height} l + T_{\text{sift_down}} r y l' \quad \text{by (15.3)} \]

In order to upper-bound \( T_{\text{sift_down}} r y l' \) via (15.4) we need \( \text{braun} \langle r, y, l' \rangle \), which follows from \( \text{braun} t \) via (15.2) and (15.1). Thus

\[ \leq h l + \max (h r) (h l') + 1 \]

\[ \leq h l + \max (h r) (h l) + 1 \quad \text{by (15.5)} \]

\[ \leq 2 \cdot \max (h l) (h r) + 1 \leq 2 \cdot h t + 1 \]

\[ \square \]
15.4 Chapter Notes
Our implementation of priority queues via Braun trees is due to Paulson [1996] who credits it to Okasaki.
Binomial Heaps

Peter Lammich

Binomial heaps are another common implementation of mergeable priority queues, which supports efficient \(O(\log n)\) insert, get_min, del_min, and merge operations.

The basic building blocks of a binomial heap are binomial trees, which are defined recursively as follows: a binomial tree of rank \(r\) is a node with \(r\) children that are binomial trees of ranks \(r - 1, \ldots, 0\), in that order. Figure 16.1 shows an example binomial tree. It can be shown that a binomial tree of rank \(r\) has \(\binom{r}{l}\) nodes on level \(l\) (see Exercise 16.1). Hence the name.

![Figure 16.1](image)

A binomial tree of rank 3. The node labels depict the rank of each node. A node of rank \(r\) has child nodes of ranks \(r - 1, \ldots, 0\).

To define binomial trees, we first define a more general datatype and the usual syntax for nodes:

```plaintext
datatype 'a tree = Node nat 'a ('a tree list)
<r, x, ts> ≡ Node r x ts
```

Apart from the list of children, a node stores a rank and a root element:

```
rank <r, x, ts> = r    root <r, x, ts> = x
```
Chapter 16  Binomial Heaps

This datatype contains all binomial trees, but also some non-binomial trees. To carve out the binomial trees, we define an invariant, which reflects the informal definition above:

\[
\text{btree} :: 'a \text{ tree} \Rightarrow \text{ bool} \\
\text{btree} \, (r, \_, ts) = ((\forall t \in \text{ set } ts. \text{ btree} \, t) \land \text{ map rank } ts = \text{ rev } [0..<r])
\]

Additionally, we require the heap property, i.e., that the root element of each subtree is a minimal element in that subtree:

\[
\text{heap} :: 'a \text{ tree} \Rightarrow \text{ bool} \\
\text{heap} \, (_, x, ts) = (\forall t \in \text{ set } ts. \text{ heap} \, t \land x \leq \text{ root } t)
\]

Thus, a binomial tree is a tree that satisfies both the structural and the heap invariant. The two invariants are combined in a single predicate:

\[
\text{bheap} :: 'a \text{ tree} \Rightarrow \text{ bool} \\
\text{bheap} \, t = (\text{btree} \, t \land \text{heap} \, t)
\]

A binomial heap is a list of binomial trees

\[
\text{type_synonym} \ 'a \text{ trees} = 'a \text{ tree list}
\]

with strictly ascending rank:

\[
\text{invar} :: 'a \text{ trees} \Rightarrow \text{ bool} \\
\text{invar} \, ts = ((\forall t \in \text{ set } ts. \text{ bheap} \, t) \land \text{ sorted_wrt } (\leq) \, \text{ (map rank } ts))
\]

Note that \text{sorted_wrt} states that a list is sorted w.r.t. the specified relation, here \text{ (\leq)}. It is defined in Appendix A.

16.1  Size

The following functions return the multiset of elements in a binomial tree and in a binomial heap:
16.2 Implementation of ADT Priority Queue

Most operations on binomial heaps are linear in the length of the heap. To show that the length is bounded by the number of heap elements, we first observe that the number of elements in a binomial tree is already determined by its rank. A binomial tree of rank $r$ has $2^r$ nodes:

$$btree t \rightarrow |mset\_tree t| = 2^{\text{rank } t}$$

This proposition is proved by induction on the tree structure. A tree of rank 0 has one element, and a tree of rank $r+1$ has subtrees of rank 0, 1, \ldots, $r$. By the induction hypothesis, these have $2^0, 2^1, \ldots, 2^r$ elements, i.e., $2^{r+1} - 1$ elements together. Including the element at the root, there are $2^{r+1}$ elements.

The length of a binomial heap is bounded logarithmically in the number of its elements:

$$\text{invar } ts \rightarrow |ts| \leq \lg (|mset\_trees ts| + 1) \tag{16.1}$$

To prove this, recall that the heap $ts$ is strictly sorted by rank. Thus, we can underestimate the ranks of the trees in $ts$ by 0, 1, \ldots, $|ts| - 1$. This means that they must have at least $2^0, 2^1, \ldots, 2^{[|ts|]-1}$ elements, i.e., at least $2^{[|ts|]} - 1$ elements together, which yields the desired bound.

16.2 Implementation of ADT Priority Queue

Obviously, the empty binomial heap is [] and a binomial heap is_empty iff it is [].

Correctness is trivial. The remaining operations are more interesting.

16.2.1 Insertion

A crucial property of binomial trees is that we can link two binomial trees of rank $r$ to form a binomial tree of rank $r+1$, simply by prepending one tree as the first child of the other. To preserve the heap property, we add the tree with the bigger root element below the tree with the smaller root element. This linking of trees is illustrated in Figure 16.2. Formally:

$$\text{link} :: \text{a tree} \Rightarrow \text{a tree} \Rightarrow \text{a tree}$$
Chapter 16  Binomial Heaps

![Figure 16.2](image)

Figure 16.2  Linking two binomial trees of rank 2 to form a binomial tree of rank 3, by linking the left tree as first child of the right tree, as indicated by the dashed line. We assume that the root element of the left tree is greater than or equal to the root element of the right tree, such that the heap property is preserved.

\[
\text{link } (\langle r, x_1, t_{s_1} \rangle =: t_1, \langle r', x_2, t_{s_2} \rangle =: t_2) =
\begin{cases}
\text{if } x_1 \leq x_2 \text{ then } \langle r + 1, x_1, t_2 \# t_{s_1} \rangle & \text{else } \langle r + 1, x_2, t_1 \# t_{s_2} \rangle
\end{cases}
\]

By case distinction, we can easily prove that \text{link} preserves the invariant and that the resulting tree contains the elements of both arguments.

\[
\text{bheap } t_1 \land \text{bheap } t_2 \land \text{rank } t_1 = \text{rank } t_2 \implies \text{bheap } (\text{link } t_1, t_2)
\]

\[
\text{mset_tree } (\text{link } t_1, t_2) = \text{mset_tree } t_1 + \text{mset_tree } t_2
\]

The link operation forms the basis of inserting a tree into a heap: if the heap does not contain a tree with the same rank, we can simply insert the tree at the correct position in the heap. Otherwise, we merge the two trees and recursively insert the result. For our purposes, we can additionally assume that the rank of the tree to be inserted is smaller than or equal to the lowest rank in the heap, which saves us a case in the following definition:

\[
\text{ins_tree :: 'a tree } \Rightarrow \text{'a trees } \Rightarrow \text{'a trees}
\]

\[
\text{ins_tree } t \ [] = [t]
\]

\[
\text{ins_tree } t_1 \ (t_2 \# \ ts)
\]

\[
= \text{if rank } t_1 < \text{rank } t_2 \text{ then } t_1 \# t_2 \# ts \text{ else } \text{ins_tree } (\text{link } t_1, t_2) \ ts
\]

Invariant preservation and functional correctness of \text{ins_tree} is easily proved by induction using the respective properties for \text{link}:
### 16.2 Implementation of ADT *Priority Queue*

#### 16.2.1 Insertion

A single element is inserted as a one-element (rank 0) tree:

\[
\begin{align*}
\text{insert} &:: 'a \Rightarrow 'a \text{ trees} \\
\text{insert } x \text{ ts} & = \text{ins_tree} \langle 0, x, \[] \rangle \text{ ts}
\end{align*}
\]

The above definition meets the specification for insert required by the *Priority Queue* ADT:

\[
\begin{align*}
invar t & \rightarrow invar \ (\text{insert } x \ t) \\
\text{mset_trees} \ (\text{insert } x \ t) & = \{ x \} + \text{mset_trees} \ t
\end{align*}
\]

#### 16.2.2 Merging

Recall the merge algorithm used in top-down merge sort (Section 2.4). It merges two sorted lists by repeatedly taking the smaller list head. We use a similar idea for merging two heaps: if the rank of one list’s head is strictly smaller, we choose it. If both ranks are equal, we link the two heads and insert the resulting tree into the merged remaining heaps. Thus, the resulting heap will be strictly ordered by rank. Formally:

\[
\begin{align*}
\text{merge} &:: 'a \text{ trees} \Rightarrow 'a \text{ trees} \Rightarrow 'a \text{ trees} \\
\text{merge } ts_1 \ [\ ] & = ts_1 \\
\text{merge } [\ ] \ ts_2 & = ts_2 \\
\text{merge } (t_1 \ # \ ts_1 \ =: \ h_1) \ (t_2 \ # \ ts_2 \ =: \ h_2) & = (\text{if rank } t_1 < \text{rank } t_2 \ \text{then } t_1 \ # \ \text{merge } ts_1 \ h_2 \\
& \ \text{else if rank } t_2 < \text{rank } t_1 \ \text{then } t_2 \ # \ \text{merge } h_1 \ ts_2 \\
& \ \text{else } \text{ins_tree} \ \langle \text{link } t_1 \ t_2 \rangle \ (\text{merge } ts_1 \ ts_2)
\end{align*}
\]

The *merge* function can be regarded as an algorithm for adding two sparse binary numbers. This intuition is explored in Exercise 16.2.

We show that the merge operation preserves the invariant and adds the elements:

\[
\begin{align*}
invar ts_1 \wedge invar ts_2 & \rightarrow invar \ (\text{merge } ts_1 \ ts_2) \\
\text{mset_trees} \ (\text{merge } ts_1 \ ts_2) & = \text{mset_trees} \ ts_1 + \text{mset_trees} \ ts_2
\end{align*}
\]

The proof is straightforward, except for preservation of the binomial heap invariant. We first show that merging two heaps does not decrease the lowest rank in these heaps.
This ensures that prepending the head with smaller rank to the merged remaining heaps results in a sorted heap. Moreover, when we link two heaps of equal rank, this ensures that the linked tree’s rank is smaller than or equal to the ranks in the merged remaining trees, as required by the ins_tree function. We phrase this property as preservation of lower rank bounds, i.e., a lower rank bound of both heaps is still a lower bound for the merged heap:

\[
t' \in \text{set} (\text{merge } ts_1 ts_2) \land (\forall t_1 \in \text{set } ts_1. \text{ rank } t < \text{ rank } t_1) \land (\forall t_2 \in \text{set } ts_2. \text{ rank } t < \text{ rank } t_2) \implies \text{rank } t < \text{ rank } t'
\]

The proof is by straightforward induction, relying on an analogous bounding lemma for ins_tree.

### 16.2.3 Finding a Minimal Element

For a binomial tree, the root node always contains a minimal element. Unfortunately, there is no such property for the whole heap—the minimal element may be at the root of any of the heap’s trees. To get a minimal element from a non-empty heap, we look at all root nodes:

```haskell
get_min :: 'a trees => 'a
get_min [t] = root t
get_min (t # ts) = min (root t) (get_min ts)
```

Correctness of this operation is proved by a simple induction:

\[
\text{mset_trees } ts \neq \{\} \land \text{invar } ts \implies 
\text{get_min } ts = \text{Min_mset (mset_trees } ts)
\]

### 16.2.4 Deleting a Minimal Element

To delete a minimal element, we first need to find one and then remove it. Removing the root node of a tree with rank \( r \) leaves us with a list of its children, which are binomial trees of ranks \( r-1, \ldots, 0 \). Reversing this list yields a valid binomial heap, which we merge with the remaining trees in the original heap:

```haskell
del_min :: 'a trees => 'a trees
del_min ts = (case get_min_rest ts of (ts1, ts2) => merge (rev ts1) ts2)
```
Here, the auxiliary function \texttt{get\_min\_rest} splits a heap into a tree with minimal root element, and the remaining trees.

\begin{verbatim}
get_min_rest :: 'a trees \Rightarrow 'a tree \times 'a trees
get_min_rest [t] = (t, [])
get_min_rest (t # ts) = (let (t', ts') = get_min_rest ts
  in if root t \leq root t' then (t, ts) else (t', t # ts'))
\end{verbatim}

We prove that, for a non-empty heap, \texttt{del\_min} preserves the invariant and deletes the minimal element:

\[
\begin{align*}
  ts \neq [] \land \text{invar ts} & \rightarrow \text{invar (del\_min ts)} \\
  ts \neq [] & \rightarrow \text{mset\_trees ts} = \text{mset\_trees (del\_min ts)} + \{ \text{get\_min ts} \}
\end{align*}
\]

The proof is straightforward. For invariant preservation, the key is to show that \texttt{get\_min\_rest} preserves the invariants:

\[
\begin{align*}
  \text{get\_min\_rest ts} = (t', ts') \land ts \neq [] \land \text{invar ts} & \rightarrow \text{bheap t'} \\
  \text{get\_min\_rest ts} = (t', ts') \land ts \neq [] \land \text{invar ts} & \rightarrow \text{invar ts'}
\end{align*}
\]

To show that we actually remove a minimal element, we show that \texttt{get\_min\_rest} selects the same tree as \texttt{get\_min}:

\[
\begin{align*}
  ts \neq [] \land \text{get\_min\_rest ts} = (t', ts') & \rightarrow \text{root t'} = \text{get\_min ts}
\end{align*}
\]

### 16.3 Running Time Analysis

The running time functions are shown in Appendix B.7. Intuitively, the operations are linear in the length of the heap, which in turn is logarithmic in the number of elements (see Section 16.1).

The running time analysis for \texttt{insert} is straightforward. The running time is dominated by \texttt{ins\_tree}. In the worst case, it iterates over the whole heap, taking constant time per iteration. By straightforward induction, we show

\[
T_{\text{ins\_tree}} t ts \leq |ts| + 1
\]

and thus

\[
\text{invar ts} \rightarrow T_{\text{insert}} x ts \leq \log (|\text{mset\_trees ts}| + 1) + 1
\]

The running time analysis for merge is more interesting. In each recursion, we need constant time to compare the ranks. However, if the ranks are equal, we link the trees and insert them into the merger of the remaining heaps. In the worst case, this costs
linear time in the length of the merger. A naive analysis would estimate $|\text{merge } ts_1 ts_2| \leq |ts_1| + |ts_2|$, and thus yield a quadratic running time in the length of the heap.

However, we can do better: we observe that every link operation in $\text{ins\_tree}$ reduces the number of trees in the heap. Thus, over the whole merge, we can only have linearly many link operations in the combined size of both heaps.

To formalize this idea, we estimate the running time of $\text{ins\_tree}$ and $\text{merge}$ together with the length of the result:

$$T_{\text{ins\_tree}} t ts + |\text{ins\_tree } t ts| = 2 + |ts|$$
$$T_{\text{merge}} ts_1 ts_2 + |\text{merge } ts_1 \text{ ts}_2| \leq 2 \cdot (|ts_1| + |ts_2|) + 1$$

Both estimates can be proved by straightforward induction, and from the second estimate we easily derive a bound for $\text{merge}$:

$$\text{invar } ts_1 \land \text{invar } ts_2 \rightarrow T_{\text{merge}} ts_1 \text{ ts}_2 \leq 4 \cdot \lg (|\text{mset\_trees } ts_1| + |\text{mset\_trees } ts_2| + 1) + 1$$

From the bound for $\text{merge}$ and (16.1) we can easily derive a bound for $\text{del\_min}$:

$$\text{invar ts } \land ts \neq [] \rightarrow T_{\text{del\_min}} ts \leq 6 \cdot \lg (|\text{mset\_trees } ts| + 1) + 2$$

The only notable point is that we use a linear time bound for reversing a list, as explained in Section 1.5.1:

$$T_{\text{rev}} :: 'a list \Rightarrow \text{nat}$$
$$T_{\text{rev}} xs = |xs| + 1$$

### 16.4 Exercises

**Exercise 16.1.** A node in a tree is on level $n$ if it is $n$ edges away from the root. Define a function $\text{nod} :: \text{nat } \Rightarrow \text{'a tree } \Rightarrow \text{nat}$ such that $\text{nod } n t$ is the number of nodes on level $n$ in tree $t$ and show that a binomial tree of rank $r$ has $\binom{r}{l}$ nodes on level $l$. In Isabelle, $\binom{r}{l}$ is written $\text{rank } t \text{ choose } l$ and thus you should prove

$$\text{btree } t \rightarrow \text{nod } l \text{ t } = \text{rank } t \text{ choose } l$$

Hint: You might want to prove separately that

$$\sum_{i=0}^{i<r} \binom{i}{n} = \binom{r}{n+1}$$

**Exercise 16.2.** Sparse binary numbers represent a binary number by a list of the positions of set bits, sorted in ascending order. Thus, the list $\{1, 3, 4\}$ represents the number $11010$. In general, $\{p_1, \ldots, p_n\}$ represents $2^{p_1} + \cdots + 2^{p_n}$.

Implement sparse binary numbers in Isabelle, using the type $\text{nat list}$. 
1. Define a function \( \text{invar	extunderscore sn} :: \text{nat list} \Rightarrow \text{bool} \) that checks for strictly ascending bit positions, a function \( \text{num	extunderscore of} :: \text{nat list} \Rightarrow \text{nat} \) that converts a sparse binary number to a natural number, and a function \( \text{add} :: \text{nat list} \Rightarrow \text{nat list} \Rightarrow \text{nat list} \) to add sparse binary numbers.

2. Show that \( \text{add} \) preserves the invariant and actually performs addition as far as \( \text{num	extunderscore of} \) is concerned.

3. Define a running time function for \( \text{add} \) and show that it is linear in the list lengths.

   Hint: The bit positions in sparse binary numbers are analogous to binomial trees of a certain rank in a binomial heap. The \( \text{add} \) function should be implemented similarly to the \( \text{merge} \) function, using a \( \text{carry} \) function to insert a bit position into a number (similar to \( \text{ins	extunderscore tree} \)). Correctness and running time can be proved similarly.

---

16.5 Chapter Notes

Binomial queues were invented by Vuillemin [1978]. Functional implementations were given by King [1994] and Okasaki [1998]. A functional implementation was verified by Meis et al. [2010], a Java implementation by Müller [2018].
Part IV

Advanced Design and Analysis Techniques
You probably have seen this function before:

\[
\begin{align*}
\text{fib} :: \text{nat} & \Rightarrow \text{nat} \\
\text{fib} \ 0 & = 0 \\
\text{fib} \ 1 & = 1 \\
\text{fib} \ (n + 2) & = \text{fib} \ (n + 1) + \text{fib} \ n
\end{align*}
\]

It computes the well-known Fibonacci numbers. You may also have noticed that calculating \( \text{fib} \ 50 \) already causes quite some stress for your computer and there is no hope for \( \text{fib} \ 500 \) to ever return a result.

This is quite unfortunate considering that there is a very simple imperative program to compute these numbers efficiently:

```c
int fib(n) {
    int a = 0;
    int b = 1;
    for (i in 1..n) {
        int temp = b;
        b = a + b;
        a = temp;
    }
    return a;
}
```

So we seem to be caught in an adverse situation here: either we use a clear and elegant definition of \( \text{fib} \) or we get an efficient but convoluted implementation for \( \text{fib} \). Admittedly, we could just prove that both formulations are the same function, and use whichever one is more suited for the task at hand. For \( \text{fib} \), of course, it is trivial to define a functional analogue of the imperative program and to prove its
correctness. However, doing this for all recursive functions we would like to define is tedious. Instead, this chapter will sketch a recipe that allows to define such recursive functions in the natural way, while still getting an efficient implementation “for free”.

In the following, the Fibonacci function will serve as a simple example on which we can illustrate the idea. Next, we will show how to prove the correctness of the efficient implementation in an efficient way. Subsequently, we will discuss further details of the approach and how it can be applied beyond _fib_. The chapter closes with the study of two famous (and archetypical) dynamic programming algorithms: the Bellman-Ford algorithm for finding shortest paths in weighted graphs and an algorithm due to Knuth for computing optimal binary search trees.

### 17.1 Memoization

Let us consider the tree of recursive calls that are issued when computing _fib_ 5 in Fig. 17.1. We can see that the subtree for _fib_ 3 is computed twice, and that the subtree for _fib_ 2 is even computed three times. How can we avoid these repeated computations? A common solution is _memoization_: we store previous computation results in some kind of memory and consult it to potentially recall a memoized result before issuing another recursive computation.

Below you see a simple memoizing version of _fib_ that implements the memory as a map of type _nat_ → _nat_ (see Section 6.4 for the notation):

```plaintext
fib₁ :: nat ⇒ (nat → nat) ⇒ nat × (nat → nat)
fib₁ 0 m = (0, m(0 → 0))
fib₁ 1 m = (1, m(1 → 1))
```
\[ \text{fib}_1 (n + 2) \, m \]

\[ = \begin{cases} \text{let } (i, m) = \text{case } m \, n \, \text{of } \text{None} \Rightarrow \text{fib}_1 \, n \, m \mid \text{Some } i \Rightarrow (i, m); & \text{(j, m)} = \\
\text{case } m \, (n + 1) \, \text{of } \text{None} \Rightarrow \text{fib}_1 \, (n + 1) \, m \mid \text{Some } j \Rightarrow (j, m) & \text{in } (i + j, m(n + 2 \leftrightarrow i + j)) \end{cases} \]

And indeed, we can ask Isabelle to compute (via the \textit{value} command) \text{fib}_1 50 or even \text{fib}_1 500 and we get the result within a split second.

However, we are not yet happy with this code. Carrying the memory around means a lot of additional weight for the definition of \text{fib}_1, and proving that this function computes the same value as \text{fib} is not completely trivial (how would you approach this?). Let us streamline the definition first by pulling out the reading and writing of memory into a function \textit{memo} (for a type \textit{'k} of keys and a type \textit{'v} of values):

\[ \text{memo} :: \]

\[ 'k \Rightarrow (('k \rightarrow 'v) \Rightarrow 'v \times ('k \rightarrow 'v)) \Rightarrow ('k \rightarrow 'v) \Rightarrow 'v \times ('k \rightarrow 'v) \]

\[ \text{memo } k \, f \, m = \begin{cases} \text{case } m \, k \, \text{of } \text{None} \Rightarrow \text{let } (v, m) = f \, m \, \text{in } (v, m(k \mapsto v)) ; & \mid \text{Some } v \Rightarrow (v, m) \end{cases} \]

\[ \text{fib}_2 :: \text{nat} \Rightarrow (\text{nat} \rightarrow \text{nat}) \Rightarrow \text{nat} \times (\text{nat} \rightarrow \text{nat}) \]

\[ \text{fib}_2 \, 0 = \text{memo} \, 0 \, (\lambda m. \, (0, m)) \]

\[ \text{fib}_2 \, 1 = \text{memo} \, 1 \, (\lambda m. \, (1, m)) \]

\[ \text{fib}_2 \, (n + 2) = \text{memo} \, (n + 2) \]

\[ (\lambda m. \, \text{let } (i, m) = \text{fib}_2 \, n \, m; \]

\[ (j, m) = \text{fib}_2 \, (n + 1) \, m \]

\[ \text{in } (i + j, m)) \]

This already looks a lot more like the original definition but it still has one problem: we have to thread the memory through the program explicitly. This can be become rather tedious for more complicated programs and diverges from the original shape of the program, complicating the proofs.
17.1.1 Enter the Monad

Let us examine the type of \texttt{fib} more closely. We can read it as the type of a function that, given a natural number, returns a \textit{computation}. Given an initial memory, it computes a pair of a result and an updated memory. We can capture this notion of “stateful” computations in a data type:

\begin{verbatim}
datatype ('s, 'a) state = State ('s => 'a x 's)
\end{verbatim}

A value of type ('s, 'a) state represents a stateful computation that returns a result of type 'a and operates on states of type 's. The constant \texttt{run_state} forces the evaluation of a computation starting from some initial state:

\begin{verbatim}
run_state :: ('s, 'a) state => 's => 'a x 's
run_state (State f) s = f s
\end{verbatim}

The advantage of this definition may not seem immediate. Its value only starts to show when we see how it allows us to \textit{chain} stateful computations. To do so, we only need to define two constants: \texttt{return} to pack up a result in a computation, and \texttt{bind} to chain two computations after each other.

\begin{verbatim}
return :: 'a => ('s, 'a) state
return x = State (\s. (x, s))

bind :: ('s, 'a) state => ('a => ('s, 'b) state) => ('s, 'b) state
bind a f = State (\s. let (x, s) = run_state a s in run_state (f x) s)
\end{verbatim}

We add a little syntax on top and write \([x]\) for \texttt{return} \(x\), and \(a >>= f\) instead of \texttt{bind a f}. The “identity” computation \([x]\) simply leaves the given state unchanged and produces \(x\) as a result. The chained computation \(a >>= f\) starts with some state \(s\), runs \(a\) on it to produce a pair of a result \(x\) and a new state \(s'\), and then evaluates \(f x\) to produce another computation that is run on \(s'\).

We have now seen how to pass state around but we are not yet able to interact with it. For this purpose we define \texttt{get} and \texttt{set} to retrieve and update the current state, respectively:
get :: ('s, 's) state
get = State (λs. (s, s))

set :: 's ⇒ ('s, unit) state
set s' = State (λ_ (()), s'))

Let us reformulate fib with the help of these concepts:

memo1 ::
  'k ⇒ ('k → 'v, 'v) state ⇒ ('k → 'v, 'v) state
memo1 k a
= get >>=
  (λm. case m k of
      None ⇒ a >>= (λv. set (m(k ↦ v)) >>= (λ_. {v}))
     | Some x ⇒ {x})

fib3 :: nat ⇒ (nat → nat, nat) state
fib3 0 = {0}
fib3 1 = {1}
fib3 (n + 2)
= memo1 (n + 2) (fib3 n >>= (λi. fib3 (n + 1) >>= (λj. {i + j}))))

Can you see how we have managed to hide the whole handling of state behind the scenes? The only explicit interaction with the state is now happening inside of memo1. This is sensible as this is the only place where we really want to recall a memoized result or to write a new value to memory.

While this is great, we still want to polish the definition further: the syntactic structure of the last case of fib3 still does not match fib exactly. To this end, we lift function application f x to the state monad:

(_) :: ('s, 'a ⇒ ('s, 'b) state) state ⇒ ('s, 'a) state ⇒ ('s, 'b) state
f_m . x_m = (f_m >>= (λf. x_m >>= (λx. f x)))

We can now spell out our final memoizing version of fib where (_) replaces ordinary function applications in the original definition:
196 Chapter 17 Dynamic Programming

\[
\begin{align*}
\text{fib}_4 & :: \text{nat} \Rightarrow (\text{nat} \rightarrow \text{nat}, \text{nat}) \text{ state} \\
\text{fib}_4 \ 0 &= \langle 0 \rangle \\
\text{fib}_4 \ 1 &= \langle 1 \rangle \\
\text{fib}_4 \ (n + 2) &= \text{memo}_1 \ (n + 2) \ (\langle \lambda i. \langle \lambda j. \{i + j\} \rangle \rangle \ . \ (\text{fib}_4 \ n) \ . \ (\text{fib}_4 \ (n + 1)))
\end{align*}
\]

You may wonder why we added that many additional computations in this last step. On the one hand, we have gained the advantage that we can now closely follow the syntactic structure of \text{fib} to prove that \text{fib}_4 is correct (notwithstanding that \text{memo}_1 will need a special treatment, of course). On the other hand, we can remove most of these additional computations in a final post-processing step.

17.1.2 Memoization and Dynamic Programming

Let us recap what we have seen so far in this chapter. We noticed that the naive recursive formulation of the Fibonacci numbers leads to a highly inefficient implementation. We then showed how to work around this problem by using memoization to obtain a structurally similar but efficient implementation. After all this, you may wonder why this chapter is titled \textit{Dynamic Programming} and not \textit{Memoization}.

Dynamic programming relies on two main principles. First, to find an optimal solution for a problem by computing it from optimal solutions for “smaller” instances of the same problem, i.e. recursion. Second, to \textit{memoize} these solutions for smaller problems in, e.g. a table. Thus we could be bold and state:

\[
\text{dynamic programming} = \text{recursion} + \text{memoization}
\]

A common objection to this equation would be that memoization should be distinguished from \textit{tabulation}. In this view, the former only computes “necessary” solutions for smaller sub-problems, while the latter just “blindly” builds solutions for sub-problems of increasing size, many of which might be unnecessary. The benefit of tabulation could be increased performance, for instance due to improved caching. We believe that this distinction is largely irrelevant to our approach. First, in this book we focus on asymptotically efficient solutions, not constant-factor optimizations. Second, in many dynamic programming algorithms memoization would actually compute solutions for the same set of sub-problems as tabulation does. No matter which of the two approaches is used in the implementation, the hard part is usually to come up with a recursive solution that can efficiently make use of sub-problems in the first place.

There are problems, however, where clever tabulation instead of naive memoization is necessary to achieve an \textit{asymptotically optimal solution} in terms of \textit{memory}
consumption. One instance of this is the Bellman-Ford algorithm presented in Section 17.4. On this example, we will show that our approach is also akin to tabulation. It can easily be introduced as a final “post-processing” step.

Some readers may have noticed that our optimized implementations of \( \texttt{fib} \) are not really optimal as they use a map for memoization. Indeed it is possible to swap in other memory implementations as long as they provide a \textit{lookup} and an \textit{update} method. One can even make use of imperative data structures like arrays. As this is not the focus of this book, the interested reader is referred to the literature that is provided at the end of this chapter. Here, we will just assume that the maps used for memoization are implemented as red-black trees (and Isabelle’s code generator can be instructed to do so).

For the remainder of this chapter, we will first outline how to prove that \( \texttt{fib}_4 \) is correct. Then, we will sketch how to apply our approach of memoization beyond \( \texttt{fib} \). Afterwards, we will study some prototypical examples of dynamic programming problems and show how to apply the above formula to them.

17.2 Correctness of Memoization

We now want to prove that \( \texttt{fib}_4 \) is correct. But what is it exactly that we want to prove? We surely want \( \texttt{fib}_4 \) to produce the same result as \( \texttt{fib} \) when run with an empty memory (in this chapter we write the empty map \( \lambda \_ \_. \texttt{None} \) simply as \( \texttt{empty} \)):

\[
\text{fst} \ (\text{run\_state} \ (\texttt{fib}_4 \ n) \ \text{empty}) = \text{fib} \ n
\]  

(17.1)

If we were to make a naive attempt at this prove, we would probably start with an induction on the computation of \( \texttt{fib} \) just to realize that the induction hypotheses are not strong enough to prove the recursion case, since they demand an empty memory. We can attempt generalization as a remedy:

\[
\text{fst} \ (\text{run\_state} \ (\texttt{fib}_4 \ n) \ m) = \text{fib} \ n
\]

However, this statement does not hold anymore for every memory \( m \).

What do we need to demand from \( m \)? It should only memoize values that are consistent with \( \texttt{fib} \):

\[
\text{type} \_\text{synonym} \ 'a \text{ mem} = \ (\text{nat} \rightarrow \text{nat}, 'a) \text{ state}
\]

\[
\text{cmem} :: (\text{nat} \rightarrow \text{nat}) \Rightarrow \text{bool}
\]

\[
\text{cmem} \ m = (\forall n \in \text{dom} \ m. \ m \ n = \text{Some} \ (\text{fib} \ n))
\]

\[
\text{dom} :: ('k \rightarrow 'v) \Rightarrow 'k \text{ set}
\]
Note that, from now, we use the type ‘a mem to denote memoized values of type ‘a that have been “wrapped up” in our memoizing state monad. Using cmem, we can formulate a general notion of equivalence between a value \( v \) and its memoized version \( a \), written \( v \gg a \): starting from a consistent memory \( m \), \( a \) should produce another consistent memory \( m' \), and the result \( v \).

Thus we want to prove

\[
\text{fib } n \gg \text{fib}_4 \ n
\]

via computation induction on \( n \). For the base cases we need to prove statements of the form \( v \gg (u) \), which follow trivially after unfolding the involved definitions. For the induction case, we can unfold \( \text{fib}_4 \ (n + 2) \), and get rid of \( \text{memo}_1 \) by applying the following rule (which we instantiate with \( a = \text{fib}_4 \ n \)):

\[
\text{fib } n \gg a \longrightarrow \text{fib } n \gg \text{memo}_1 \ n \ a
\]  

(17.3)

For the remainder of the proof, we now want to unfold \( \text{fib} \ (n + 2) \) and then follow the syntactic structure of \( \text{fib}_4 \) and \( \text{fib} \) in lockstep. To do so, we need to find a proof rule for function application. That is, what do we need in order to prove \( f \ a \gg f \ _m \) . \( x_m \)? For starters, \( x \gg x_m \) seems reasonable to demand. But what about \( f \) and \( f_m \)? If \( f \) has type \( ‘a \Rightarrow ‘b \), then \( f_m \) is of type \( ‘a \Rightarrow ‘b \text{ mem} \) \( \text{mem} \). Intuitively, we want to state something along these lines:

“\( f_m \) is a memoized function that, when applied to a value \( x \), yields a memoized value that is equivalent to \( f \ x \).”

This goes beyond what we can currently express with \((\gg)\) as \( v \gg a \) merely states that “\( a \) is a memoized value equivalent to \( v \).” What we need is more liberty in our choice of equivalence. That is, we want to use statements \( v \gg R \ a \), with the meaning: “\( a \) is a memoized value that is related to \( v \) by \( R \).” The formal definition is analogous to \((\gg)\) (and \((\gg) = (\gg (=))\):
17.2 Correctness of Memoization

\[(\forall \cdot) :: 'a \Rightarrow ('a \Rightarrow 'b \Rightarrow \text{bool}) \Rightarrow 'b \text{ mem} \Rightarrow \text{bool} \]

\[v \triangleright R \Rightarrow s \rightarrow \ (\forall m. \ \text{cmem} m \rightarrow \ \text{let} (v', m') = \text{run_state} s m \ \text{in} \ v \triangleright v' \land \text{cmem} m')\]

However, we still do not have a means of expressing the second part of our sentence. To this end, we use the function relator \((\Rightarrow)\):

\[(\Rightarrow) :: ('a \Rightarrow 'c \Rightarrow \text{bool}) \Rightarrow ('b \Rightarrow 'd \Rightarrow \text{bool}) \Rightarrow ('a \Rightarrow 'b) \Rightarrow ('c \Rightarrow 'd) \Rightarrow \text{bool} \]

\[R \Rightarrow S = (\lambda f g. \ \forall x y. \ R x y \rightarrow S (f x) (g y))\]

Spelled out, we have \((R \Rightarrow S) f g\) if for any values \(x\) and \(y\) that are related by \(R\), the values \(f x\) and \(g y\) are related by \(S\).

We can finally state a proof rule for application:

\[x \triangleright x_m \land f \triangleright \Rightarrow (\Rightarrow) f_m \rightarrow f x \triangleright f_m \cdot x_m\]  \hspace{1cm} (17.4)

In our concrete example, we apply it once to the goal

\[\text{fib} (n + 1) + \text{fib} n \triangleright \{ \lambda a. \ \{ \lambda b. \ (a + b) \} \} \cdot (\text{fib}_4 (n + 1)) \cdot (\text{fib}_4 n)\]

solve the first premise with the induction hypotheses, and arrive at

\[(+ \ (\text{fib} (n + 1)) \triangleright \Rightarrow (\Rightarrow) \{ \lambda a. \ \{ \lambda b. \ (a + b) \} \} \cdot (\text{fib}_4 (n + 1))\]

Our current rule for application (17.4) does not match this goal. Thus we need to generalize it. In addition, we need a new rule for \text{return}, and a rule for \((\Rightarrow)\). To summarize, we need the following set of theorems about our consistency relation, applying them wherever they match syntactically to finish the proof of (17.2):

\[R x y \rightarrow x \triangleright R \Rightarrow y\]

\[x \triangleright R x_m \land f \triangleright S f_m \rightarrow f x \triangleright S f_m \cdot x_m\]

\[(\forall x y. \ R x y \rightarrow S (f x) (g y)) \rightarrow (R \Rightarrow S) f g\]

The theorem we aimed for initially

\[\text{fst (run_state (\text{fib}_4 n) \text{ empty})} = \text{fib} n\]  \hspace{1cm} (17.1)

is now a trivial corollary of \text{fib} n \triangleright \text{fib}_4 n. Note that by reading the equation from right to left, we have an easy way to make the memoization transparent to an end-user of \text{fib}. 

Theorem we aimed for initially

\[\text{fst (run_state (\text{fib}_4 n) \text{ empty})} = \text{fib} n\]  \hspace{1cm} (17.1)
17.3 Details of Memoization*

In this section, we will look at some further details of the memoization process and sketch how it can be applied beyond \( \text{fib} \). First note that our approach of memoization hinges on two rather independent components: We transform the original program to use the state monad, to thread (an \textit{a priori} arbitrary) state through the program. Only at the call sites of recursion, we then introduce the memoization functionality by issuing lookups and updates to the memory (as implemented by \texttt{memo1}). We will name this first process \textit{monadification}. For the second component, many different memory implementations can be used, as long as we can define \texttt{memo1} and prove its characteristic theorem (17.3). For details on this, the reader is referred to the literature. Here, we want to turn our attention towards monadification.

To discuss some of the intricacies of monadification, let us first stick with \( \text{fib} \) for a bit longer and consider the following alternative definition (which is mathematically equivalent but not the same program):

\[
\text{fib} \ n = (\text{if } n = 0 \text{ then } 0 \text{ else } 1 + \text{sum_list (map fib [0..<n - 1])})
\]

We have not yet seen how to handle two ingredients of this program: constructs like \textit{if-then-else} or case-combinators; and higher-order functions such as \texttt{map}.

It is quite clear how \textit{if-then-else} can be lifted to the state monad:

\[
\text{if}_m : \text{bool mem} \Rightarrow \text{a mem} \Rightarrow \text{a mem} \Rightarrow \text{'a mem}
\]

\[
\text{if}_m \ b_m \ x_m \ y_m = b_m \gg (\lambda b. \text{if } b \text{ then } x_m \text{ else } y_m)
\]

By following the structure of the terms, we can also deduce a proof rule for \( \text{if}_m \):

\[
\begin{align*}
& b \vdash b_m \wedge x \vdash_R x_m \wedge y \vdash_R y_m \\
& (\text{if } b \text{ then } x \text{ else } y) \vdash_R \text{if}_m \ b_m \ x_m \ y_m
\end{align*}
\]

However, suppose we want to apply this proof rule to our new equation for \( \text{fib} \). We will certainly need the knowledge of whether \( n = 0 \) to make progress in the correctness proof. Thus we make our rule more precise:

\[
\begin{align*}
& b \vdash b_m \wedge (b \rightarrow x \vdash_R x_m) \wedge (\neg b \rightarrow y \vdash_R y_m) \\
& (\text{if } b \text{ then } x \text{ else } y) \vdash_R \text{if}_m \ b_m \ x_m \ y_m
\end{align*}
\]

How can we lift \texttt{map} to the state monad level? Consider its defining equations:

\footnote{If you are just interested in the dynamic programming algorithms of the following sections, this section can safely be skipped on first reading.}
\[ \begin{align*}
map f [] &= [] \\
map f (x \# xs) &= f x \# \map f xs
\end{align*} \]

We can follow the pattern we used to monadify \( \text{fib} \) to monadify \( \text{map} \):

\[ \begin{align*}
\map_{m'} f [] &= \langle [] \rangle \\
\map_{m'} f (x \# xs) &= \langle \lambda a. \langle \lambda b. \langle a \# b \rangle \rangle \rangle \cdot (\langle f \rangle \cdot \langle x \rangle) \cdot (\map_{m'} f xs)
\end{align*} \]

We have obtained a function \( \map_{m'} \) of type

\[ ('a \Rightarrow 'b \text{ mem}) \Rightarrow 'a \text{ list} \Rightarrow 'b \text{ list mem} \]

This is not yet compatible with our scheme of lifting function application to \( (\cdot) \). We need a function of type

\[ (('a \Rightarrow 'b \text{ mem}) \Rightarrow ('a \text{ list} \Rightarrow 'b \text{ list mem}) \text{ mem}) \text{ mem} \]

because \( \text{map} \) has two arguments and we need one layer of the state monad for each of its arguments. Therefore we simply define

\[ \map_m = \langle \lambda f. \map_{m'} f \rangle \]

For inductive proofs about the new definition of \( \text{fib} \), we also need the knowledge that \( \text{fib} \) is recursively applied only to smaller values than \( n \) when computing \( \text{fib} n \). That is, we need to know which values \( f \) is applied to in \( \map f xs \). We can encode this knowledge in a proof rule for \( \map \):

\[ \begin{align*}
zs &= ys \land (\forall x. x \in \text{set} ys \rightarrow f x \lor f_m x) \rightarrow \\
\map f zs \triangleright \text{list_all2} R \map_m \cdot \langle f_m \rangle \cdot \langle ys \rangle
\end{align*} \]

The relator \( \text{list_all2} \) lifts \( R \) to a pairwise relation on lists:

\[ \text{list_all2} R xs ys = (|zs| = |ys| \land (\forall i < |zs| . \ R (xs ! i) (ys ! i))) \]

To summarize, here is a fully memoized version of the alternative definition of \( \text{fib} \):

\[ \begin{align*}
\text{fib}_m :: \text{nat} \Rightarrow \text{nat mem} \\
\text{fib}_m = \text{memo}_1 n \\
(\text{if}_m \langle n = 0 \rangle \langle 0 \rangle)
\end{align*} \]
Chapter 17  Dynamic Programming

\[
(\langle \lambda a. \{1 + a\} \rangle .
(\langle \lambda a. \{\text{sum\_list} a\} \rangle . (\text{map}_m . \{\text{fib}_m\} . \{[0..<n - 1]\})))
\]

The correctness proof for \(\text{fib}_m\) is analogous to the one for \(\text{fib}_4\), once we have proved the new rules discussed above.

At the end of this section, we note that the techniques that were sketched above also extend to case-combinators and other higher-order functions. Most of the machinery for monadification and the corresponding correctness proofs can be automated in Isabelle [Wimmer et al. 2018b]. Finally note that none of the techniques we used so far are specific to \(\text{fib}\). The only parts that have to be adopted are the definitions of \(\text{memo}\) and \(\text{cmem}\). In Isabelle, this can be done by simply instantiating a locale.

This concludes the discussion of the fundamentals of our approach towards verified dynamic programming. We now turn to the study of two typical examples of dynamic programming algorithms: the Bellman-Ford algorithm and an algorithm for computing optimal binary search trees.

17.4 The Bellman-Ford Algorithm

Calculating shortest paths in weighted graphs is a classic algorithmic task that we all encounter in everyday situations, such as planning the fastest route to drive from \(A\) to \(B\). In this scenario we can view streets as edges in a graph and nodes as street crossings. Every edge is associated with a weight, e.g. the time to traverse a street. We are interested in the path from \(A\) to \(B\) with minimum weight, corresponding to the fastest route in the example. Note that in this example it is safe to assume that all edge weights are non-negative.

Some applications demand negative edge weights as well. Suppose, we transport ourselves a few years into the future, where we have an electric car that can recharge itself via solar cells while driving. If we aim for the most energy-efficient route from \(A\) to \(B\), a very sunny route could then incur a negative edge weight.

The **Bellman-Ford algorithm** is a classic dynamic programming solution to the *single-destination shortest path problem* in graphs with negative edge weights. That is, we are given a directed graph with negative edge weights and some target vertex (known as *sink*), and we want to calculate the weight of the shortest (i.e. minimum weight) paths from every vertex to the sink. Figure 17.2 shows an example of such a graph.

Formally, we will take a simple view of graphs. We assume that we are given a number of nodes numbered \(0, \ldots, n\), and some sink \(t \in \{0..n\}\) (thus \(n = t = 4\) in the example).
17.4 The Bellman-Ford Algorithm

Figure 17.2 Example of a weighted directed graph

Edge weights are given by a function $W : \text{int} \Rightarrow \text{int} \Rightarrow \text{int extended}$. The type int extended extends the natural numbers with positive and negative infinity:

$$\text{datatype } 'a\,\text{extended} = \text{Fin}'a \mid \infty \mid -\infty$$

We refrain from giving the explicit definition of addition and comparison on this domain, and rely on your intuition instead. A weight assignment $W_{ij} = 1$ means that there is no edge from $i$ to $j$. The purpose of $-\infty$ will become clear later.

17.4.1 Deriving a Recursive Solution

The main idea of the algorithm is to consider paths in order of increasing length in the number of edges. In the example, we can immediately read off the weights of the shortest paths to the sink that use only one edge: only nodes 2 and 3 are directly connected to the sink, with edge weights 3 and 2, respectively; for all others the weight is infinite. How can we now calculate the minimum weight paths (to the sink) with at most two edges? For node 3, the weight of the shortest path with at most two edges is: either the weight of the path with one edge; or the weight of the edge from node 3 to node 2 plus the weight of the path with one edge from node 2 to the sink. Because $-2 + 3 = 1 \leq 2$, we get a new minimum weight of 1 for node 3. Following the same scheme, we can iteratively calculate the minimum path weights given in table 17.1.

The analysis we just ran on the example already gives us a clear intuition on all we need to deduce a dynamic program: a recursion on sub-problems, in this case to compute the weight of shortest paths with at most $i + 1$ edges from the weights of shortest paths with at most $i$ edges. To formalize this recursion, we first define the notion of a minimum weight path from some node $v$ to $t$ with at most $i$ edges, denoted as $OPT_{i} v$: 
### Table 17.1

The minimum weights of paths from vertices $v = 0 \ldots 4$ to $t$ that use at most $i = 0 \ldots 4$ edges.

<table>
<thead>
<tr>
<th>$i/v$</th>
<th>0</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>$\infty$</td>
<td>$\infty$</td>
<td>$\infty$</td>
<td>$\infty$</td>
<td>0</td>
</tr>
<tr>
<td>1</td>
<td>$\infty$</td>
<td>$\infty$</td>
<td>3</td>
<td>2</td>
<td>0</td>
</tr>
<tr>
<td>2</td>
<td>5</td>
<td>6</td>
<td>3</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>3</td>
<td>5</td>
<td>5</td>
<td>3</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>4</td>
<td>4</td>
<td>5</td>
<td>3</td>
<td>1</td>
<td>0</td>
</tr>
</tbody>
</table>

The definition of $OPT$ is given by:

$$OPT :: \mathbb{nat} \Rightarrow \mathbb{nat} \Rightarrow \mathbb{int\ extended}$$

$$OPT i v = \text{Min} (\{\text{weight} (v \# xs @ [t]) \mid 1 \leq i \land \text{set} \ xs \subseteq \{0..n\} \} \cup \{\text{if} \ t = v \text{ then } 0 \text{ else } \infty\})$$

The definition of `weight` is given by:

$$weight :: (\mathbb{nat} \Rightarrow \mathbb{nat} \Rightarrow \mathbb{int\ extended}) \Rightarrow \mathbb{nat\ list} \Rightarrow \mathbb{int\ extended}$$

$$weight _ [\_] = 0$$

$$weight W (v \# w \# xs) = W v w + weight W (w \# xs)$$

If $i = 0$, things are simple:

$$OPT 0 v = (\text{if} \ t = v \text{ then } 0 \text{ else } \infty)$$

A shortest path that constitutes $OPT (i + 1) v$ uses either at most $i$ or exactly $i + 1$ edges. That is, $OPT (i + 1) v$ is either $OPT i v$, or the weight of the edge from $v$ to any of its neighbours $w$ plus $OPT i w$:

$$OPT (i + 1) v = \text{min} (OPT i v) (\text{Min} \{W v w + OPT i w \mid w \leq n\})$$

Proof. We prove this equality by proving two inequalities:

(\text{lhs} \leq \text{rhs}) For this direction, we essentially need to show that every path on the rhs is covered by the lhs, which is trivial.

(\text{lhs} \geq \text{rhs}) We skip the cases where $OPT (i + 1) v$ is trivially 0 or $\infty$ (i.e. where it is given by the singleton set in the definition of $OPT$). Thus consider some $xs$ such that $OPT (i + 1) v = weight (v \# xs @ [t], |xs| \leq i, \text{ and set} \ xs \subseteq \{0..n\}$. The cases where $|xs| < i$ or $i = 0$ are trivial. Otherwise, we have $OPT (i + 1) v = W v (\text{hd} \ xs) + weight (xs @ [t])$ by definition of $weight$,
and $OPT \ i \ (hd \ xs) \leq \ weight \ (xs @ [i])$ by definition of $OPT$. Therefore, we can show:

$$OPT \ (i + 1) \ v \geq W \ v \ (hd \ xs) + OPT \ i \ (hd \ xs) \geq rhs$$

We can turn these equations into a recursive program:

```plaintext
bf :: nat ⇒ nat ⇒ int
bf 0 v = (if t = v then 0 else ∞)
bf (i + 1) v
      = min_list (bf i v # map (λw. W v w + bf i w) [0..<n + 1])
```

It is obvious that we can prove correctness of $bf$ by induction:

$$bf \ i \ v = OPT \ i \ v$$

### 17.4.2 Negative Cycles

Have we solved the initial problem now? The answer is “not quite” because we have ignored one additional complication. Consider our example table 17.1 again. The table stops at path length five because no shorter paths with more edges exist. For this example, five corresponds to the number of nodes, which bounds the length of the longest simple path. However, is it the case that we will never find shorter non-simple paths in other graphs? The answer is “no”. If a graph contains a negative reaching cycle, i.e. a cycle with a negative sum of edge weights from which the sink is reachable, then we can use it arbitrarily often to find shorter and shorter paths.

Luckily, we can use the Bellman-Ford algorithm to detect this situation by examining the relationship of $OPT \ n$ and $OPT \ (n + 1)$. The following proposition summarizes the key insight:

The graph contains a negative reaching cycle if and only if there exists a $v \leq n$ such that $OPT \ (n + 1) \ v < OPT \ n \ v$

**Proof.** If there is no negative reaching cycle, then all shortest paths are either simple or contain superfluous cycles of weight 0. Thus, we have $OPT \ (n + 1) \ v = OPT \ n \ v$ for all $v \leq n$.

Otherwise, there is a negative reaching cycle $ys = a \ # \ xs @ [a]$ with $weight \ ys < 0$. Working towards a contradiction, assume that $OPT \ n \ v \leq OPT \ (n + 1) \ v$ for all $v \leq n$. Using the recursion we proved above, this implies $OPT \ n \ v \leq W \ v \ u + OPT \ n \ u$ for all $u, v \leq n$. By applying this inequality to the nodes in $a \ # \ xs$, we can prove the inequality
This implies $0 \leq \text{weight } ys$, which yields the contradiction.

This means we can use $bf$ to detect the existence of negative reaching cycles by computing one more round, i.e. $bf (n + 1) v$ for all $v$. If nothing changes in this step, we know that there are no negative reaching cycles and that $bf n$ correctly represents the shortest path weights. Otherwise, there has to be a negative reaching cycle.

Finally, we can use memoization to obtain an efficient implementation that solves the single-destination shortest path problem. Applying our memoization technique from above, we first obtain a memoizing version $bf_m$ of $bf$. We then define the following program:

```haskell
bellman_ford :: ((nat × nat, int extended) mapping, int extended list option) state
bellman_ford
= iter_bf (n, n) >>=
  (λ_. map m' (bf_m n) [0..<n + 1] >>=
    (λxs. map_m' (bf_m (n + 1)) [0..<n + 1] >>=
      (λys. (if zs = ys then Some zs else None))))
```

Here, $iter_bf (n, n)$ just computes the values from $bf_m 0 0$ to $bf_m n n$ in a row-by-row manner. Using the reasoning principles that were described above (for $fib$), we can then prove that $bellman_ford$ indeed solves its intended task correctly ($shortest v$ is the length of the shortest path from $v$ to $t$):

$$
(\forall i \leq n. \forall j \leq n. -\infty < W i j) \implies
\text{fst (run_state bellman_ford empty)}
= (\text{if contains_negative_reaching_cycle then None}
  \text{ else Some (map shortest [0..<n + 1])})
$$

Here, $shortest$ is defined analogously to $OPT$ but for paths of unbounded length.

### 17.5 Optimal Binary Search Trees

In this book, we have studied various tree data structures that guarantee logarithmic running time bounds for operations such as lookups and updates into the tree. These bounds were usually worst-case and did not take into account any information about the actual series of queries that are to be issued to the data structure. In this section, instead, we want to focus on binary search trees that minimize the amount of work...
that needs to be done when the distribution of keys in a sequence of lookup operations is known in advance.

More formally, we want to study the following problem. We are given a list \([i..j]\) of integers ranging from \(i\) to \(j\) and a function \(p :: \text{int} \Rightarrow \text{nat}\) that maps each key in the range to a frequency with which this key is searched for. Our goal is to find a binary search tree that minimizes the expected number of comparisons when presented with a sequence of lookup operations for keys in the range \([i..j]\) that adhere to the distribution given by \(p\).

As an example, consider the range \([1..5]\) with probabilities \([10, 30, 15, 25, 20]\). This tree

![Tree](image)

incurs an expected value of 2.15 comparison operations. However, the minimal expected value is 2 and is achieved by this tree:

![Tree](image)

Our task is equivalent to minimizing the weighted path length (or cost) as we did for Huffman encodings (Chapter 24). Recall that the weighted path length is the sum of the frequencies of every node in the tree multiplied by its depth in the tree. It fulfills the following (recursive) equations:

\[
\begin{align*}
cost(\langle \rangle) &= 0 \\
cost(l, k, r) &= (\sum_{k \in \text{set_tree} \ l \ p \ k}) + \text{cost} \ l + p \ k + \text{cost} \ r + (\sum_{k \in \text{set_tree} \ r \ p \ k})
\end{align*}
\]

The difference of our task compared to finding an optimal Huffman encoding is the constraint that the resulting tree needs to be sorted, making it hard to deploy a similar greedy solution. Instead, we want to come up with a dynamic programming solution and thus need to find a way to subdivide the problem.
17.5.1 Deriving a Recursive Solution

The key insight into the problem is that subtrees of optimal binary search trees are also optimal. The left and right subtrees of the root must be optimal, since if we could improve either one, we would also get a better tree for the complete range of keys. This motivates the following definition:

\[
\begin{align*}
\text{wpl } W_{i,j} & = 0 \\
\text{wpl } W_{i,j} (l, k, r) & = \text{wpl } W_{i,k} (k - 1) l + \text{wpl } W_{k+1,j} r + W_{i,j} \\
W_{i,j} & = (\sum_k = i \ p \ k)
\end{align*}
\]

It is easy to see that \text{wpl } W_{i,j} is just a reformulation of \text{cost } t:

\[
\text{inorder } t = [i..j] \mapsto \text{wpl } W_{i,j} t = \text{cost } t
\]

We can actually forget about the original frequencies \( p \) and just optimize \text{wpl } W_{i,j} for some fixed weight function \( W :: \text{int } \Rightarrow \text{int } \Rightarrow \text{nat} \).

The binary search tree \( t \) that contains the keys \([i..j]\) and minimizes \text{wpl } W_{i,j} t has some root \( k \) with \([i..j] = [i..k - 1] @ k # [j + 1..k]\). Its left and right subtrees need to be minimal again, i.e. minimize \text{wpl } W_{i,k} (k - 1) and \text{wpl } W_{k+1,j} (k + 1) j. This yields the following recursive functions for computing the minimal weighted path length (\text{min\_wpl}) and a corresponding binary search tree (\text{opt\_bst}):

\[
\begin{align*}
\text{min\_wpl} :: \text{int } \Rightarrow \text{int } \Rightarrow \text{nat} \\
\text{min\_wpl } i \ j &= (\text{if } j < i \text{ then } 0 \text{ else } \text{min\_list} (\map (\lambda k. \text{min\_wpl } i \ (k - 1) + \text{min\_wpl } k + 1 \ j + W \ i \ j) [i..j]))) \\
\text{opt\_bst} :: \text{int } \Rightarrow \text{int } \Rightarrow \text{int } \text{tree} \\
\text{opt\_bst } i \ j &= (\text{if } j < i \text{ then } () \text{ else } \text{argmin} (\text{wpl } W \ i \ j) (\map (\lambda k. (\text{opt\_bst } i \ (k - 1), k, \text{opt\_bst } (k + 1) \ j)) [i..j]))
\end{align*}
\]

Here \text{argmin } f \ xs returns the rightmost \( x \in \text{set } xs \) such that \( f \ x \) is minimal among \( xs \) (i.e. \( f \ x < f \ y \) for all \( y \in \text{set } xs \)).

To prove that \text{min\_wpl} and \text{opt\_bst} are correct, we want to show two propositions: \text{min\_wpl } i \ j should be a lower bound of \text{wpl } W \ i \ j \ t for any search tree \( t \) for \([i..j]\),
and \( \text{min\_wpl} \ i \ j \) should correspond to the weight of an actual search tree, namely \( \text{opt\_bst} \ i \ j \). Formally, we prove the following propositions:

\[
\text{inorder} \ t = [i..j] \quad \rightarrow \quad \text{min\_wpl} \ i \ j \leq \text{wpl} \ W \ i \ j \ t
\]

\[
\text{inorder} \ (\text{opt\_bst} \ i \ j) = [i..j]
\]

\[
\text{wpl} \ W \ i \ j \ (\text{opt\_bst} \ i \ j) = \text{min\_wpl} \ i \ j
\]

The three propositions are easily proved by computation induction on \( \text{wpl} \), \( \text{opt\_bst} \) and \( \text{min\_wpl} \), respectively.

If \( W \) is constructed from \( p \) as above, we can derive the following correctness theorems referring to the original problem:

\[
\text{inorder} \ t = [i..j] \quad \rightarrow \quad \text{min\_wpl} \ W \ i \ j \leq \text{cost} \ t
\]

\[
\text{cost} \ (\text{opt\_bst} \ W \ i \ j) = \text{min\_wpl} \ W \ i \ j
\]

### 17.5.2 Memoization

We can apply the memoization techniques that were discussed above to efficiently compute \( \text{min\_wpl} \) and \( \text{opt\_bst} \). The only remaining caveat is that \( W \) also needs to be computed efficiently from the distribution \( p \). If we just use the defining equality \( W \ i \ j = (\sum_{k} = i \ p \ k) \), the computation of \( W \) is unnecessarily costly. Another way is to memoize \( W \) itself, using the following recursion:

\[
W \ p \ i \ j = (\text{if } i \leq j \text{ then } W \ p \ (j - 1) + p \ j \text{ else } 0)
\]

This yields a memoizing version \( W_{m}' \) and a theorem that connects it to \( W \):

\[
W \ p \ i \ j \triangleright W_{m}' \ p \ i \ j
\]

We can now iterate \( W_{m}' \ p \ i \ n \) for \( i = 0 \ldots n \) to pre-compute all relevant values of \( W \ p \ i \ j \):

\[
W_{c} \ p \ n = \text{snd} \ (\text{run\_state} \ (\text{map}_{m}' \ (\lambda i. \ W_{m}' \ p \ i \ n) \ [0..n]) \text{ empty})
\]

Using the correctness theorem for \( \text{map}_{m}' \) from above, it can easily be shown that this yields a consistent memory:

\[
\text{cmem} \ (W_{c} \ p \ n)
\]

We can show the following equation for computing \( W \)

\[
W \ p \ i \ j = (\text{case} \ (W_{c} \ p \ n) \ (i, \ j) \text{ of None } \Rightarrow W \ p \ i \ j \mid \text{Some } x \Rightarrow x)
\]

Note that the \( \text{None} \) branch will only be triggered when indices outside of \( 0 \ldots n \) are accessed. Finally, we can use \( W_{c} \) to pass the pre-computed values of \( W \) to \( \text{opt\_bst} \):
17.5.3 Optimizing the Recursion

While we have applied some trickery to obtain an efficient implementation of the simple dynamic programming algorithm expressed by \textit{opt\_bst}, we still have not arrived at the solution that is currently known to be most efficient. The most efficient known algorithm to compute optimal binary search trees due to Knuth [Knuth 1971] is a slight variation of \textit{opt\_bst} and relies on the following observation.

Let \( R_{i\ j} \) denote the maximal root of any optimal binary search for \([i..j]\):

\[
R_{i\ j} = \arg\min (\lambda k. \ w_{i\ j} + \text{min\_upl}_{i\ k-1} + \text{min\_upl}_{k+1\ j}) [i..j]
\]

It can be shown that \( R_{i\ j} \) is bounded by \( R_{i\ (j-1)} \) and \( R_{(i+1)\ j} \):

\[
i < j \rightarrow R_{i\ (j-1)} \leq R_{i\ j} \land R_{i\ j} \leq R_{(i+1)\ j}
\]

The proof of this fact is rather involved and the details can be found in the references provided at the end of this section.

With this knowledge, we can make the following optimization to \textit{opt\_bst}:

\[
\text{opt\_bst'} :: (\text{int} \Rightarrow \text{nat}) \Rightarrow \text{int} \Rightarrow \text{int} \Rightarrow \text{int\ tree}
\]

\[
\text{opt\_bst'} p i j \equiv
\]

\[
\begin{align*}
\text{let } M = W_c \ p j; \\
W = \lambda i\ j. \ \text{case } M (i, j) \text{ of } \text{None } \Rightarrow W p i j | \text{Some } x \Rightarrow x \\
\text{in } \text{opt\_bst } W i j
\end{align*}
\]

You may wonder whether this change really results in an asymptotic runtime improvement. Indeed, it can be shown that it improves the algorithm's runtime by a
factor of $O(n)$. For a fixed search tree size $d = i - j$, the total number of recursive computations is given by the following telescoping series:

$$d \leq n \rightarrow 
\sum j = d \cdot n \ \text{let} \ i = j - d \ \text{in} \ R (i + 1) j - R i (j - 1) + 1 \\
= R (n - d + 1) n - R 0 (d - 1) + n - d + 1$$

This quantity is bounded by $2 \cdot n$, which implies that the overall number of recursive calls is bounded by $O(n^2)$.

### 17.6 Chapter Notes

The original $O(n^2)$ algorithm for Binary Search Trees is due to Knuth [Knuth 1971]. Yao later explained this optimization more elegantly in his framework of "quadrilateral inequalities" [Yao 1980]. Nipkow and Somogyi follow Yao's approach in their Isabelle formalization [Nipkow and Somogyi 2018], on which the last subsection of this chapter is based. The other parts of this chapter are based on a paper by Wimmer et al. [Wimmer et al. 2018b] and its accompanying Isabelle formalization [Wimmer et al. 2018a]. The formalization also contains further examples of dynamic programming algorithms, including solutions for the Knapsack and the minimum edit distance problems, and the CYK algorithm.
Amortized Analysis

Consider a $k$-bit binary counter and a sequence of increment (by 1) operations on it where each one starts from the least significant bit and keeps flipping the 1s until a 0 is encountered (and flipped). Thus the worst-case running time of an increment is $O(k)$ and a sequence of $n$ increments takes time $O(nk)$. However, this analysis is very coarse: in a sequence of increments there are many much faster ones (for half of them the least significant bit is 0!). It turns out that a sequence of $n$ increments takes time $O(n)$. Thus the average running time of each increment is $O(1)$. Amortized analysis is the analysis of the running time of a sequence of operations on some data structure by upper-bounding the average running time of each operation.

As the example of the binary counter shows, the amortized running time for a single call of an operation can be much better than the worst-case time. Thus amortized analysis is unsuitable in a real-time context where worst-case bounds on every call of an operation are required.

Amortized analysis of some data structure is valid if the user of that data structure never accesses old versions of the data structure (although in a functional language one could). The binary counter shows why that invalidates amortized analysis: start from 0, increment the counter until all bits are 1, then increment that counter value again and again, without destroying it. Each of those increments takes time $O(k)$ and you can do that as often as you like, thus subverting the analysis. In an imperative language you can easily avoid this “abuse” by making the data structure stateful: every operation modifies the state of the data structure. This shows that amortized analysis has an imperative flavour. In a purely functional language, monads can be used to restrict access to the latest version of a data structure.

18.1 The Potential Method

The potential method is a particular technique for amortized analysis. The key idea is to define a potential function $\Phi$ from the data structure to non-negative numbers. The potential of the data structure is like a savings account that cheap calls can pay into (by increasing the potential) to compensate for later expensive calls (which decrease the potential). In a nutshell: the less “balanced” a data structure is, the higher its potential should be because it will be needed to pay for the impending restructuring.
The amortized running time (or complexity) is defined as the actual running time plus the difference in potential, i.e. the potential after the call minus the potential before it. If the potential increases, the amortized running time is higher than the actual running time and we pay the difference into our savings account. If the potential decreases, the amortized running time is lower than the actual running time and we take something out of our savings account to pay for the difference.

More formally, we are given some data structure with operations \( f, g \), etc with corresponding time functions \( T_f, T_g \) etc. We are also given a potential function \( \Phi \). The amortized running time function \( A_f \) for \( f \) is defined as follows:

\[
A_f s = T_f s + \Phi (f s) - \Phi s
\]  

(18.1)

where \( s \) is the data structure under consideration; \( f \) may also have additional parameters. Given a sequence of data structure states \( s_0, \ldots, s_n \) where \( s_{i+1} = f_i s_i \), it is not hard to see that

\[
\sum_{i=0}^{n-1} A_f s_i = \sum_{i=0}^{n-1} T_f s_i + \Phi s_n - \Phi s_0
\]

If we assume (for simplicity) that \( \Phi s_0 = 0 \), then it follows immediately that the amortized running time of the whole sequence is an upper bound of the actual running time (because \( \Phi \) is non-negative). This observation becomes useful if we can bound \( A_f s \) by some closed term \( u_f s \). Typical examples for \( u_f s \) are constants, logarithms etc. Then we can conclude that \( f \) has constant, logarithmic etc amortized complexity. Thus the only proof obligation is

\[
A_f s \leq u_f s
\]

possibly under the additional assumption \texttt{invar} \( s \) if the data structure comes with an invariant \texttt{invar}.

In the sequel we assume that \( s_0 \) is some fixed value, typically "empty", and that its potential is 0.

How do we analyze operations that combine two data structures, e.g. the union of two sets? Their amortized complexity can be defined in analogy to (18.1):

\[
A_f s_1 s_2 = T_f s_1 s_2 + \Phi (f s_1 s_2) - (\Phi s_1 + \Phi s_2)
\]

So far we implicitly assumed that all operations return the data structure as a result, otherwise \( \Phi (f s) \) does not make sense. How should we analyze so-called observer functions that do not modify the data structure but return a value of some other type? Amortized analysis does not make sense here because the same observer can be applied multiple times to the same data structure value without modifying it. Classical worst-case complexity is needed, unless the observer does modify the data structure as a side effect or by returning a new value. Then one can perform an amortized analysis that ignores the returned observer value (but not the time it takes to compute it).
Now we study two important examples of amortize analyses. More complex applications are found in later chapters.

18.2 Binary Counter

The binary counter is represented by a list of Booleans where the head of the list is the least significant bit. The increment operation and its running time are easily defined:

\[
\text{incr} :: \text{bool list} \Rightarrow \text{bool list} \\
\text{incr} [] = [\text{True}] \\
\text{incr} (\text{False} \# bs) = \text{True} \# bs \\
\text{incr} (\text{True} \# bs) = \text{False} \# \text{incr} bs
\]

\[
T_{\text{incr}} :: \text{bool list} \Rightarrow \text{real} \\
T_{\text{incr}} [] = 1 \\
T_{\text{incr}} (\text{False} \# _) = 1 \\
T_{\text{incr}} (\text{True} \# bs) = T_{\text{incr}} bs + 1
\]

The potential of a counter is the number of True’s because they increase \( T_{\text{incr}} \):

\[
\Phi :: \text{bool list} \Rightarrow \text{real} \\
\Phi bs = |\text{filter} (\lambda x. x) bs|
\]

Clearly the potential is never negative.

The amortized complexity of \( \text{incr} \) is 2:

\[
T_{\text{incr}} bs + \Phi (\text{incr} bs) - \Phi bs = 2
\]

This can be proved automatically by induction on \( bs \).

18.3 Dynamic Tables

A dynamic table is an abstraction of a dynamic array that can grow and shrink subject to a specific memory management. At any point the table has a certain size (the number of cells) but some cells may be unoccupied or free. As long as there are free cells, inserting a new element into the table takes constant time. When the table overflows, the whole table has to be copied into a larger table, which takes linear time. Similarly, elements can be deleted from the table in constant time, but when too many elements have been deleted, the table is contracted to save space. Contraction involves copying into a smaller table. This is an abstraction of a dynamic array, where the index
bounds can grow and shrink. It is an abstraction because we ignore the actual contents of the table and abstract the table to a pair \((n, l)\) where \(l\) is its size and \(n < l\) the number of occupied cells. The empty table is represented by \((0, 0)\).

Below we do not comment on the formal proofs because they are essentially just case analyses (as dictated by the definitions) plus linear arithmetic.

### 18.3.1 Insertion

The key observation is that doubling the size of the table upon overflow leads to an amortized cost of 3 per insertion: 1 for inserting the element, plus 2 towards the later cost of copying a table of size \(l\) upon overflow (because only the \(l/2\) elements that lead to the overflow pay for it).

Insertion always increments \(n\) by 1. The size increases from 0 to 1 with the first insertion and doubles with every further overflow:

\[
\text{ins} :: \text{nat} \times \text{nat} \Rightarrow \text{nat} \times \text{nat} \\
\text{ins} (n, l) = \begin{cases} 
(n + 1, \text{if } n < l \text{ then } l \text{ else if } l = 0 \text{ then } 1 \text{ else } 2 \cdot l) 
\end{cases}
\]

\[
\text{T}_\text{ins} :: \text{nat} \times \text{nat} \Rightarrow \text{real} \\
\text{T}_\text{ins} (n, l) = (\text{if } n < l \text{ then } 1 \text{ else } n + 1)
\]

This guarantees the load factor \(n/l\) is always between 1/2 and 1:

\[
\text{invar} :: \text{nat} \times \text{nat} \Rightarrow \text{bool} \\
\text{invar} (n, l) = (l/2 \leq n \land n \leq l)
\]

The potential of a table \((n, l)\) is \(2 \cdot (n - l/2) = 2 \cdot n - l\) following the intuitive argument at the beginning of the Insertion section.

\[
\Phi :: \text{nat} \times \text{nat} \Rightarrow \text{real} \\
\Phi (n, l) = 2 \cdot n - l
\]

The potential is always non-negative because of the invariant.

Note that in our informal explanatory text we use / freely and assume we are working with real numbers. In the formalization we often prefer multiplication over division because the former is easier to reason about.
18.3 Dynamic Tables

18.3.2 Insertion and Deletion

A naive implementation of deletion simply removes the element but never contracts the table. This works (Exercise 18.2) but is a waste of space. It is tempting to think we should contract once the load factor drops below $1/2$. However, this can lead to the following fluttering. Starting with a full table (of size $l = 2^k$ for an arbitrary $k$) one insertion causes an overflow, two deletions cause a contraction, another insertion causes an overflow, and so on. The cost of each overflow and contraction is $l$ but there are at most two operations to pay for it. Thus the amortized cost of both insertion and deletion cannot be constant. It turns out that it works if we allow the load factor to drop to $1/4$ before we contract the table to half its size:

\[
\text{del} :: \text{nat} \times \text{nat} \Rightarrow \text{nat} \times \text{nat}
\]
\[
\text{del} (n, l) = (n - 1, \text{if } n = 1 \text{ then } 0 \text{ else if } 4 \cdot (n - 1) < l \text{ then } l \div 2 \text{ else } l)
\]

\[
T_{\text{del}} :: \text{nat} \times \text{nat} \Rightarrow \text{real}
\]
\[
T_{\text{del}} (n, l) = (\text{if } n = 1 \text{ then } 1 \text{ else if } 4 \cdot (n - 1) < l \text{ then } n \text{ else } 1)
\]

Now the load factor is always between $1/4$ and $1$. It turns out that the lower bound is not needed in the proofs and we settle for a simpler invariant:

\[
invar :: \text{nat} \times \text{nat} \Rightarrow \text{bool}
\]
\[
invar (n, l) = (n \leq l)
\]

The potential distinguishes two cases:

\[
\Phi :: \text{nat} \times \text{nat} \Rightarrow \text{real}
\]
\[
\Phi (n, l) = (\text{if } n < l/2 \text{ then } l/2 - n \text{ else } 2 \cdot n - l)
\]

The condition $2 \cdot n \geq l$ concerns the case when we are heading up for an overflow and has been dealt with above. Conversely, $2 \cdot n < l$ concerns the case where we are heading down for a contraction. That is, we start at $(l, 2 \cdot l)$ (where the potential is $0$) and $l/2$ deletions lead to $(l/2, 2 \cdot l)$ where a contraction requires $l/2$ credits, and indeed $\Phi (l/2, 2 \cdot l) = l/2$. Since $l/2$ is spread over $l/2$ deletions, the amortized cost of a single deletion is $2$, $1$ for the real cost and $1$ for the savings account.

Note that the case distinction in the definition of $\Phi$ ensures that the potential is always $\geq 0$ — the invariant is not even needed.
18.4 Exercises

Exercise 18.1. Generalize the binary counter to a base $b$ counter, $b \geq 2$. Prove that there is a constant $c$ such that the amortized complexity of incrementation is at most $c$ for every $b \geq 2$.

Exercise 18.2. Prove that in the dynamic table with naive deletion (where deletion decrements $n$ but leaves $l$ unchanged), insertion has an amortized cost of at most 3 and deletion of at most 1.

Exercise 18.3. Modify deletion as follows. Contraction happens when the load factor would drop below $1/3$, i.e. when $3 \cdot (n - 1) < l$. Then the size of the table is multiplied by $2/3$, i.e. reduced to $(2 \cdot l) \div 3$. Prove that insertion and deletion have constant amortized complexity using the potential $\Phi(n, l) = |2 \cdot n - l|$.

18.5 Chapter Notes

Amortized analysis is due to Tarjan [1985]. Introductions to it can be found in most algorithm textbooks. This chapter is based on work by Nipkow [2015] and Nipkow and Brinkop [2019] which also formalizes the meta-theory of amortized analysis.
19

Queues

Alejandro Gómez-Londoño and Tobias Nipkow

19.1 Queue Specification

A queue can be viewed as a glorified list with function \textsl{enq} for adding an element to the end of the list and function \textsl{first} for accessing and \textsl{deq} for removing the first element. This is the full ADT:

\begin{verbatim}
ADT Queue =
interface  empty :: 'q
    enq :: 'a \Rightarrow 'q \Rightarrow 'q
    deq :: 'q \Rightarrow 'q
    first :: 'q \Rightarrow 'a
    is_empty :: 'q \Rightarrow bool
abstraction list :: 'q \Rightarrow 'a list
invariant invar :: 'q \Rightarrow bool
specification list empty = []
invar q \rightarrow list (enq x q) = list q @ [x]
invar q \rightarrow list (deq q) = tl (list q)
invar q \land list q \neq [] \rightarrow first q = hd (list q)
invar q \rightarrow is_empty q = (list q = [])
invar empty
invar q \rightarrow invar (enq x q)
invar q \rightarrow invar (deq q)
\end{verbatim}

A trivial implementation is as a list, but then \textsl{enq} is linear in the length of the queue. To improve this we consider two more sophisticated implementations. First, a simple implementation where every operation has amortized constant complexity. Second, a tricky “real time” implementation where every operation has worst-case constant complexity.

19.2 Queues as Pairs of Lists

The queue is implemented as a pair of lists \((fs, rs)\), the front and rear lists. Function \textsl{enq} adds elements to the head of the rear \(rs\) and \textsl{deq} removes elements from the head.
of the front \(fs\). When \(fs\) becomes empty, it is replaced by \(rev\ \rs\) (and \(rs\) is emptied) — the reversal ensures that now the oldest element is at the head. Hence \(rs\) is really the reversal of the rear of the queue but we just call it the rear. The abstraction function is obvious:

\[
\text{list} :: \text{'a list} \times \text{'a list} \Rightarrow \text{'a list} \\
\text{list} (fs, rs) = fs @ rev rs
\]

Clearly \(enq\) and \(deq\) are constant-time until the front becomes empty. Then we need to reverse the rear which takes linear time (if it is implemented by \(itrev\), see Section 1.5.1). But we can pay for this linear cost up front by paying a constant amount for each call of \(enq\). Thus we arrive at amortized constant time. See below for the formal treatment.

The implementation is shown in Figure 19.1. Of course \(empty = ([], [])\). Function \(norm\) performs the reversal of the rear once the front becomes empty. Why does not only \(deq\) but also \(enq\) call \(norm\)? Because otherwise \(enq\ \ x_n \ (\ldots (enq \ x_1 \ empty) ...)\) would result in \(([], [x_n, \ldots, x_1])\) and \(first\) would become an expensive operation because

\[
\begin{align*}
\text{norm} :: \text{'a list} \times \text{'a list} & \Rightarrow \text{'a list} \times \text{'a list} \\
\text{norm} (fs, rs) &= (\text{if } fs = [] \text{ then } (\text{rev } rs, []) \text{ else } (fs, rs)) \\
\text{enq} :: \text{'a} \Rightarrow \text{'a list} \times \text{'a list} \Rightarrow \text{'a list} \times \text{'a list} \\
\text{enq} \ a (fs, rs) &= \text{norm} (fs, a @ rs) \\
\text{deq} :: \text{'a list} \times \text{'a list} \Rightarrow \text{'a list} \times \text{'a list} \\
\text{deq} (fs, rs) &= (\text{if } fs = [] \text{ then } (fs, rs) \text{ else } \text{norm} (\text{tl } fs, rs)) \\
\text{first} :: \text{'a list} \times \text{'a list} \Rightarrow \text{'a} \\
\text{first} (a @ _, _) &= a \\
\text{is_empty} :: \text{'a list} \times \text{'a list} \Rightarrow \text{bool} \\
\text{is_empty} (fs, _) &= (fs = [])
\end{align*}
\]
it would require the reversal of the rear. Thus we need to avoid queues \([\text{[]} \rightarrow rs\) where \(rs \neq \text{[]}\). Thus \(norm\) guarantees the following invariant:

\[
\text{invar :: 'a list \times 'a list \to bool}
\]
\[
\text{invar (fs, rs) = (fs = \text{[]} \longrightarrow rs = \text{[]})}
\]

Functional correctness, i.e. proofs of the properties in the ADT \(Queue\), are straightforward. Let us now turn to the amortized running time analysis. The time functions are shown in Appendix B.8.

For the amortized analysis we define the potential function

\[
\Phi :: 'a list \times 'a list \to \text{nat}
\]
\[
\Phi (_, rs) = |rs|
\]

because \(|rs|\) is the amount we have accumulated by charging 1 for each \(enq\). This is enough to pay for the eventual reversal. Now it is easy to prove that both \(enq\) and \(deq\) have amortized constant running time:

\[
T_{\text{enq}} a (fs, rs) + \Phi (enq a (fs, rs)) - \Phi (fs, rs) \leq 2
\]
\[
T_{\text{deq}} (fs, rs) + \Phi (deq (fs, rs)) - \Phi (fs, rs) \leq 1
\]

The two observer functions \(\text{first}\) and \(\text{is_empty}\) have constant running time.

**Exercise 19.1.** A min-queue is a queue that supports an operation \(\text{min}_q\) that returns the minimal element in the queue. Formally, the ADT \(Queue\) is extended as follows: we assume \(\text{'}a :: \text{linorder}\), extend the interface with \(\text{min}_q :: \text{'}q \Rightarrow \text{'}a\) and the specification with

\[
\text{invar q \wedge list q \neq [] \to min}_q q = \text{Min (set (list q))}
\]

Implement and verify a min-queue with amortized constant time operations. Hint: follow the pair-of-lists idea above but store additional information that allows you to return the minimal element in constant time.

**19.3 A Real Time Implementation**

This section presents the Hood-Melville queue, a tricky implementation that improves upon the representation in the previous section by preemptively performing reversals over a number of operations before they are required.
19.3.1 Stepped Reversal

Breaking down a reversal operation into multiple steps can be done using the following function:

\[
\text{rev\_step} :: 'a list \times 'a list \Rightarrow 'a list \times 'a list
\]
\[
\text{rev\_step} (x \# xs, ys) = (xs, x \# ys)
\]
\[
\text{rev\_step} ([], ys) = ([], ys)
\]

where \(x \# xs\) is the list being reversed, and \(x \# ys\) is the partial reversal result. Thus, to reverse a list of size 3 one should call \text{rev\_step} 3 times:

\[
\text{rev\_step} ([1, 2, 3], []) = ([2, 3], [1])
\]
\[
\text{rev\_step} (\text{rev\_step} ([1, 2, 3], [])) = ([3], [2, 1])
\]
\[
\text{rev\_step} (\text{rev\_step} (\text{rev\_step} ([1, 2, 3], []))) = ([], [3, 2, 1])
\]

Note that each call to \text{rev\_step} takes constant time since its definition is non-recursive.

Using the notation \(f^n\) for the \(n\)-fold composition of function \(f\) we can state a simple inductive lemma:

**Lemma 19.1.** \(\text{rev\_step}^n (xs, ys) = ([], \text{rev} \ xs @ ys)\)

As a special case this implies \(\text{rev\_step}^n (xs, []) = ([], \text{rev} \ xs)\).

19.3.2 A Real Time Intuition

Hood-Melville queues are similar to those presented in Section 19.2 in that they use a pair of lists \((f, r)\) (front and rear — for succinctness we drop the s’s now) to achieve constant running time \(\text{deq}\) and \(\text{enq}\). However, they avoid a costly reversal operation once \(f\) becomes empty by preemptively computing a new front \(fr = f @ \text{rev} \ r\) one step at a time using \text{rev\_step} as enqueueing and dequeueing operations occur. The process that generates \(fr\) consists of three phases:

1. Reverse \(r\) to form \(r'\), which is the tail end of \(fr\)
2. Reverse \(f\) to form \(f'\)
3. Reverse \(f'\) onto \(r'\) to form \(fr\)

All three phases can be described in terms of \text{rev\_step} as follows:

1. \(r' = \text{snd} (\text{rev\_step}^1 (r, []))\)
2. \(f' = \text{snd} (\text{rev\_step}^1 (f, []))\)
3. \(fr = \text{snd} (\text{rev\_step}^1 (f', r'))\)
Phases (1) and (2) are independent and can be performed at the same time, hence, when starting from this configuration

\[
\begin{array}{cccc}
  f & f' & r & r' \\
  q_0 & \ldots & q_m & q_{m+1} & \ldots & q_n \\
\end{array}
\]

after \( \text{max} |f| |r| \) steps of reversal the state would be the following:

\[
\begin{array}{cccc}
  f & f' & r & r' \\
  q_m & \ldots & q_0 & q_{m+1} & \ldots & q_n \\
\end{array}
\]

Phase (3) reverses \( f' \) onto \( r' \) \( fr \) to obtain the same result as a call to \textit{list}:

\[
fr = \text{snd} (\text{rev}_\text{step} f'[[f', r']]) \quad \text{by definition of} \ fr
\]
\[
= \text{rev} f' @ r' \quad \text{using Lemma 19.1}
\]
\[
= \text{rev} f' @ \text{snd} (\text{rev}_\text{step} r'[[r, []]]) \quad \text{by definition of} \ r'
\]
\[
= \text{rev} f' @ \text{rev} r \quad \text{using Lemma 19.1}
\]
\[
= \text{rev} (\text{snd} (\text{rev}_\text{step} f'[[f, []]])) @ \text{rev} r' \quad \text{by definition of} \ f'
\]
\[
= \text{rev} (\text{rev} f) @ \text{rev} r \quad \text{using Lemma 19.1}
\]
\[
= f @ \text{rev} r \quad \text{by rev involution}
\]

The resulting front list \( fr \) contains all elements previously in \( f \) and \( r \):

\[
\begin{array}{cccc}
  \text{fr} & \\
  q_0 & \ldots & q_m & q_{m+1} & \ldots & q_n \\
\end{array}
\]

\[
\begin{array}{ccc}
  f & \text{rev} r \\
\end{array}
\]

A Hood-Melville queue spreads all reversal steps across queue-altering operations requiring careful bookkeeping. To achieve this gradual reversal, additional lists \textit{front} and \textit{rear} are used for enqueuing and dequeuing, while internal operations rely only on \( f, f', r, \) and \( r' \). At the start of the reversal process \( \text{rear} \) is copied into \( r \) and emptied; similarly, \( \text{front} \) is copied into \( f \), but its contents are kept as they might need to be dequeued. Moreover, to avoid using elements from \( f \) or \( f' \) that may have been removed from \( \text{front} \), a counter \( d \) records the number of dequeuing operations that have occurred since the reversal process started; this way, only \(|f| - d\) elements are appended into \( r \) to form \( fr \). Once the reversal finishes \( fr \) become the new \( \text{front} \) and the internal lists are cleared. When the queue is not being reversed all operations are performed in a manner similar to previous implementations. The configuration of a queue at the beginning of the reversal process is as follows:
19.3.3 The Reversal Strategy

A crucial detail of this implementation is determining at which point the reversal process should occur. The strategy is to start once \( |\text{rear}| \) becomes larger than \( |\text{front}| \), and ensure that all reversal steps are done before \( \text{front} \) runs out of elements or \( \text{rear} \) becomes larger than the new front (\( \text{fr} \)).

With this strategy, once \( |\text{rear}| = n+1 \) and \( |\text{front}| = n \), the reversal processes start. The first two phases take \( n + 1 \) steps (\( \max |\text{front}| |\text{rear}| \)) to generate \( f' \) and \( r' \), and the third phase produces \( \text{fr} \) in \( n \) steps. A complete reversal takes \( 2 \cdot n + 1 \) steps. Because the queue can only perform \( n \) \( \text{deq} \) operations before \( \text{front} \) is exhausted, \( 2 \cdot n + 1 \) steps must be performed in at most \( n \) operations. This can be achieved by performing the first two steps in the operation that causes \( \text{rear} \) to become larger than \( \text{front} \) and two more steps in each subsequent operation. Therefore, \( 2 \cdot (n + 1) \) steps can occur before \( \text{front} \) is emptied, allowing the reversal process to finish in time.

Finally, since at most \( n \) \( \text{enq} \) or \( \text{deq} \) operations can occur during reversal, the largest possible \( \text{rear} \) has length \( n \) (only \( \text{enq} \) ops), while the smallest possible \( \text{fr} \) has length \( n + 1 \) (only \( \text{deq} \) ops). Thus, after the reversing process has finished the new front (\( \text{fr} \)) is always larger than \( \text{rear} \).

19.3.4 Implementation

Queues are implemented using the following record type:

\[
\text{record } 'a\ \text{queue} = \begin{array}{l}
\text{lenf} :: \text{nat} \\
\text{front} :: 'a\ \text{list} \\
\text{status} :: 'a\ \text{status} \\
\text{rear} :: 'a\ \text{list} \\
\text{lenr} :: \text{nat}
\end{array}
\]
In a nutshell, a record is a product type with named fields and "built-in" construction, selection, and update operations. Values of 'a queue are constructed using make :: nat ⇒ 'a list ⇒ 'a status ⇒ 'a list ⇒ nat ⇒ 'a queue were each argument corresponds to one of the fields of the record in canonical order. Additionally, given a queue q we can obtain the value in field front with \( q; front := [] \), and update its content using \( q; (j; front := [], rear := []; \)\. Multiple updates can be composed as \( q; (j; front := [], rear := []; \). All values in the queue along with its internal state are stored in the various fields of 'a queue. Fields front and rear contain the lists over which all queue operations are performed. The length of front and rear is recorded in lenf and lenr (respectively) to avoid calling length whose complexity is not constant. Finally, status tracks the current reversal phase of the queue in a 'a status value.

```haskell
datatype 'a status =
  Idle |
  Rev nat ('a list) ('a list) ('a list) ('a list) |
  App nat ('a list) ('a list) |
  Done
```

Each value of 'a status represents either a phase of reversal or the queue's normal operation. Constructor Idle signals that no reversal is being performed. Status Rev ok f f' r r' corresponds to phases (1) and (2) where the lists f, f', r, and r' are used for the reversal steps of the front and the rear. The App ok f' r' case corresponds to phase (3) where both lists are appended to form the new front (fr). In both App and Rev, the first argument ok :: nat keeps track of the number of elements in f' that have not been removed from the queue, effectively ok = |f| - d, where d is the number of deq operations that have occurred so far. Lastly, Done fr marks the end of the reversal process and contains only the new front list fr.

In the implementation, all of the steps of reversal operations in the queue are performed by functions exec and invalidate; they ensure at each step that the front list being computed is kept consistent w.r.t. the contents and operations in the queue.

Function exec :: 'a status ⇒ 'a status performs the incremental reversal of the front list by altering the queue's status one step at a time in accordance with the reversal phases. Following the strategy described in Section 19.3.3, all queue operations call exec twice to be able to finish the reversal in time. On Idle queues exec has no effect. The implementation of exec is an extension of rev_step with specific considerations for each status value and is defined as follows:
If the status is $Rev \ ok \ f' \ r'$, then $exec$ performs two (or one if $f = []$) simultaneous reversal steps from $f$ and $r$ into $f'$ and $r'$; moreover $ok$ is incremented if a new element has been added to $f'$. Once $f$ is exhausted and $r$ is a singleton list, the remaining element is moved into $r'$ and the status is updated to the next phase of reversal. In the $App \ ok \ f' \ r'$ phase, $exec$ moves elements from $f'$ to $r'$ until $ok = 0$, at which point $r'$ becomes the new front by transitioning into $Done \ r'$. In all other cases $exec$ behaves as the identity function. As is apparent from its implementation, a number of assumptions are required for $exec$ to function properly and eventually produce $Done$. These assumption are discussed in Section 19.3.5.

If an element is removed from the queue during the reversal process, it also needs to be removed from the new front list ($fr$) being computed. Function $invalidate$ is used to achieve this:

By decreasing the value of $ok$, the number of elements from $f'$ that are moved into $r'$ in phase (3) is reduced, since $exec$ might produce $Done$ early, once $ok = 0$, ignoring the remaining elements of $f'$. Furthermore, since $f'$ is a reversal of the front list, elements left behind in its tail correspond directly to those being removed from the queue.

The rest of the implementation is shown below. Auxiliary function $exec2$, as its name suggests, applies $exec$ twice and updates the queue accordingly if $Done$ is returned.
19.3 A Real Time Implementation

```
exec2 q = (case exec (exec (status q)) of
        Done fr ⇒ q[status = Idle, front = fr] |
        newstatus ⇒ q[status = newstatus])

check :: 'a queue ⇒ 'a queue
check q
= (if lenr q ≤ lenf q then exec2 q
   else let newstate = Rev 0 (front q) [] (rear q) []
        in exec2
             (q[lenf := lenf q + lenr q, status := newstate, rear := [],
              lenr := 0]))

empty :: 'a queue
empty = make 0 [] Idle [] 0

first :: 'a queue ⇒ 'a
first q = hd (front q)

enq :: 'a ⇒ 'a queue ⇒ 'a queue
enq x q = check (q[rear := x # rear q, lenr := lenr q + 1])

deq :: 'a queue ⇒ 'a queue
deq q
= check
   (q[lenf := lenf q - 1, front := tl (front q),
    status := invalidate (status q)])
```

The two main queue operations, `enq` and `deq`, alter `front` and `rear` as expected, with additional updates to `lenf` and `lenr` to keep track of the their length. To perform all “internal” operations, both functions call `check`. Additionally, `deq` uses `invalidate` to mark elements as removed.

Function `check` calls `exec2` if `lenr` is not larger than `lenf`. Otherwise a reversal process is initiated: `rear` is emptied and `lenr` is set to 0; `lenf` is increased to the size of the whole queue since, conceptually, all elements are now in the soon-to-be-computed front; the status `newstate` is initialized as described at the beginning of Section 19.3.2.
The time complexity of this implementation is clearly constant, since there are no recursive functions.

19.3.5 Functional Correctness

To show this implementation is an instance of the ADT Queue, we need a number of invariants to ensure the consistency of a queue values are preserved by all operations.

Initially, as hinted by the definition of exec, values of type a status should have specific properties to guarantee a Done result after a (finite) number of calls to exec. The predicate inv_st defines these properties as follows:

\[
\text{inv}_\text{st} :: 'a \text{ status} \Rightarrow \text{bool}
\]

\[
\text{inv}_\text{st} (\text{Rev } \text{ok } f \, f' \, r \, r') = ([f] + 1 = |r| \land |f'| = |r'| \land \text{ok} \leq |f'|)
\]

\[
\text{inv}_\text{st} (\text{App } \text{ok } f' \, r') = (\text{ok} \leq |f'| \land |f'| < |r'|)
\]

\[
\text{inv}_\text{st} \, \text{Idle} = \text{True}
\]

\[
\text{inv}_\text{st} (\text{Done } \_ \_) = \text{True}
\]

First, \text{inv}_\text{st} ensures for pattern, \text{Rev } \text{ok } f \, f' \, r \, r' that arguments \text{f} and \text{r} follow the reversal strategy, and counter \text{ok} is only ever increased as elements are added to \text{f}'. Similarly, for \text{App } \text{ok } f' \, r', it must follow that \text{r'} remains larger than \text{f'}, and \text{|f'|} provides an upper bound for \text{ok}. All other patterns trivially fulfill the invariant.

The queue invariant invar is an extension of \text{inv}_\text{st} and considers all the other fields in the queue:

\[
invar :: 'a \text{ queue} \Rightarrow \text{bool}
\]

\[
invar \, q = \left(\text{lenf } q = |\text{front_list } q| \land \text{lenr } q = |\text{rear_list } q| \land \text{lenr } q \leq \text{lenf } q \land \left(\begin{cases}
\text{case } \text{status } q \text{ of} \\
\text{Rev } \text{ok } f \, f' \, r \, r' \Rightarrow \\
\quad 2 \cdot \text{lenr } q \leq |f| \land \text{ok} \neq 0 \land 2 \cdot |f| + \text{ok} + 2 \leq 2 \cdot |\text{front } q| \\
\quad |\text{App } \text{ok } f' \, r' \Rightarrow 2 \cdot \text{lenr } q \leq |r| \land \text{ok} + 1 \leq 2 \cdot |\text{front } q| \\
\quad |\_ \Rightarrow \text{True} ) \land \\
\quad (\exists \text{rest. } \text{front_list } q = \text{front } q @ \text{rest}) \land \\
\quad (\exists \text{fr. } \text{status } q = \text{Done } \text{fr}) \land \\
\quad \text{inv}_\text{st} (\text{status } q))
\right)
\]

The condition \text{lenr } q = |\text{rear_list } q| ensures \text{lenr} is equal to the length of the queue's rear, where function \text{rear_list } q, defined as \text{(rev } \circ \text{rear} ) q, produces the rear list in canonical order. Likewise, \text{lenf } q = |\text{front_list } q| matches \text{lenf} to the queue's front.
However, function \texttt{front\_list} warrants special attention as it must compute the list representing the front of the queue even during a reversal:

\begin{verbatim}
front_list :: 'a queue \Rightarrow 'a list
front_list q = (case status q of
  Idle \Rightarrow front q |
  Rev ok f f' r r' \Rightarrow rev (take ok f') \circ f \circ rev r \circ r' |
  App ok f' x \Rightarrow rev (take ok f') \circ x |
  Done f \Rightarrow f)
\end{verbatim}

For case \texttt{App ok f' r'}, the front list corresponds to the final result of the stepped reversal (19.1) but only elements in \( f' \) that are still in the queue, denoted by \texttt{take ok f'}, are considered. Analogously for \texttt{Rev ok f f' r r'}, both stepped reversal results are appended and only relevant elements in \( f' \) are used, however, rear lists \( r \) and \( r' \) are reversed again to achieve canonical order.

Continuing with \texttt{invar}, inequality \( \text{lenr} q \leq \text{lenf} q \) is the main invariant in our reversal strategy, and by the previous two equalities must hold even as internal operations occur. Furthermore, \( \exists \text{rest}. \) \texttt{front\_list q = front q \circ rest} ensures \texttt{front q} is contained within \texttt{front\_list q}, thus preventing any mismatch between the internal state and the queue’s front. Given that \texttt{exec2} is the only function that manipulates a queue’s \texttt{status}, it holds that \( \exists \text{fr.} \) \texttt{status q = Done fr} since any internal \texttt{Done} result is replaced by \texttt{Idle}.

The case distinction on \texttt{status q} places size bounds on internal lists \texttt{front} and \texttt{rear} ensuring the front does not run out of elements and the rear never grows beyond \( \text{lenr} q \leq \text{lenf} q \). In order to clarify some of formulations used in this part of \texttt{invar}, consider the following correspondences, which hold once the reversal process starts:

- \texttt{lenr q} corresponds to the number of \texttt{enq} operations performed so far, and \( 2 \cdot \text{lenr} q \) denotes the \texttt{exec} applications in those operations.
- \(|\text{front} q|\) corresponds to the number of \texttt{deq} operations that can be performed before \texttt{front q} is exhausted. Therefore, \( 2 \cdot |\text{front} q|\) is the minimum number of \texttt{exec} applications the queue will be able to do at any given point.
- On \texttt{Rev ok f f' r r'} status, \(|f'|\) corresponds to the number of \texttt{exec}'s performed so far and the internal length of front being constructed. Expression \(|r|\) is the analogous for a \texttt{App ok f r}.
- From a well formed \texttt{App ok f r} it takes \( ok + 1 \) applications of \texttt{exec} to reach \texttt{Done}. Since, the base case of \texttt{App} is obtained after \( ok \) applications, and the transition into \texttt{Done} takes an extra step.
• From a well formed $\text{Rev } ok \ f \ f' \ r \ r'$ it takes $2 \cdot |f'| + ok + 2$ applications of $\text{exec}$ to reach $\text{Done}$. Since, the base case of $\text{Rev}$ is obtained after $|f'|$ applications (incrementing $ok$ by the same amount), the transitioning into $\text{App}$ takes one step, and $ok + |f'|$ extra steps are need to reach $\text{Done}$ from $\text{App}$.

In the $\text{Rev } ok \ f \ f' \ r \ r'$ case, $2 \cdot \text{lenr } q \leq |f'|$ ensures $f'$ grows larger with every $\text{enq}$ operation and the internal list is at least twice the length of the queue's rear. Additionally, the value of $ok$ cannot be 0 as this either marks the beginning of a reversal which calls $\text{exec2}$ immediately, or signals that elements in $\text{front } q$ have run out. Finally, to guarantee the reversal process can finish before the $\text{front } q$ is exhausted the number of $\text{exec}$ applications before reaching $\text{Done}$ must be less than the minimum number of applications possible, denoted by $2 \cdot |f| + ok + 2 \leq 2 \cdot |\text{front } q|$.

Case $\text{App } ok \ f \ r$ has similar invariants, with equation $2 \cdot \text{lenr } q \leq |r|$ bounding the growth of $r$ as it was previously done with $f'$. Moreover, $ok + 1 \leq 2 \cdot |\text{front } q|$ ensures $\text{front } q$ is not exhausted before the reversal is completed.

With the help of $\text{invar}$ and this abstraction function

```plaintext
list :: 'a queue => 'a list
list q = front_list q @ rear_list q
```

all properties of the $\text{Queue}$ ADT can be proved. The proofs are mostly by cases on the $\text{status}$ field followed by reasoning about lists. It is essential that the invariant characterizes all cases precisely.

### 19.4 Chapter Notes

The representation of queues as pairs of lists is due to Burton [1982]. Hood-Melville queues are due to Hood and Melville [1981]. The implementation is based on the presentation by Okasaki [1998, Sect. 8.2.1.].

The idea underlying Hood-Melville queues can be generalized to double-ended queues. This was explained by Hood [1982, Sect. 4.2], rediscovered in more detail by Chuang and Goldberg [1993] and verified by Tóth and Nipkow [2023].

Okasaki [1998] shows how both single and double-ended real-time queues can be defined more simply with the help of lazy evaluation. However, reasoning about the running time under lazy evaluation is nontrivial, as the verification by Pottier et al. [2024] of the amortized running time of some queue implementations shows.
20

Splay Trees

Tobias Nipkow

Splay trees are fascinating self-organizing search trees. Self-organizing means that the tree structure is modified upon access (including \texttt{isin} queries) to improve the performance of subsequent operations. Concretely, every splay tree operation moves the element concerned to the root. Thus splay trees excel in applications where a small fraction of the entries are the targets of most of the operations. In general, splay trees perform as well as any static binary search tree.

Splay trees have two drawbacks. First, their performance guarantees (logarithmic running time of each operation) are only amortized. Self-organizing does not mean self-balancing: splay trees can become unbalanced, in contrast to, for example, red-black trees. Second, because \texttt{isin} modifies the tree, splay trees are less convenient to use in a purely functional language.

20.1 Implementation

The central operation on splay trees is the \texttt{splay} function shown in Figure 20.1. It rotates the given element \(x\) to the root of the tree if \(x\) is already in the tree. Otherwise the last element found before the search for \(x\) hits a leaf is rotated to the root.

Function \texttt{isin} has a trivial implementation in terms of \texttt{splay}:

\begin{verbatim}
isin : 'a tree \Rightarrow 'a \Rightarrow bool
isin t x = (case splay x t of \{} \Rightarrow False | \_, a, \_ \Rightarrow x = a)
\end{verbatim}

Except that \texttt{splay} creates a new tree that needs to be returned from a proper \texttt{isin} as well to achieve the amortized logarithmic complexity (see the discussion of observer functions at the end of Section 18.1). This is why splay trees are inconvenient in functional languages. For the moment we ignore this aspect and stick with the above \texttt{isin} because it has the type required by the \texttt{Set ADT}.

The implementation of \texttt{insert} \(x\) \(t\) in Figure 20.2 is straightforward: let \(\langle l, a, r\rangle = \texttt{splay} \ x \ t\); if \(a = x\), return \(\langle l, a, r\rangle\); otherwise make \(x\) the root of a suitable recombination of \(l, a\) and \(r\).
splay x \langle AB, b, CD \rangle
= (\text{case} \ cmp \ x \ b \ of
  \langle \rangle \Rightarrow \langle AB, b, CD \rangle | 
  \langle A, a, B \rangle \Rightarrow 
    \text{case} \ cmp \ x \ a \ of 
      \langle \rangle \Rightarrow \langle A, a, \langle B, b, CD \rangle \rangle | 
      \langle A, a', A_2 \rangle \Rightarrow \langle A_1, a', \langle A_2, a, \langle B, b, CD \rangle \rangle \rangle | 
      \langle B_1, b', B_2 \rangle \Rightarrow \langle \langle A, a, \langle B, b, CD \rangle \rangle, b', \langle B_2, b, CD \rangle \rangle | 
    \text{else case splay x A of} \ 
    \langle A, a, \langle B, b, CD \rangle \rangle | 
    \langle B_1, b', B_2 \rangle \Rightarrow \langle \langle A, a, \langle B, b, CD \rangle \rangle, b', \langle B_2, b, CD \rangle \rangle | 
  \langle C, c, D \rangle \Rightarrow 
    \text{case} \ cmp \ x \ c \ of 
      \langle \rangle \Rightarrow \langle AB, b, CD \rangle | 
      \langle C_1, c', C_2 \rangle \Rightarrow \langle AB, b, C_1 \rangle, c', \langle C_2, c, D \rangle | 
      \langle AB, b, C \rangle, c, D \rangle | 
    \text{else case splay x C of} \ 
    \langle A_1, a', C_2 \rangle \Rightarrow \langle AB, b, C_1 \rangle, c', \langle C_2, c, D \rangle | 
    \langle AB, b, C \rangle, c, D \rangle | 
    \langle AB, b, \langle C, D \rangle \rangle | 
  \langle D_1, d, D_2 \rangle \Rightarrow \langle AB, b, C \rangle, c, \langle D_1, d, D_2 \rangle |)

Figure 20.1 Function splay

The implementation of delete x t in Figure 20.3 starts similarly: let \langle l, a, r \rangle = splay x t; if a \neq x, return \langle l, a, r \rangle. Otherwise follow the deletion-by-replacing paradigm (Section 5.2.1): if l \neq \langle \rangle, splay the maximal element m in l to the root and replace x with it. Note that splay_max returns a tree that is just a glorified pair: if t \neq \langle \rangle then splay_max t is of the form \langle t', m, \rangle. The definition splay_max \langle \rangle = \langle \rangle is not really needed (splay_max is always called with non-\langle \rangle argument) but some lemmas can be stated more slickly with this definition.
insert :: 'a ⇒ 'a tree ⇒ 'a tree
insert x t
= (if t = () then ()(, x, ()
else case splay x t of
  (l, a, r) ⇒ case cmp x a of
    LT ⇒ ⟨l, x, ⟨l, a, r⟩⟩ |
    EQ ⇒ ⟨l, a, r⟩ |
    GT ⇒ ⟨⟨l, a⟩, x, r⟩)

Figure 20.2  Function insert

delete :: 'a ⇒ 'a tree ⇒ 'a tree
delete x t
= (if t = () then ()
else case splay x t of
  (l, a, r) ⇒
    if x ≠ a then ⟨l, a, r⟩
    else if l = () then r
      else case splay_max l of ⟨l', m, _⟩ ⇒ ⟨l', m, r⟩)
splay_max :: 'a tree ⇒ 'a tree
splay_max () = ()
splay_max ⟨A, a, ⟨⟩⟩ = ⟨A, a, ⟨⟩⟩
splay_max ⟨A, a, ⟨B, b, CD⟩⟩ = (if CD = ⟨⟩ then ⟨⟨A, a, B⟩, b, ⟨⟩⟩
  else case splay_max CD of ⟨C, c, D⟩ ⇒ ⟨⟨A, a, B⟩, b, C⟩, c, D⟩)

Figure 20.3  Functions delete and splay_max
20.2 Correctness

The inorder approach of Section 5.4 applies. Because the details are a bit different (everything is reduced to splay) we present the top-level structure.

The following easy inductive properties are used implicitly in a number of subsequent proofs:

\[ \text{splay } a \ t = \langle \rangle \iff t = \langle \rangle \]
\[ \text{splay\_max } t = \langle \rangle \iff t = \langle \rangle \]

Correctness of isin

\[ \text{sorted } (\text{inorder } t) \to \text{isin } t \ x = (x \in \text{set } (\text{inorder } t)) \]

follows directly from this easy inductive property of splay:

\[ \text{splay } x \ t = \langle l, a, r \rangle \land \text{sorted } (\text{inorder } t) \to (x \in \text{set } (\text{inorder } t)) = (x = a) \]

Correctness of insert and delete

\[ \text{sorted } (\text{inorder } t) \to \text{inorder } (\text{insert } x \ t) = \text{ins\_list } x \ (\text{inorder } t) \]
\[ \text{sorted } (\text{inorder } t) \to \text{inorder } (\text{delete } x \ t) = \text{del\_list } x \ (\text{inorder } t) \]

relies on the following characteristic inductive properties of splay:

\[ \text{inorder } (\text{splay } x \ t) = \text{inorder } t \tag{20.1} \]
\[ \text{sorted } (\text{inorder } t) \land \text{splay } x \ t = \langle l, a, r \rangle \to \text{sorted } (\text{inorder } l \ @ \ [a] = \text{inorder } t \land r = \langle \rangle) \]

Correctness of delete also needs the inductive proposition

\[ \text{splay\_max } t = \langle l, a, r \rangle \land \text{sorted } (\text{inorder } t) \to \text{inorder } l \ @ [a] = \text{inorder } t \land r = \langle \rangle \]

Note that \[ \text{inorder } (\text{splay } x \ t) = \text{inorder } t \] is also necessary to justify the proper isin that returns the newly created tree as well.

Automation of the above proofs requires the lemmas in Figure 5.2 together with a few additional lemmas about sorted, ins\_list and del\_list that can be found in the Isabelle proofs.

Recall from Section 5.4 that correctness of insert and delete implies that they preserve \( \text{bst} = \text{sorted} \circ \text{inorder} \). Similarly, (20.1) implies that splay preserves bst. Thus we may assume the invariant bst in the amortized analysis.

These two easy size lemmas are used implicitly below:

\[ |\text{splay } a \ t| = |t| \quad |\text{splay\_max } t| = |t| \]
20.3 Amortized Analysis

This section shows that splay, insertion and deletion all have amortized logarithmic complexity.

We define the potential $\Phi$ of a tree as the sum of the potentials $\varphi$ of all nodes:

$$
\Phi \colon 'a\ tree \Rightarrow real
$$

$$
\Phi \langle \rangle = 0
$$

$$
\Phi \langle l, a, r \rangle = \varphi \langle l, a, r \rangle + \Phi l + \Phi r
$$

$$
\varphi t \equiv \log_2 |t|
$$

The central result is the amortized complexity of splay. Function $T_{splay}$ is shown in Appendix B.9. We follow (18.1) and define

$$
A_{splay} a t = T_{splay} a t + \Phi (splay a t) - \Phi t
$$

First we consider the case where the element is in the tree:

**Theorem 20.1.** $bst t \land \langle l, x, r \rangle \in subtrees t \rightarrow

$$
A_{splay} x t \leq 3 \cdot (\varphi t - \varphi \langle l, x, r \rangle) + 1
$$

**Proof** by induction on the computation of splay. The base cases involving $\langle \rangle$ are impossible. For example, consider the call splay $x t$ where $t = \langle \rangle, b, C \rangle$ and $x < b$: from $\langle l, x, r \rangle \in subtrees t$ it follows that $x \in set\_tree t$ but because $bst t$ and $x < b$ this implies that $x \in set\_tree \langle \rangle$, a contradiction. There are three feasible base cases. The case $t = \langle \_, x, \_ \rangle$ is easy. We consider one of the two other symmetric cases. Let $t = \langle \langle A, x, B \rangle, b, C \rangle$ and $t' = splay x t = \langle A, x, \langle B, b, C \rangle \rangle$.

$$
A_{splay} x t = \Phi t' - \Phi t + 1
$$

by definition of $A_{splay}$ and $T_{splay}$

$$
= \varphi t' + \varphi \langle B, b, C \rangle - \varphi t - \varphi \langle A, x, B \rangle + 1
$$

by definition of $\Phi$

$$
= \varphi \langle B, b, C \rangle - \varphi \langle A, x, B \rangle + 1
$$

by definition of $\varphi$

$$
\leq \varphi t - \varphi \langle A, x, B \rangle + 1
$$

because $\varphi \langle B, b, C \rangle \leq \varphi t$

$$
\leq 3 \cdot (\varphi t - \varphi \langle A, x, B \rangle) + 1
$$

because $\varphi \langle A, x, B \rangle \leq \varphi t$

$$
= 3 \cdot (\varphi t - \varphi \langle l, x, r \rangle) + 1
$$

because $bst t \land \langle l, x, r \rangle \in subtrees t$
There are four inductive cases. We consider two of them, the other two are symmetric variants. First the so-called zig-zig case:

This is the case where \( x < a < b \) and \( A \neq \langle \rangle \). On the left we have the input and on the right the output of splay \( x \). Because \( A \neq \langle \rangle \), splay \( x A = \langle A_1, a', A_2 \rangle =: A' \) for some \( A_1, a' \) and \( A_2 \). The intermediate tree is obtained by replacing \( A \) by \( A' \). This tree is shown for illustration purpose only; in the algorithm the right tree is constructed directly from the left one. Let \( X = \langle l, x, r \rangle \). Clearly \( X \in \text{subtrees} A \). We abbreviate compound trees like \( \langle A, a, B \rangle \) by the names of their subtrees, in this case \( AB \). First note that

\[
\varphi A_1 A_2 BC = \varphi ABC
\]

because \( |A'| = |\text{splay } x A| = |A| \). We can now prove the claim:

\[
A_{\text{splay}} x ABC = T_{\text{splay}} x A + 1 + \varphi A_1 A_2 BC - \varphi ABC
\]

\[
= T_{\text{splay}} x A + 1 + \varphi A_1 + \varphi A_2 + \varphi A_2 BC + \varphi BC - \varphi A - \varphi AB
\]

by (\*) and definition of \( \varphi \)

\[
= T_{\text{splay}} x A + \varphi A' - \varphi A' - \varphi A + \varphi A_2 BC + \varphi BC - \varphi AB - \varphi A' + 1
\]

\[
= A_{\text{splay}} x A + \varphi A_2 BC + \varphi BC - \varphi AB - \varphi A' + 1
\]

\[
\leq 3 \cdot \varphi A + \varphi A_2 BC + \varphi BC - \varphi AB - \varphi A' - 3 \cdot \varphi X + 2
\]

by IH and \( X \in \text{subtrees} A \)

\[
= 2 \cdot \varphi A + \varphi A_2 BC + \varphi BC - \varphi AB - 3 \cdot \varphi X + 2
\]

because \( \varphi A = \varphi A' \)

\[
< \varphi A + \varphi A_2 BC + \varphi BC - 3 \cdot \varphi X + 2
\]

because \( \varphi A < \varphi AB \)

\[
< \varphi A_2 BC + 2 \cdot \varphi ABC - 3 \cdot \varphi X + 1
\]

because \( 1 + \lg x + \lg y < 2 \cdot \lg (x + y) \) if \( x, y > 0 \)

\[
< 3 \cdot (\varphi ABC - \varphi X) + 1
\]

because \( \varphi A_2 BC < \varphi ABC \)
Now we consider the so-called zig-zag case:

\[
\begin{array}{ccc}
A & B & C \\
\downarrow & \downarrow & \downarrow \\
A & B_1 & B_2 \\
\end{array}
\rightarrow
\begin{array}{ccc}
a & b & C \\
\downarrow & \downarrow & \downarrow \\
A & B & C \\
\end{array}
\rightarrow
\begin{array}{ccc}
b & a & C \\
\downarrow & \downarrow & \downarrow \\
A & B_1 & B_2 \\
\end{array}
\]

This is the case where \( a < x < b \) and \( B \neq \emptyset \). On the left we have the input and on the right the output of \( \text{splay} \ x \). Because \( B \neq \emptyset \), \( \text{splay} \ x \ B = \langle B_1, b', B_2 \rangle =: B' \) for some \( B_1, b' \) and \( B_2 \). The intermediate tree is obtained by replacing \( B \) by \( B' \). Let \( X = \langle l, x, r \rangle \). Clearly \( X \in \text{subtrees} \ B \). The proof is very similar to the zig-zig case, the same naming conventions apply and we omit some details:

\[
A_{\text{splay}} \ x \ ABC = T_{\text{splay}} \ x \ A + 1 + \Phi \ AB_1B_2C - \Phi \ ABC
\]

\[
= A_{\text{splay}} \ x \ B + \varphi \ AB_1 + \varphi \ B_2C - \varphi \ AB - \varphi \ B' + 1
\]

using \( \varphi \ AB_1B_2C = \varphi \ ABC \)

\[
\leq 3 \cdot \varphi \ B + \varphi \ AB_1 + \varphi \ B_2C - \varphi \ AB - \varphi \ B' - 3 \cdot \varphi \ X + 2
\]

by IH and \( X \in \text{subtrees} \ B \)

\[
= 2 \cdot \varphi \ B + \varphi \ AB_1 + \varphi \ B_2C - \varphi \ AB - 3 \cdot \varphi \ X + 2
\]

because \( \varphi \ B = \varphi \ B' \)

\[
< \varphi \ B + \varphi \ AB_1 + \varphi \ B_2C - 3 \cdot \varphi \ X + 2
\]

because \( \varphi \ B < \varphi \ AB \)

\[
< \varphi \ B + 2 \cdot \varphi \ ABC - 3 \cdot \varphi \ X + 1
\]

because \( 1 + \lg \ x + \lg \ y < 2 \cdot \lg \ (x + y) \) if \( x, y > 0 \)

\[
< 3 \cdot (\varphi \ ABC - \varphi \ X) + 1
\]

because \( \varphi \ B < \varphi \ ABC \)

Because \( \varphi \langle l, x, r \rangle \geq 1 \), the above theorem implies

**Corollary 20.2.** \( \text{bst} \ t \land x \in \text{set_tree} \ t \longrightarrow A_{\text{splay}} \ x \ t \leq 3 \cdot (\varphi \ t - 1) + 1 \)

If \( x \) is not in the tree we show that there is a \( y \) in the tree such that splaying with \( y \) would produce the same tree in the same time:

**Lemma 20.3.** \( t \neq \langle \rangle \land \text{bst} \ t \longrightarrow \\
(\exists y \in \text{set_tree} \ t. \ \text{splay} \ y \ t = \text{splay} \ x \ t \land T_{\text{splay}} \ y \ t = T_{\text{splay}} \ x \ t) \)

Element \( y \) is the last element in the tree that the search for \( x \) encounters before it hits a leaf. Naturally, the proof is by induction on the computation of \( \text{splay} \).

Combining this lemma with Corollary 20.2 yields the final unconditional amortized complexity of \( \text{splay} \) on BSTs:

**Corollary 20.4.** \( \text{bst} \ t \longrightarrow A_{\text{splay}} \ x \ t \leq 3 \cdot \varphi \ t + 1 \)
The “− 1” has disappeared to accommodate the case $t = \emptyset$.

The amortized analysis of insertion is straightforward now. From the amortized complexity of $splay$ it follows that

**Lemma 20.5.** $bst \ t \rightarrow T_{\text{insert}} \ x \ t + \Phi \ (\text{insert} \ x \ t) - \Phi \ t \leq 4 \cdot \varphi \ t + 2$

We omit the proof which is largely an exercise in simple algebraic manipulations.

The amortized analysis of deletion is similar but a bit more complicated because of the additional function $splay_{\text{max}}$ whose amortized running time is defined as usual:

$$A_{splay_{\text{max}}} \ t = T_{splay_{\text{max}}} \ t + \Phi \ (splay_{\text{max}} \ t) - \Phi \ t$$

Like in the analysis of $A_{splay}$, an inductive proof yields

$$t \neq \emptyset \rightarrow A_{splay_{\text{max}}} \ t \leq 3 \cdot (\varphi \ t - 1) + 1$$

from which

$$A_{splay_{\text{max}}} \ t \leq 3 \cdot \varphi \ t + 1$$

follows by a simple case analysis. The latter proposition, together with Corollary 20.4, proves the amortized logarithmic complexity of $delete$

$$bst \ t \rightarrow T_{\text{delete}} \ a \ t + \Phi \ (\text{delete} \ a \ t) - \Phi \ t \leq 6 \cdot \varphi \ t + 2$$

in much the same way as for $insert$ (Lemma 20.5).

A running time analysis of $isin$ is trivial because $isin$ is just $splay$ followed by a constant-time test.

### 20.4 Exercises

**Exercise 20.1.** Find a sequence of numbers $n_1, n_2, \ldots, n_k$ such that the insertion of these numbers one by one creates a splay tree of height $k$.

### 20.5 Chapter Notes

Splay trees were invented and analyzed by Sleator and Tarjan [1985] for which they received the 1999 ACM Paris Kanellakis Theory and Practice Award [Kanellakis]. In addition to the amortized complexity as shown above they proved that splay trees perform as well as static BSTs (the Static Optimality Theorem) and conjectured that, roughly speaking, they even perform as well as any other BST-based algorithm. This Dynamic Optimality Conjecture is still open.

This chapter is based on earlier publications [Nipkow 2015, 2016, Nipkow and Brinkop 2019, Schoenmakers 1993].
Skew Heaps

Tobias Nipkow

Skew heaps are heaps in the sense of Section 13.1 and implement mergeable priority queues. Skew heaps can be viewed as a self-adjusting form of leftist heaps that attempts to maintain balance by unconditionally swapping all nodes in the merge path when merging two heaps.

### 21.1 Implementation of ADT `PriorityQueueMerge`

The central operation is `merge`:

```haskell
merge :: 'a tree ⇒ 'a tree ⇒ 'a tree
merge ∅ t = t
merge t ∅ = t
merge (∧t₁, a₁, r₁) = t₁ ∨ (∧t₂, a₂, r₂) = t₂
= (if a₁ ≤ a₂ then merge t₂ r₁, a₁, l₁) else (merge t₁ r₂, a₂, l₂)
```

The remaining operations (`{}, insert, get_min` and `del_min`) are defined as in Section 13.1.

The following properties of `merge` have easy inductive proofs:

- `|merge t₁ t₂| = |t₁| + |t₂|
- `mset_tree (merge t₁ t₂) = mset_tree t₁ + mset_tree t₂`
- `heap t₁ ∧ heap t₂ → heap (merge t₁ t₂)`

Now it is straightforward to prove the correctness of the implementation w.r.t. the ADT `PriorityQueueMerge`.

Skew heaps attempt to maintain balance, but this does not always work:

**Exercise 21.1.** Find a sequence of numbers `n₁, n₂, ... n_k` such that the insertion of theses numbers one by one creates a tree of height `k`. Prove that this sequence will produce a tree of height `k`.

Nevertheless, insertion and deletion have amortized logarithmic complexity.
21.2 Amortized Analysis

The key is the definition of the potential. It counts the number of right-heavy (\(rh\)) nodes:

\[
\Phi :: 'a tree \Rightarrow int
\]
\[
\Phi \langle \rangle = 0
\]
\[
\Phi \langle l, \_, r \rangle = \Phi l + \Phi r + rh l r
\]

\(rh :: 'a tree \Rightarrow 'a tree \Rightarrow nat\)
\[
rh l r = (if \ |l| < |r| then 1 else 0)
\]

The rough intuition: because \(merge\) descends along the right spine, the more right-heavy nodes a tree contains, the longer \(merge\) takes.

Two auxiliary functions count the number of right-heavy nodes on the left spine (\(lrh\)) and left-heavy (= not right-heavy) nodes on the right spine (\(rlh\)):

\[
lrh :: 'a tree \Rightarrow nat
\]
\[
lrh \langle \rangle = 0
\]
\[
lrh \langle l, \_, r \rangle = rh l r + lrh l
\]

\[
rlh :: 'a tree \Rightarrow nat
\]
\[
rlh \langle \rangle = 0
\]
\[
rlh \langle l, \_, r \rangle = 1 - rh l r + rlh r
\]

The following properties have automatic inductive proofs:

\[
2^{lrh t} \leq |t| + 1 \quad 2^{rlh t} \leq |t| + 1
\]

They imply

\[
lrh t \leq lg |t|_1 \quad rlh t \leq lg |t|_1 \quad (21.1)
\]

Now we are ready for the amortized analysis. All time functions can be found in Appendix B.10. The key lemma is an upper bound of the amortized complexity of \(merge\) in terms of \(lrh\) and \(rlh\):

**Lemma 21.1.** \(T_{merge} t_1 t_2 + \Phi (merge t_1 t_2) - \Phi t_1 - \Phi t_2 \leq lrh (merge t_1 t_2) + rlh t_1 + rlh t_2 + 1\)
Proof by induction on the computation of `merge`. We consider only the node-node case: let \( t_1 = (l_1, a_1, r_1) \) and \( t_2 = (l_2, a_2, r_2) \). W.l.o.g. assume \( a_1 \leq a_2 \). Let \( m = \text{merge} \ t_2 \ r_1 \).

\[
T_{\text{merge}} t_1 \ t_2 = \Phi (\text{merge} \ t_1 \ t_2) - \Phi t_1 - \Phi t_2
\]
\[
= T_{\text{merge}} t_2 \ r_1 + 1 + \Phi m + \Phi l_1 + rh m l_1 - \Phi t_1 - \Phi t_2
\]
\[
= T_{\text{merge}} t_2 \ r_1 + 1 + \Phi m + rh m l_1 - \Phi r_1 - rh l_1 r_1 - \Phi t_2
\]
\[
\leq lrh m + rh t_2 + rh t_1 + rh m l_1 + 2 - rh l_1 r_1 \quad \text{by IH}
\]
\[
= lrh m + rh l_1 t_2 + rh t_1 + rh m l_1 + 1
\]
\[
= lrh (\text{merge} \ t_1 \ t_2) + rh l_1 t_2 + rh t_1 + rh l_1 t_2 + 1
\]

As a consequence we can prove the following logarithmic upper bound on the amortized complexity of `merge`:

\[
T_{\text{merge}} t_1 \ t_2 + \Phi (\text{merge} \ t_1 \ t_2) - \Phi t_1 - \Phi t_2
\]
\[
\leq lrh (\text{merge} \ t_1 \ t_2) + rh l_1 t_2 + rh t_1 + rh l_1 t_2 + 1 \quad \text{by Lemma 21.1}
\]
\[
\leq \lg |\text{merge} t_1 \ t_2| + \lg |t_1| + \lg |t_2| + 1 \quad \text{by (21.1)}
\]
\[
\leq \lg (|t_1| + |t_2|) + 2 \cdot \lg (|t_1| + |t_2|) + 1
\]
\[
= 3 \cdot \lg (|t_1| + |t_2|) + 1
\]

The amortized complexity of insertion and deletion follows easily from the complexity of `merge`:

\[
T_{\text{insert}} a \ t + \Phi (\text{insert} \ a \ t) - \Phi \ t \leq 3 \cdot \lg (|t_1| + 2) + 1
\]
\[
T_{\text{del_min}} t + \Phi (\text{del_min} \ t) - \Phi \ t \leq 3 \cdot \lg (|t_1| + 2) + 1
\]

21.3 Chapter Notes

Skew heaps were invented by Sleator and Tarjan [1986] as one of the first self-organizing data structures. Their presentation was imperative. Our presentation follows earlier work by Nipkow [2015] and Nipkow and Brinkop [2019] based on the functional account by Kaldewaij and Schoenmakers [1991].
Pairing Heaps

Tobias Nipkow

The pairing heap is another form of a self-adjusting priority queue. Section 22.1 presents an intuitive version of pairing heaps based on lists. In the rest of the chapter we change to a slightly different presentation that leads to a more succinct amortized analysis.

22.1 Implementation via Lists

A pairing heap is a heap in the sense that it is a tree with the minimal element at the root — except that it is not a binary tree but a tree where each node has a list of children:

\[
\text{datatype } \text{'a heap} = \text{Empty} | \text{'a heap list}
\]

The abstraction function to multisets and the invariant follow the heap paradigm:

\[
\begin{align*}
\text{mset_heap} : \text{'a heap} & \rightarrow \text{'a multiset} \\
\text{mset_heap Empty} & = \{ \} \\
\text{mset_heap (Hp x hs)} & = \{x\} + \sum_s \text{(mset (map mset_heap hs))}
\end{align*}
\]

\[
\begin{align*}
\text{pheap} : \text{'a heap} & \rightarrow \text{bool} \\
\text{pheap Empty} & = \text{True} \\
\text{pheap (Hp x hs)} & = (\forall h \in \text{set hs}. (\forall y \mid mset\_heap h. x \leq y) \land \text{pheap h})
\end{align*}
\]

Note that \text{pheap} is sufficient for functional correctness. Additionally, \text{Empty} does not occur inside a non-empty heap. The amortized analysis, where this additional invariant would be required, is performed on a slightly different model without \text{Empty}.

The implementations of \text{empty} and \text{get_min} are obvious, and \text{insert} follows the standard heap paradigm:
Chapter 22  Pairing Heaps

\[ empty = \text{Empty} \]

\[ \text{get\_min} :: 'a \text{ heap} \Rightarrow 'a \]

\[ \text{get\_min} (\text{Hp } x \_\_) = x \]

\[ \text{insert} :: 'a \Rightarrow 'a \text{ heap} \Rightarrow 'a \text{ heap} \]

\[ \text{insert } x \ h = \text{merge} (\text{Hp } x \ [\_]) \ h \]

Function \text{merge} is not recursive (as in binary heaps) but simply adds one of the two heaps to the front of the top-level heaps of the other, depending on the root value:

\[ \text{merge} :: 'a \text{ heap} \Rightarrow 'a \text{ heap} \Rightarrow 'a \text{ heap} \]

\[ \text{merge } h \ \text{Empty} = h \]

\[ \text{merge } \text{Empty} \ h = h \]

\[ \text{merge} (\text{Hp } x \ hsx =: hx) (\text{Hp } y \ hsy =: hy) \]

\[ = (\text{if } x < y \text{ then } \text{Hp } x (\text{hy } \# \hsx) \text{ else } \text{Hp } y (hx \# \hsy)) \]

Thus \text{merge} and \text{insert} have constant running time. All the work is offloaded on \text{del\_min} which just calls \text{merge\_pairs}:

\[ \text{del\_min} :: 'a \text{ heap} \Rightarrow 'a \text{ heap} \]

\[ \text{del\_min } \text{Empty} = \text{Empty} \]

\[ \text{del\_min} (\text{Hp } \_ \hs) = \text{merge\_pairs} \ hs \]

\[ \text{merge\_pairs} :: 'a \text{ heap list} \Rightarrow 'a \text{ heap} \]

\[ \text{merge\_pairs} [\_] = \text{Empty} \]

\[ \text{merge\_pairs} [h] = h \]

\[ \text{merge\_pairs} (h_1 \# h_2 \# \hs) = \text{merge} (\text{merge} h_1 \ h_2) (\text{merge\_pairs} \ hs) \]

Function \text{merge\_pairs} is a compact way of expressing a two pass algorithm: on the first pass from left to right, it merges pairs of adjacent heaps (hence “pairing heap”) and on the second pass it merges the results in a cascade from right to left. By reformulating the definition in terms of these two passes, we obtain a more readable formulation with the same running time:
del_min :: 'a heap ⇒ 'a heap

\[ \text{del_min Empty} = \text{Empty} \]

\[ \text{del_min (Hp x hs)} = \text{pass}_2 (\text{pass}_1 \text{hs}) \]

\[ \text{pass}_1 :: \text{a heap list ⇒ a heap list} \]

\[ \text{pass}_1 (h_1 \# h_2 \# \text{hs}) = \text{merge} h_1 h_2 \# \text{pass}_1 \text{hs} \]

\[ \text{pass}_1 \text{hs} = \text{hs} \]

\[ \text{pass}_2 :: \text{a heap list ⇒ a heap} \]

\[ \text{pass}_2 [] = \text{Empty} \]

\[ \text{pass}_2 (h \# \text{hs}) = \text{merge} h (\text{pass}_2 \text{hs}) \]

The proof of \( \text{pass}_2 (\text{pass}_1 \text{hs}) = \text{merge\_pairs} \text{hs} \) is an easy induction.

Clearly \text{del\_min} can take linear time but it will turn out that the constant-time \text{insert} saves enough to guarantee amortized logarithmic complexity for both insertion and deletion.

We base the correctness proofs on the \text{merge\_pairs} version of \text{del\_min}. From the following lemmas (all proofs are routine inductions) the properties in the specifications \text{Priority\_Queue(\_Merge)} follow easily.

\[ h \neq \text{Empty} \rightarrow \text{get\_min} h \in h \text{mset\_heap} h \]

\[ h \neq \text{Empty} \land \text{pheap} h \land x \in h \text{mset\_heap} h \rightarrow \text{get\_min} h \leq x \]

\[ \text{mset\_heap (merge} h_1 h_2) = \text{mset\_heap} h_1 + \text{mset\_heap} h_2 \]

\[ \text{mset\_heap (merge\_pairs} \text{hs}) \]

\[ = \sum_{x} (\text{image\_mset mset\_heap} (\text{mset} \text{hs})) \]

\[ h \neq \text{Empty} \rightarrow \]

\[ \text{mset\_heap (del\_min} h) = \text{mset\_heap} h - \{\text{get\_min} h\} \]

\[ \text{pheap} h_1 \land \text{pheap} h_2 \rightarrow \text{pheap} (\text{merge} h_1 h_2 ) \]

\[ (\forall h \in \text{set} \text{hs. pheap} h) \rightarrow \text{pheap} (\text{merge\_pairs} \text{hs}) \]

\[ \text{pheap} h \rightarrow \text{pheap} (\text{del\_min} h) \]

### 22.2 Amortized Analysis

The amortized analysis of pairing heaps is slightly simplified if we replace the above type of heaps by trees as follows: a heap \( \text{Hp} x \text{hs} \) is expressed as the tree \( \langle \text{hs}, x, \langle \rangle \rangle \) and a list of heaps \([\text{Hp} x_1 \text{hs}_1, \text{Hp} x_2 \text{hs}_2, \ldots] \) is expressed as the tree \( \langle \text{hs}_1, x_1, \langle \text{hs}_2, \ldots \rangle \rangle \).
empty = ()

get_min :: 'a tree ⇒ 'a
get_min (_, x, _) = x

link :: 'a tree ⇒ 'a tree
link (hsx, x, ⟨hsy, y, hs⟩) = (if x < y then ⟨hsy, y, hs⟩, x, hs) else ⟨hsx, x, hsy⟩, y, hs⟩
link hp = hp

pass₁ :: 'a tree ⇒ 'a tree
pass₁ (hsx, x, ⟨hsy, y, hs⟩) = link (hsx, x, ⟨hsy, y, pass₁ hs⟩)
pass₁ hp = hp

pass₂ :: 'a tree ⇒ 'a tree
pass₂ (hsx, x, hs) = link (hsx, x, pass₂ hs)
pass₂ () = ()

get_min :: 'a tree ⇒ 'a
get_min (_, x, _) = x

merge :: 'a tree ⇒ 'a tree ⇒ 'a tree
merge () hp = hp
merge hp () = hp
merge (hsx, x, ⟨⟩) ⟨hsy, y, ⟨⟩⟩ = link (hsx, x, ⟨hsy, y, ⟨⟩⟩)

insert :: 'a ⇒ 'a tree ⇒ 'a tree
insert x hp = merge ⟨⟩, x, ⟨⟩ hp

Figure 22.1  Pairing heaps via trees

This simplifies the analysis because we now have to deal only with a single type, trees.
The code for the tree representation of pairing heaps is shown in Figure 22.1. We work with the pass_1/pass_2 version of del_min. The correctness proof is very similar to what we saw in the previous section. We merely display the two invariants:

```
is_root :: 'a tree ⇒ bool
is_root hp = (case hp of () ⇒ True | (_, _, r) ⇒ r = ())
```

```
pheap :: 'a tree ⇒ bool
pheap () = True
pheap (l, x, r) = ((∀y∈set_tree l. x ≤ y) ∧ pheap l ∧ pheap r)
```

Now we turn to the amortized analysis. The potential of a tree is the sum of the logarithms of the sizes of the subtrees:

```
Φ :: 'a tree ⇒ real
Φ () = 0
Φ (l, x, r) = lg |⟨l, x, r⟩| + Φ l + Φ r
```

These easy inductive size properties are frequently used implicitly below:

- |link hp| = |hp|
- |pass_1 hp| = |hp|
- |pass_2 hp| = |hp|
- is_root h_1 ∧ is_root h_2 ⇒ |merge h_1 h_2| = |h_1| + |h_2|

### 22.2.1 Potential Differences

We can now analyze the differences in potential caused by all the queue operations. In a separate step we will derive their amortized complexities.

For insertion, the following upper bound follows trivially from the definitions:

**Lemma 22.1.** is_root hp ⇒ Φ (insert x hp) − Φ hp ≤ lg (|hp| + 1)

For merge it needs a bit more work:

**Lemma 22.2.** h_1 = ⟨hs_1, x_1, ⟨⟩⟩ ∧ h_2 = ⟨hs_2, x_2, ⟨⟩⟩ ⇒
Φ (merge h_1 h_2) − Φ h_1 − Φ h_2 ≤ lg (|h_1| + |h_2|) + 1

**Proof.** From

Φ (merge h_1 h_2)
= Φ (link ⟨hs_1, x_1, h_2⟩)
Chapter 22  
Pairing Heaps

\[ \Phi \cdot h_1 + \Phi \cdot h_2 + \lg (|h_1| + |h_2| + 1) + \lg (|h_1| + |h_2| + 2) \]
\[ = \Phi \cdot h_1 + \Phi \cdot h_2 + \lg (|h_1| + |h_2| + 1) + \lg (|h_1| + |h_2|) \]

it follows that

\[ \Phi (\text{merge } h_1, h_2) - \Phi h_1 - \Phi h_2 \]
\[ = \lg (|h_1| + |h_2| + 1) - \lg (|h_1| + |h_2|) \]
\[ - \lg (|h_1| + 1) - \lg (|h_2| + 1) \]
\[ \leq \lg (|h_1| + |h_2|) + 1 \]

because \( \lg (1 + x + y) \leq 1 + \lg (1 + x) + \lg (1 + y) \) if \( x, y \geq 0 \)

Now we come to the core of the proof, the analysis of \( \text{del_min} \). Its running time is linear in the number of nodes reachable by descending to the right (starting from the left child of the root). We denote this metric by \( \text{len} \):

\[
\text{len} :: \text{'a tree } \Rightarrow \text{ nat}
\]
\[
\text{len } \langle \rangle = 0
\]
\[
\text{len } \langle \_, \_, r \rangle = 1 + \text{len } r
\]

Therefore we have to show that the potential change compensates for this linear work. Our main goal is this:

**Theorem 22.3.** \( \Phi (\text{del_min } \langle hs, x, \langle \rangle \rangle) - \Phi \langle hs, x, \langle \rangle \rangle \)
\[ \leq 2 \cdot \lg (|hs| + 1) - \text{len } hs + 2 \]

It will be proved in two steps: First we show that \( \text{pass}_1 \) frees enough potential to compensate for the work linear in \( \text{len } hs \) and increases the potential only by a logarithmic term. Then we show that the increase due to \( \text{pass}_2 \) is also only at most logarithmic. Combining these results one easily shows that the amortized running time of \( \text{del_min} \) is indeed logarithmic.

First we analyze the potential difference caused by \( \text{pass}_1 \):

**Lemma 22.4.** \( \Phi (\text{pass}_1 hs) - \Phi hs \leq 2 \cdot \lg (|hs| + 1) - \text{len } hs + 2 \)

*Proof* by induction on the computation of \( \text{pass}_1 \). The base cases are trivial. We focus on the induction step. Let \( t = \langle hs_1, x, \langle hs_2, y, hs \rangle \rangle, n_1 = |hs_1|, n_2 = |hs_2| \) and \( m = |hs| \).

\[ \Phi (\text{pass}_1 t) - \Phi t \]
\[ = \lg (n_1 + n_2 + 1) - \lg (n_2 + m + 1) + \Phi (\text{pass}_1 hs) - \Phi hs \]
\[ \leq \lg (n_1 + n_2 + 1) - \lg (n_2 + m + 1) + 2 \cdot \lg (m + 1) - \text{len } hs + 2 \text{ by IH} \]
\[ \leq 2 \cdot \lg (n_1 + n_2 + m + 1) - \lg (n_2 + m + 1) + \lg (m + 1) - \text{len } hs \]

because \( \lg x + \lg y + 2 \leq 2 \cdot \lg (x + y) \) if \( x, y > 0 \)
\[ \leq 2 \cdot \lg (n_1 + n_2 + m + 2) - \text{len } hs \]
22.2 Amortized Analysis

\[ t = 2 \cdot \lg |t| - \text{len } t + 2 \]
\[ \leq 2 \cdot \lg (|t| + 1) - \text{len } t + 2 \]

Now we turn to pass 2:

**Lemma 22.5.** \( h s \neq \langle \rangle \rightarrow \Phi (\text{pass}_2 \ h s) - \Phi \ h s \leq \lg |h s| \)

**Proof** by induction on \( h s \). The base cases are trivial. The induction step (for \( \langle h s_1, x, h s \rangle \)) is trivial if \( h s = \langle \rangle \). Assume \( h s = \langle h s_2, y, r \rangle \). Now we need one more property of \( \text{pass}_2 \):

\[ \exists h s_3 \ z. \ \text{pass}_2 \langle h s_2, y, r \rangle = \langle h s_3, z, \langle \rangle \rangle \]

The proof is a straightforward induction on \( r \). This implies \(|h s_3| + 1 = |h s|\) and thus

\[ \Phi (\text{link} \langle h s_1, x, \text{pass}_2 \ h s \rangle) - \Phi \ h s_1 - \Phi (\text{pass}_2 \ h s) \]
\[ = \lg (|h s_1| + |h s| + 1) + \lg (|h s_1| + |h s|) - \lg |h s| \]
\[ \leq \lg (|h s_1| + |h s|) \]

Thus the overall claim follows:

\[ \Phi (\text{pass}_2 \langle h s_1, x, h s \rangle) - \Phi \langle h s_1, x, h s \rangle \]
\[ = \Phi (\text{link} \langle h s_1, x, \text{pass}_2 \ h s \rangle) - \Phi h s_1 - \Phi \ h s - \lg (|h s_1| + |h s| + 1) \]
\[ = \Phi (\text{pass}_2 \ h s) - \Phi \ h s + \lg (|h s_1| + |h s|) - \lg |h s| \]
\[ \leq \lg (|h s_1| + |h s|) \]

**Corollary 22.6.** \( \Phi (\text{pass}_2 \ h s) - \Phi \ h s \leq \lg (|h s| + 1) \)

Finally we can prove Theorem 22.3:

\[ \Phi (\text{del}_\text{min} \langle h s, x, \langle \rangle \rangle) - \Phi \langle h s, x, \langle \rangle \rangle \]
\[ = \Phi (\text{pass}_2 (\text{pass}_1 \ h s)) - \lg (|h s| + 1) - \Phi \ h s \]
\[ \leq \Phi (\text{pass}_1 \ h s) - \Phi \ h s \]
\[ \leq 2 \cdot \lg (|h s| + 1) - \text{len } h s + 2 \]

**22.2 Amortized Running Times**

The running time functions are displayed in Appendix B.11. It is now straightforward to derive these amortized running times:

\[ \text{is\_root} \ h \rightarrow T_{\text{insert}} \ a \ h + \Phi (\text{insert} \ a \ h) - \Phi \ h \leq \lg (|h| + 1) + 1 \]
\[ \text{is\_root} \ h_1 \land \text{is\_root} \ h_2 \rightarrow \]
\[ T_{\text{merge}} \ h_1 \ h_2 + \Phi (\text{merge} \ h_1 \ h_2) - \Phi \ h_1 - \Phi \ h_2 \leq \lg (|h_1| + |h_2| + 1) + 2 \]

They follow from the corresponding Lemmas 22.1 and 22.2.

Combining this inductive upper bound for the running time of the two passes

\[ T_{\text{pass}_2} (\text{pass}_1 \ h s_1) + T_{\text{pass}_1} \ h s_1 \leq \text{len } h s_1 + 2 \]

with Theorem 22.3 yields the third and final amortized running time:
Thus we have proved that insertion, merging and deletion all have amortized logarithmic running times.

22.3 Chapter Notes
Pairing heaps were invented by Fredman et al. [1986] as a simpler but competitive alternative to Fibonacci heaps. The authors gave the amortized analysis presented above and conjectured that it can be improved. Later research confirmed this [Iacono 2000, Iacono and Yagnatinsky 2016, Pettie 2005] but the final analysis is still open. An empirical study [Larkin et al. 2014] showed that pairing heaps do indeed outperform Fibonacci heaps in practice. This chapter is based on an article by Nipkow and Brinkop [2019].
Part V

Selected Topics
Nothing could be simpler than searching for occurrences of a string in a text file, yet we have two sophisticated algorithms for doing this: one by Knuth, Morris and Pratt (KMP), the other by Boyer and Moore. Both were published in 1977, when 1 MB was thought to be a lot of memory. Nowadays strings can be orders of magnitude longer, making the need for efficiency all the greater. Bioinformatics requires searching truly gigantic strings: of nucleotides (when working with genomes) and amino acids (in the case of proteins). Here we look at KMP, the simpler of the two.

The naive algorithm aligns the pattern $p$ with the text string $a$, comparing corresponding characters from left to right, and in case of a mismatch, shifting one position along $a$ and starting again. This is actually fine under plausible assumptions. The alphabet surely has more than one character, and if furthermore the characters in the string are random then the expected length of a partial match will be finite, since it involves the sum of a geometric series. Ergo, linear time.

But if the text is not random then the worst-case time is $O(mn)$, where $m = ||p||$ and $n = ||a||$. For suppose that $p$ and $a$ both have the form $\ldots xx \ldots xy$, consisting entirely of the letter $x$ except having a single $y$ at the end. The naive algorithm will make $m$ comparisons, failing at the last one; then it will shift $p$ one position along $a$ even though there is no hope of a match. This wasteful search will continue until $a$ is exhausted.

The idea of KMP is to exploit the knowledge gained from the partial match, never re-comparing characters that matched. At the first mismatched character, it shifts $p$ as far to the right as safely possible. To do so, it consults a precomputed table, based on the pattern $p$, identifying repeated substrings for which the current, failed partial match could become the first part of a full match.

In the case of our example, the successful match of the first part of the pattern, namely $\ldots xx \ldots$, means we already know the previous $m - 1$ characters of $a$, so instead of shifting one position along and checking $p$ from the beginning, we can check from where we left off, i.e. its penultimate character. The search will still fail until the final
Chapter 23

Knuth–Morris–Pratt String Search

If y is reached, but without any superfluous comparisons. The algorithm takes $\Theta(m + n)$ time, where the $\Theta(m)$ part comes from the pre-computation of the table.

23.1 Preliminaries: Difference Arrays

Our task is to take an imperative algorithm designed nearly half a century ago and express it in a functional style, retaining the possibility of efficient execution. Strictly speaking, there are two algorithms: the computation of the table, and the string search using the table. Neither would normally be seen as functional, but both algorithms are simple while loops, easily expressed as tail-recursive functions. Arrays are used, and random access is necessary. However, in the building phase, the table entries are added one after another, and the search does no array updates at all.

Because the original algorithms are imperative, their use of arrays is single-threaded. That means there is a single thread of updates starting from the initial value to the final array. It implies that updates can be done without copying: the previous array value can safely be destroyed. This conception can be realised by an ordinary array as supported by the hardware, augmented with a difference structure to deal with any array accesses that are not single-threaded. Provided there are none of those, performance can be good.

This data structure is called a difference array, and is part of the Collections framework [Lammich 2009]. This chapter uses the following notation for array operations:

- $A[n]$ to look up an array element (indexed from 0)
- $A[n := x]$ to update an array
- $|A|$ for the number of elements
- $\text{array } x n$ to create an $n$-element array, all elements filled with $x$

All but the last of these is assumed to take constant time.

23.2 Matches between Strings

A key concept is that of an $n$-character match between two strings $a$ and $b$, starting at positions $i$ and $j$, respectively (indexed from 0).

```
matches :: 'a array ⇒ nat ⇒ 'a array ⇒ nat ⇒ nat ⇒ bool
matches a i b j n
= (i + n ≤ |a| ∧ j + n ≤ |b| ∧ (∀ k<n. a !! (i + k) = b !! (j + k)))
```
Most of its properties are obvious. It always holds when \( n = 0 \), provided \( i \) and \( j \) lie within the range of their respective strings. A simple but valuable fact is weakening to get a shorter match: if \( \text{matches } a \, i \, b \, j \, n \) and \( k \leq n \) then

\[
\text{matches } a \, i \, b \, j \, k \quad \text{and} \quad \text{matches } a \, (i + k) \, b \, (j + k) \, (n - k).
\]

Sometimes we look for matches between the pattern \( p \) with the text \( a \), but when building the table we will be matching prefixes of \( p \) with other sections of \( p \).

### 23.3 The Next-Match Table

As noted above, the table identifies repetitions in the pattern that open the possibility that the current failed match may yet form part of a successful match. For example, suppose our search pattern \( p \) is \( xyzxyzxzxy \). And suppose we have matched \( xyzx \) in the string followed by a mismatch. The point is that the final \( x \) could be the start of an occurrence of \( p \) in the string. Similarly, if we have matched \( xyzxy \), \( xyzxyz \) or \( xyzxyzx \), the underlined section is a partial match of \( p \) and the search for a full match should continue from that point. But if we match \( xyzxyzx \), no suffix of this matches a prefix of \( p \). Finally, matching \( xyzxyzxzx \) let us use the final \( x \) as the start of a match. (Matching the whole of \( p \) would leave \( xy \) as the start of another possible match, but the algorithm below stops after the first.) Figure 23.1 illustrates the situation.

The corresponding next-match table is

\[
\begin{array}{ccccccccccc}
x & y & z & \underline{x} & y & z & \underline{x} & z & \underline{x} & y \\
0 & 0 & 0 & 1 & 2 & 3 & 4 & 0 & 1 \end{array}
\]

These numbers are indices into \( p \), numbering from 0. So for example 4 above tells us that at the position shown, we have successfully matched the first four characters of \( p \) and should start comparing at \( p[4] \), which is \( y \).

Now we are ready for the following predicate, which defines the next available match following a failed comparison:
is_next :: 'a array ⇒ nat ⇒ nat ⇒ bool

is_next p j n
= (n < j ∧ matches p (j − n) p 0 n ∧
   (∀m. n < m < j −→ ¬matches p (j − m) p 0 m))

In other words, n is the largest possible that is less than j and with an n-character match of a prefix of p with a substring of p ending at j.

The following two lemmas capture the essence of this. First, if the first j characters of the pattern already match (ending at position i in the text), and n is the next match, then indeed the first n characters of p match the text (again ending at i).

Lemma 23.1. matches a (i − n) p 0 n, provided

• matches a (i − j) p 0 j
• is_next p j n
• j ≤ i

Proof. We have matches a (i − n) p (j − n) n by weakening the given assumption. Moreover, we have matches p (j − n) p 0 n by the definition of is_next. The conclusion is immediate by transitivity.

The second lemma considers the same situation (a j-character match ending at i) and tells us that the ‘next match’, n, is really maximal: there does not exist a full match of p ending at k for any k, where i − j < k < i − n.

Lemma 23.2. ¬ matches a k p 0 ||p||, provided

• matches a (i − j) p 0 j
• is_next p j n
• j ≤ i
• i − j < k < i − n

Proof. Let m denote i − k. Then ¬ matches a (i − m) p 0 m by the definition of is_next and weakening. Further weakening using m < ||p|| yields the desired ¬ matches a (i − m) p 0 ||p||.

Therefore, using the next-match table to shift the pattern along will give us a partial match, which we can hope to complete, safe in the knowledge that there are no matches starting in the skipped-over region. All we have to do is build this table.
### 23.4 Building the Table: Loop Body and Invariants

Although this is a book of functional algorithms, here we basically have a while loop. Maintaining $j < i \leq ||p||$, it builds a match of the first $j$ characters of $p$ with a substring of $p$ ending at $i$, meanwhile filling the next table $nxt$ with the corresponding $j$ values. At a mismatch, it consults its own table—exactly as the main string search will do—for the longest possible match that still holds. In the imperative pseudo-code, $m$ denotes $||p||$, the length of $p$.

```plaintext
nxt[1] := 0; i := 1; j := 0;
while i < m-1 do
    if p[i] = p[j] then
        begin i := i+1; j := j+1; nxt[i] := j end
    else
        if j = 0 then begin i := i+1; nxt[i] := 0 end
        else j := nxt[j]
```

The loop body, expressed as a function, takes the pattern $p$ and the three loop variables $nxt$, $i$, $j$:

```plaintext
buildtab_step ::
    'a array ⇒ nat array ⇒ nat ⇒ nat ⇒ nat array × nat × nat
buildtab_step p nxt i j
= (if p !! i = p !! j then (nxt[i + 1 ::= j + 1], i + 1, j + 1)
     else if j = 0 then (nxt[i + 1 ::= 0], i + 1, j) else (nxt, i, nxt !! j))
```

To verify the while loop requires defining the loop invariant: a property of the loop variables that holds initially and is preserved in each iteration.

```plaintext
buildtabInvariant :: 'a array ⇒ nat array ⇒ nat ⇒ nat ⇒ bool
buildtabInvariant p nxt i j
= (||nxt|| = ||p|| ∧ i ≤ ||p|| ∧ j < i ∧ matches p (i - j) p 0 j ∧
   (∀ k. 0 < k ≤ i −→ is_next p k (nxt !! k)) ∧
   (∀ k. j + 1 < k < i + 1 −→ ¬ matches p (i + 1 − k) p 0 k))
```

It’s natural to regard this as the conjunction of six simpler invariants, some of which obviously hold, but some are nontrivial and depend on one another. The length of $nxt$ obviously doesn’t change, and since $i + 1 < ||p||$ holds prior to execution of...
the loop body, \( i \leq ||p|| \) holds and this inequality could even be strict. As for \( j < i \),
the critical case is when \( p !! i \neq p !! j \) and \( j > 0 \); the point is that \( \text{nxt} !! j < j \) by
the definition of \( \text{is}_\text{next} \) and the corresponding invariant. The invariant that we have
a match of length \( j \) has the same critical case and holds for the same reason.

We are left with two nontrivial invariants, and must prove they are preserved by
every execution of the loop body.

- That the next-match table is indeed built correctly (up to \( i \))
- That there cannot exist a match of length \( j+1 \) starting earlier in \( p \) than the
  match we have.

**Lemma 23.3.** \( \text{is}_\text{next} \; p \; k \; (\text{nxt}!! k) \), provided

\[
(nxt', i', j') = \text{buildtab}_\text{step} \; p \; \text{nxt} \; i \; j
\]

\[
\text{buildtab}_\text{invariant} \; p \; \text{nxt} \; i \; j
\]

\[
i + 1 < ||p||
\]

\[
0 < k \leq i'
\]

**Proof.** Consider \( \text{buildtab}_\text{step} \; p \; \text{nxt} \; i \; j \). If \( p !! i = p !! j \) then \( i' = i + 1 \) and \( j' = j + 1 \); then \( \text{matches} \; p \; (i - j) \; p \; 0 \; (j + 1) \) using the \( \text{matches} \) part of the invariant,

hence \( \text{is}_\text{next} \; p \; (i + 1) \; (j + 1) \) by definition and the prior invariant. Therefore, the
updated table, \( \text{nxt}' = \text{nxt}[i + 1 ::= j + 1] \), satisfies the conclusion.

So we can assume \( p !! i \neq p !! j \). If \( j = 0 \) then \( i' = i + 1 \). The character clash
implies \( \neg \text{matches} \; p \; (i - j) \; p \; 0 \; (j + 1) \) and therefore \( \text{is}_\text{next} \; p \; (i + 1) \; 0 \), validating
the updated next-match table, \( \text{nxt}' = \text{nxt}[i + 1 ::= 0] \). In the final case, when \( j > 0 \),
both \( i \) and \( \text{nxt} \) are left unchanged, making the conclusion trivial. \( \square \)

**Lemma 23.4.** \( \neg \text{matches} \; p \; (i' + 1 - k) \; p \; 0 \; k \), provided

\[
(nxt', i', j') = \text{buildtab}_\text{step} \; p \; \text{nxt} \; i \; j
\]

\[
\text{buildtab}_\text{invariant} \; p \; \text{nxt} \; i \; j
\]

\[
||p|| \geq 2
\]

\[
i + 1 < ||p||
\]

\[
j' + 1 < k < i' + 1
\]

**Proof.** Consider \( \text{buildtab}_\text{step} \; p \; \text{nxt} \; i \; j \). If \( p !! i = p !! j \) then \( i' = i + 1 \) and \( j' = j + 1 \); the conclusion follows from the same invariant for \( i \) and \( j \). So we can assume
\( p !! i \neq p !! j \). If \( j = 0 \) then we need to show

\( \neg \text{matches} \; p \; (i + 2 - k) \; p \; 0 \; k \) if \( 1 < k \) and \( k < i + 2 \).

The case \( k = 2 \) is immediate and otherwise it follows by instantiating the same
invariant with \( k - 1 \).
The remaining case is when \( p \mathrel{!!} i \neq p \mathrel{!!} j \) and \( j > 0 \). Then \( i' = i \) and \( j' = \text{nxt} \mathrel{!!} j \), so we need to show

\[ \neg \text{matches } p \mathrel{ (i + 1 - k) } p \mathrel{ 0 } k \text{ if } \text{nxt} \mathrel{!!} j + 1 < k \text{ and } k < i + 2. \]

This is trivial if \( k > j + 1 \) because the invariant holds beforehand, and if \( k = j + 1 \) because \( p \mathrel{!!} i \neq p \mathrel{!!} j \). So we can assume \( k \leq j \) and assume for contradiction that the match holds. Write \( k' = k - 1 \). Then we have

\[ \neg \text{matches } p \mathrel{ (j - k') } p \mathrel{ 0 } k', \text{ by the invariant } \text{is_next } p \mathrel{ j } (\text{nxt} \mathrel{!!} j) \]

\[ \text{matches } p \mathrel{ (j - k') } p \mathrel{ (i - k') } k', \text{ by the invariant } \text{matches } p \mathrel{ 0 } p \mathrel{ (i - j) } j \]

\[ \text{matches } p \mathrel{ (i - k') } p \mathrel{ 0 } k', \text{ weakening the negated conclusion} \]

The desired contradiction follows by the transitivity of \text{matches}. \( \Box \)

To summarize: we have proved that \text{buildtab\_invariant} is preserved by \text{buildtab}:

**Corollary 23.5.** \text{buildtab\_invariant } p \mathrel{ nxt' } i' j', \text{ provided}

- \((\text{nxt}', \text{i}', \text{j}') = \text{buildtab\_step } p \mathrel{ \text{nxt } i } j\)
- \text{buildtab\_invariant } p \mathrel{ \text{nxt } i } j
- \(i + 1 < \|p\|\)

## 23.5 Building the Table: Outer Loop

Now that we know that the loop body preserves the invariant, we are ready to define the actual function to build the next-match table. The loop itself is the obvious recursion:

```plaintext
buildtab :: 'a array ⇒ nat array ⇒ nat ⇒ nat ⇒ nat array
buildtab p nxt i j
= (if i + 1 < ||p||
    then let (nxt', i', j') = buildtab\_step p \text{nxt } i \text{ j} in buildtab p \text{nxt' } i' j'
    else nxt)
```

The key correctness property of the constructed table is not hard to prove. We must assume that the invariant holds initially.

**Lemma 23.6.** \text{is\_next } p \mathrel{ k } (buildtab p \mathrel{ \text{nxt } i } j \mathrel{!!} k), \text{ provided}

- \text{buildtab\_invariant } p \mathrel{ \text{nxt } i } j
- \(0 < k < ||p||\)
Proof by computation induction on \( \text{buildtab} \). If \( i + 1 < \|p\| \), \( \text{buildtab\_step} \) yields \((\text{nxt}', i', j')\) also satisfying the invariant (by Corollary 23.5) and by IH the result of the recursive call has the desired \( \text{is\_next} \) property. Conversely, if not \( i + 1 < \|p\| \), the invariant implies the desired property of \( \text{nxt} \).

It is convenient to define a top-level function to call \( \text{buildtab} \). It starts the loop with appropriate initial values, which can trivially be shown to establish the invariant, and catches a degenerate case to return a null table when \( p \) is trivial.

\[
\text{table} :: 'a\ \text{array} \Rightarrow \text{nat}\ \text{array} \\
\text{table} \ p = (\text{if } 1 < \|p\| \text{ then } \text{buildtab} \ p \ (\text{array} \ 0 \ \|p\|) \ 1 \ 0 \ \text{else } \text{array} \ 0 \ \|p\|)
\]

By Lemma 23.6 we have all we need to know about the table-building function:

\[0 < j < \|p\| \implies \text{is\_next} \ p \ j \ (\text{table} \ p \ !! \ j)\]  

(23.1)

### 23.6 Building the Table: Termination

It turns out that \( \text{buildtab} \) does not terminate on all inputs. For example, if \( i = 0 \), \( j = 1 \), \( \|p\| > 1 \), \( p \ !! \ i \neq p \ !! \ j \), \( p \ !! \ j = j \), then \( \text{buildtab\_step} \ p \ \text{nxt} \ i \ j = (\text{nxt}, i, j) \) and thus \( \text{buildtab} \) loops. We have not encountered non-termination before in this book and it raises two fundamental questions: is computation induction valid and can we even define \( \text{buildtab} \) in a logic of total functions, which HOL is.

Luckily, \( \text{buildtab} \) terminates on all inputs that satisfy the invariant: At every recursive call, either

- \( i \) increases by 1, with \( j \) unchanged or increased by 1, or
- \( i \) stays unchanged while \( j \) is replaced by \( \text{nxt} \ !! \ j \), and \( \text{nxt} \ !! \ j < j \) by the invariant.

In each of these cases, the integer quantity \( 2 \cdot \|p\| - 2 \cdot i + j \) decreases, and it is nonnegative because \( i \leq \|p\| \) by the invariant. Therefore, execution terminates, and the number of calls to \( \text{buildtab\_step} \) is linear in \( \|p\| \). Since each step—a couple of comparisons and a couple of assignments—clearly takes constant time, the overall running time is linear.

The proof of termination justifies the use of computation induction whenever we can assume that the invariant holds initially.

Defining functions that need non terminate is a subtle issue in a logic of total functions like HOL. Luckily, \( \text{buildtab} \) is tail-recursive (which is not a coincidence: every \text{while} loop corresponds to a tail-recursive function). That fact allows us to define \( \text{buildtab} \) without having to prove termination: it is consistent to assume the
existence of \( f \) satisfying \( f(x) = f(x+1) \), since any constant function will do, unlike the apparently similar \( f(x) = f(x+1) + 1 \).

We conclude this section with a formal counterpart of the above informal linear running time argument by means of a time function for \( \text{buildtab} \). Ironically, the very difficulty of \( \text{buildtab} \)'s termination proof complicates this step. Time functions are defined by equations of the form \( T_f \ p = T[e] + 1 \), which are not tail-recursive (if \( f \) occurs in \( e \)). For example, \( f(C\ x) = f\ x \) induces \( T_f(C\ x) = T_f\ x + 1 \). However, we can easily turn \( T_f \) into a tail-recursive function with an accumulating time parameter: \( T_f(C\ x, \ t) = T_f(x, \ t + 1) \). This leads to the following definition of \( T_{\text{buildtab}} \):

\[
T_{\text{buildtab}} :: \ 'a\ array \Rightarrow \ nat\ array \Rightarrow \ nat \Rightarrow \ nat \Rightarrow \ nat \Rightarrow \ nat
\]

\[
T_{\text{buildtab}}\ p\ \text{nxt}\ i\ j\ t = \begin{cases} 
(\text{if } i + 1 < ||p||) & \text{then let } (\text{nxt}',\ i',\ j') = \text{buildtab}_\text{step}\ p\ \text{nxt}\ i\ j \\
\text{in } T_{\text{buildtab}}\ p\ \text{nxt}'\ i'\ j'\ (t + 1) \\
\text{else } t 
\end{cases}
\]

The following result is proved similarly to Lemma 23.6.

**Lemma 23.7.** \( \text{buildtab}_\text{invariant} \ p\ \text{nxt}\ i\ j \longrightarrow \)
\( T_{\text{buildtab}}\ p\ \text{nxt}\ i\ j\ t \leq 2 \cdot ||p|| - 2 \cdot i + j + t \)

Plugging in the initial values, we find that

\[
2 \leq ||p|| \longrightarrow T_{\text{buildtab}}\ p\ (\text{array}\ 0\ ||p||)\ 1\ 0\ 0 \leq 2 \cdot (||p|| - 1)
\]

The precondition \( 2 \leq ||p|| \) is required because \( \text{buildtab}_\text{invariant} \) holds initially only in that case: \( 2 \leq ||p|| \longrightarrow \text{buildtab}_\text{invariant} \ p\ (\text{array}\ 0\ ||p||)\ 1\ 0 \)

The summary so far: we can build the next-match table, and in linear time. Now we are ready to search.

### 23.7 KMP String Search: Loop Body and Invariants

Like last time, let’s begin with a while loop and then analyse the corresponding functional version. In this pseudocode, \( m \) and \( n \) denote the lengths of \( p \) and \( a \), respectively. It closely resembles the previous algorithm, except it doesn’t build a table, and it compares \( p \) with \( a \) rather than with itself.
Chapter 23  Knuth–Morris–Pratt String Search

i := 0; j := 0; nxt := table(p);
while j<m and i<n do
  if a[i] = p[j] then
    begin i := i+1; j := j+1 end
  else
    if j = 0 then begin i := i+1 end
    else j := nxt[j];
if j=m then i-m else i

The last line returns the result of the algorithm: if \( j = m \), the whole pattern has been matched and \( i - m \) is the beginning of the (first) occurrence of the pattern; otherwise \( i \) will be \( n \), an indication that the pattern has not been found.

In the loop body, only \( i \) and \( j \) are modified, but the string, the pattern and the next-match table also need to be available. Hence the functional version takes all of them as arguments, but returns only the new values of \( i \) and \( j \):

\[
KMP\_step :: 'a array ⇒ nat array ⇒ 'a array ⇒ nat ⇒ nat × nat
\]

\[
KMP\_step p nxt a i j
= (if a !! i = p !! j then (i + 1, j + 1)
   else if j = 0 then (i + 1, 0) else (i, nxt !! j))
\]

Once again, we need an invariant relating these quantities, which must be preserved at every loop iteration. This invariant is simpler because the tough intellectual work has been done already. It asserts that there is a match between the first \( j \) characters of \( p \) and the text, ending at \( i \); moreover, there is no match of the whole of \( p \) with the text prior to that point.

\[
KMP\_invariant :: 'a array ⇒ 'a array ⇒ nat ⇒ nat ⇒ bool
\]

\[
KMP\_invariant p a i j
= (j ≤ ||p|| ∧ j ≤ i ∧ i ≤ ||a|| ∧ matches a (i - j) p 0 j ∧
  (∀ k<i - j: ¬ matches a k p 0 ||p||))
\]

This property is preserved in each step provided \( j < ||p|| \) and \( i < ||a|| \). If \( a !! i = p !! j \), or if \( j = 0 \), then the conclusion is trivial. The only interesting case is when \( a !! i ≠ p !! j \) and \( j > 0 \). Then we need to show the existence of a match of length \( nxt !! j \), but that is immediate by the already established correctness of the next-match table. Finally, we need to show \( ¬ matches a k p 0 ||p|| \) for \( k < i - nxt !! j \). We know
that \( k \neq i - j \) by the mismatch that just occurred, so either \( k < i - j \), when the result is immediate by the given invariant, or \( k > i - j \), when the result holds by Lemma 23.2.

### 23.8 KMP String Search: Outer Loop

Like last time, we express the while loop using recursion. The two active loop variables are \( i \) and \( j \), but the function takes additional arguments \( m \), \( n \) and \( nxt \) to prevent their being re-computed at every iteration. Their values will be \( \|p\|, \|a\| \), and table \( p \), respectively.

```plaintext
search :: nat ⇒ nat ⇒ nat array ⇒ 'a array ⇒ 'a array ⇒ nat ⇒ nat ⇒ nat × nat

search m n nxt p a i j
= (if j < m ∧ i < n
   then let (i', j') = KMP_step p nxt a i j in search m n nxt p a i' j'
   else (i, j))
```

The following function is the “top level” version, invoking the search loop with appropriate initial values. That includes building the table, and the loop invariant is established vacuously.

```plaintext
KMP_search :: 'a array ⇒ 'a array ⇒ nat × nat

KMP_search p a = search ||p|| ||a|| (table p) p a 0 0
```

Note that the definition of \( search \) raises the same termination problems we already faced with \( buildtab \). Termination again requires \( nxt !! j < j \). This time it follows from the correctness of table (23.1) if we know \( nxt = \text{table } p \).

### 23.9 KMP String Search: Correctness

The following predicate expresses the correctness of the result (as computed in the last line of the imperative algorithm). There are two possibilities. Termination before the end of the text string is reached (\( r < ||a|| \)) signifies success. Conversely, \( r = ||a|| \) implies failure.

```plaintext
first_occurrence :: 'a array ⇒ 'a array ⇒ nat ⇒ bool

first_occurrence p a r
= ((r < ||a|| → matches a r p 0 ||p||) ∧ (∀k< r. ¬ matches a k p 0 ||p||))
```
Lemma 23.8. \( \text{first\_occur } p \ a \ (\text{if } j' = ||p|| \ \text{then } i' - ||p|| \ \text{else } i') \), provided

- \((i', j') = \text{search } ||p|| \ ||a|| \ (\text{table } p) \ p \ a \ i \ j\)
- \(\text{KMP\_invariant } p \ a \ i \ j\)

Proof. By computation induction on \(\text{search}\). We have \(j \leq m\) and \(i \leq n\) by the invariant. If \(j < m\) and \(i < n\) then we obtain the result by IH (because \(\text{KMP\_step}\) preserves the invariant). Conversely, if \(j = m\) or \(i = n\) then the success or failure, respectively, follows by the invariant.

As a corollary we obtain correctness of \(\text{KMP\_search}\) because \(\text{KMP\_search}\) establishes \(\text{KMP\_invariant}\).

Corollary 23.9. \((i, j) = \text{KMP\_search } p \ a \rightarrow \text{first\_occur } p \ a \ (\text{if } j = ||p|| \ \text{then } i - ||p|| \ \text{else } i)\)

The proof of linearity of \(\text{search}\) is almost identical to that of Lemma 23.6, except that the quantity that decreases is \(2 \cdot ||a|| - 2 \cdot i + j\), which is nonnegative because \(i \leq ||a||\). Its initial value is \(2 \cdot ||a||\) because those of \(i\) and \(j\) are both zero. So the loop body can execute at most \(2 \cdot ||a||\) times. It’s not hard to see that this worst possible outcome occurs with the pathological string search mentioned at the beginning of this chapter. Even so, it is linear.

23.10 Chapter Notes

Acknowledgement. This development closely follows a formal verification of the Knuth–Morris–Pratt algorithm by Jean-Christophe Filliâtre using Why3. Due to the need for high performance in the era of gigabyte memories, innumerable variations exist. This version already achieves linear worst-case performance, and exhibits a pleasing symmetry between the table-building and search algorithms.

The original paper on KMP [Knuth et al. 1977], seemingly written by Knuth himself, is extremely clear. The realities of computing in the 1970s are evident in his suggestion that the string being searched might be held on an external file and that the naive search algorithm could introduce buffering issues, since after every failure of a match the algorithm would go back and rescans characters possibly no longer in main memory.
Huffman’s Algorithm

Jasmin Blanchette

Huffman’s algorithm [Huffman 1952] is a simple and elegant procedure for constructing a binary tree with minimum weighted path length—a measure of cost that considers both the lengths of the paths from the root to the leaf nodes and the weights associated with the leaf nodes. The algorithm’s main application is data compression: By equating leaf nodes with characters and weights with character frequencies, we can use it to derive optimum binary codes. A binary code is a map from characters to non-empty sequences of bits.

This chapter presents Huffman’s algorithm and its optimality proof. In a slight departure from the rest of this book, the emphasis is more on graphical intuitions and less on rigorous logical arguments.

24.1 Binary Codes

Suppose we want to encode strings over a finite source alphabet as sequences of bits. Fixed-length codes like ASCII are simple and fast, but they generally waste space. If we know the frequency \( w_a \) of each source symbol \( a \), we can save space by using shorter code words for the most frequent symbols. We say that a variable-length code is optimum if it minimizes the sum \( \sum_a w_a \delta_a \), where \( \delta_a \) is the length of the binary code word for \( a \).

As an example, consider the string ‘abacabad’. Encoding it with the code

\[
C_1 = \{ a \mapsto 0, b \mapsto 10, c \mapsto 110, d \mapsto 111 \}
\]

gives the 14-bit code word 01001100100111. The code \( C_1 \) is optimum: No code that unambiguously encodes source symbols one at a time could do better than \( C_1 \) on the input ‘abacabad’. With a fixed-length code such as

\[
C_2 = \{ a \mapsto 00, b \mapsto 01, c \mapsto 10, d \mapsto 11 \}
\]

we need at least 16 bits to encode the same string.

Binary codes can be represented by binary trees. For example, the trees
correspond to $C_1$ and $C_2$. The code word for a given symbol can be obtained as follows: Start at the root and descend toward the leaf node associated with the symbol one node at a time. Emit a 0 whenever the left child of the current node is chosen and a 1 whenever the right child is chosen. The generated sequence of 0s and 1s is the code word.

To avoid ambiguities, we require that only leaf nodes are labeled with symbols. This ensures that no code word is a prefix of another. Moreover, it is sufficient to consider only full binary trees (trees whose inner nodes all have two children), because any node with only one child can advantageously be eliminated by removing it and letting the child take its parent’s place.

Each node in a code tree is assigned a **weight**. For a leaf node, the weight is the frequency of its symbol; for an inner node, it is the sum of the weights of its subtrees. In diagrams, we often annotate the nodes with their weights.

### 24.2 The Algorithm

Huffman’s algorithm is a very simple procedure for constructing an optimum code tree for specified symbol frequencies. It works as follows: First, create a list of leaf nodes, one for each symbol in the alphabet, taking the given symbol frequencies as node weights. The nodes must be sorted in increasing order of weight. Second, pick the two trees

with the lowest weights and insert the tree
into the list so as to keep it ordered. Finally, repeat the process until only one tree is left in the list.

As an illustration, executing the algorithm for the frequencies \( f_d = 3, f_e = 11, f_f = 5, f_s = 7, f_z = 2 \) gives rise to the following sequence of states:

1. 
   \[
   \begin{array}{cccccc}
   z & d & f & s & e \\
   2 & 3 & 5 & 7 & 11 \\
   \end{array}
   \]

2. 
   \[
   \begin{array}{cccccc}
   5 & f & s & e \\
   z & d & f & s & e \\
   2 & 3 & 5 & 7 & 11 \\
   \end{array}
   \]

3. 
   \[
   \begin{array}{cccccc}
   s & f & e \\
   7 & 5 & 11 \\
   \end{array}
   \]

4. 
   \[
   \begin{array}{cccccc}
   e & 17 \\
   11 \\
   \end{array}
   \]
Chapter 24  Huffman’s Algorithm

24.3 The Implementation

The functional implementation of the algorithm relies on the following type:

```plaintext
datatype 'a tree = Leaf nat 'a | Node nat ('a tree) ('a tree)
```

Leaf nodes are of the form Leaf w a, where a is a symbol and w is the frequency associated with a, and inner nodes are of the form Node w t₁ t₂, where t₁ and t₂ are the left and right subtrees and w caches the sum of the weights of t₁ and t₂. The cachedWeight function extracts the weight stored in a node:

```plaintext
cachedWeight :: 'a tree ⇒ nat
cachedWeight (Leaf w _) = w
cachedWeight (Node w _ _) = w
```

The implementation builds on two additional auxiliary functions. The first one, uniteTrees, combines two trees by adding an inner node above them:

```plaintext
uniteTrees :: 'a tree ⇒ 'a tree ⇒ 'a tree
uniteTrees t₁ t₂ = Node (cachedWeight t₁ + cachedWeight t₂) t₁ t₂
```

The second function, insertTree, inserts a tree into a list sorted by cached weight, preserving the sort order:
24.4 Basic Auxiliary Functions Needed for the Proof

The main function that implements Huffman’s algorithm follows:

\[ \text{huffman} :: \text{'a tree list} \Rightarrow \text{'a tree} \]
\[ \text{huffman} \ [t] = t \]
\[ \text{huffman} \ (t_1 \# t_2 \# ts) = \text{huffman} \ (\text{insortTree} \ (\text{uniteTrees} \ t_1 \ t_2) \ ts) \]

The function should initially be invoked with a non-empty list of leaf nodes sorted by weight. It repeatedly unites the first two trees of the list it receives as argument until a single tree is left.

24.4 Basic Auxiliary Functions Needed for the Proof

This section introduces basic concepts such as alphabet, consistency and optimality, which are needed to state the correctness and optimality of Huffman’s algorithm. The next section introduces more specialized functions that arise in the proof.

The alphabet of a code tree is the set of symbols appearing in the tree’s leaf nodes:

\[ \text{alphabet} :: \text{'a tree} \Rightarrow \text{'a set} \]
\[ \text{alphabet} \ (\text{Leaf} \ _ \ a) = \{a\} \]
\[ \text{alphabet} \ (\text{Node} \ _ \ t_1 \ t_2) = \text{alphabet} \ t_1 \cup \text{alphabet} \ t_2 \]

A tree is consistent if for each inner node the alphabets of the two subtrees are disjoint. Intuitively, this means that a symbol occurs in at most one leaf node. Consistency is a sufficient condition for \( \delta_a \) (the length of the code word for \( a \)) to be uniquely defined. This well-formedness property appears as an assumption in many of the lemmas. The definition follows:

\[ \text{consistent} :: \text{'a tree} \Rightarrow \text{bool} \]
\[ \text{consistent} \ (\text{Leaf} \ _ \ _) = \text{True} \]
consistent \((\text{Node } t_1 \ t_2)\)
\[= (\text{alphabet } t_1 \cap \text{alphabet } t_2 = \{\} \land \text{consistent } t_1 \land \text{consistent } t_2)\]

The depth of a symbol (which we wrote as \(\delta_a\) above) is the length of the path from the root to that symbol, or equivalently the length of the code word for the symbol:

\[
\begin{align*}
\text{depth} &:: 'a \text{ tree} \Rightarrow 'a \Rightarrow \text{nat} \\
\text{depth} (\text{Leaf } _) &\_ = 0 \\
\text{depth} (\text{Node } t_1 \ t_2) a &\_ = \\
&= (\text{if } a \in \text{alphabet } t_1 \text{ then } \text{depth } t_1 \ a + 1 \\
&\quad \text{else if } a \in \text{alphabet } t_2 \text{ then } \text{depth } t_2 \ a + 1 \text{ else } 0)
\end{align*}
\]

By convention, symbols that do not occur in the tree or that occur at the root of a one-node tree are given a depth of 0. If a symbol occurs in several leaf nodes (of an inconsistent tree), the depth is arbitrarily defined in terms of the leftmost node labeled with that symbol.

The height of a tree is the length of the longest path from the root to a leaf node, or equivalently the length of the longest code word:

\[
\begin{align*}
\text{height} &:: 'a \text{ tree} \Rightarrow \text{nat} \\
\text{height} (\text{Leaf } _) &\_ = 0 \\
\text{height} (\text{Node } t_1 \ t_2) &\_ = \\
&= \max (\text{height } t_1) (\text{height } t_2) + 1
\end{align*}
\]

The frequency of a symbol (which we wrote as \(w_a\) above) is the sum of the weights attached to the leaf nodes labeled with that symbol:

\[
\begin{align*}
\text{freq} &:: 'a \text{ tree} \Rightarrow 'a \Rightarrow \text{nat} \\
\text{freq} (\text{Leaf } w \ a) b &\_ = (\text{if } b = a \text{ then } w \text{ else } 0) \\
\text{freq} (\text{Node } t_1 \ t_2) b &\_ = \text{freq } t_1 \ b + \text{freq } t_2 \ b
\end{align*}
\]

For consistent trees, the sum comprises at most one non-zero term. The frequency is then the weight of the leaf node labeled with the symbol, or 0 if there is no such node.

Two trees are comparable if they have the same alphabet and symbol frequencies. This is an important concept, because it allows us to state not only that the tree constructed by Huffman’s algorithm is optimal but also that it has the expected alphabet and frequencies.
The weight function returns the weight of a tree:

\[
weight :: \text{'}a\text{ tree} \Rightarrow \text{nat}
\]

\[
weight (\text{Leaf } w \_ ) = w
\]

\[
weight (\text{Node } \_ t_1 t_2) = weight t_1 + weight t_2
\]

In the Node case, we ignore the weight cached in the node and instead compute the tree’s weight recursively.

The cost (or weighted path length) of a consistent tree is the sum

\[
\sum_{a \in \text{alphabet } t} freq t a \cdot \text{depth } t a
\]

which we wrote as \( \sum_a w_a \delta_a \) above. It is defined recursively by

\[
cost :: \text{'}a\text{ tree} \Rightarrow \text{nat}
\]

\[
cost (\text{Leaf } \_ \_ ) = 0
\]

\[
cost (\text{Node } \_ t_1 t_2) = weight t_1 + cost t_1 + weight t_2 + cost t_2
\]

A tree is optimum iff its cost is not greater than that of any comparable tree:

\[
\text{optimum} :: \text{'}a\text{ tree} \Rightarrow \text{bool}
\]

\[
\text{optimum } t \\
= (\forall u. \text{consistent } u \land \text{alphabet } t = \text{alphabet } u \land \text{freq } t = \text{freq } u \rightarrow \\
\text{cost } t \leq \text{cost } u)
\]

Tree functions are readily generalized to lists of trees, or forests. For example, the alphabet of a forest is defined as the union of the alphabets of its trees. The forest generalizations have a subscript ‘F’ attached to their name (e.g. \textit{alphabet}_F).

### 24.5 Other Functions Needed for the Proof

The optimality proof needs to interchange nodes in trees, to replace a two-leaf subtree with weights \(w_1\) and \(w_2\) by a single leaf node of weight \(w_1 + w_2\) and vice versa, and to refer to the two symbols with the lowest frequencies. These concepts are represented by five functions: \textit{swapSyms}, \textit{swapFourSyms}, \textit{mergeSibling}, \textit{splitLeaf} and \textit{minima}.

The interchange function \textit{swapSyms} takes a tree \(t\) and two symbols \(a, b\), and exchanges the symbols:
Chapter 24  Huffman’s Algorithm

\(\text{swapSyms} :: \text{"a tree} \Rightarrow \text{"a} \Rightarrow \text{"a} \Rightarrow \text{"a tree}\)

\(\text{swapSyms} \ t \ a \ b = \text{swapLeaves} \ t \ (\text{freq} \ t \ a) \ a \ (\text{freq} \ t \ b) \ b\)

The following lemma captures the intuition that to minimize the cost, more frequent symbols should be encoded using fewer bits than less frequent ones:

**Lemma 24.1.** consistent \(t \wedge a \in \text{alphabet} \wedge b \in \text{alphabet} \wedge \text{freq} \ t \ a \leq \text{freq} \ t \ b \wedge \text{depth} \ t \ a \leq \text{depth} \ t \ b \rightarrow \text{cost} \ (\text{swapSyms} \ t \ a \ b) \leq \text{cost} \ t\)

The four-way symbol interchange function \(\text{swapFourSyms}\) takes four symbols \(a, b, c, d\) with \(a \neq b\) and \(c \neq d\), and exchanges them so that \(a\) and \(b\) occupy \(c\)'s and \(d\)'s positions. A naive definition of this function would be \(\text{swapSyms} \ (\text{swapSyms} \ t \ a \ c) \ b \ d\). This naive definition fails in the face of aliasing: If \(a = d\), but \(b \neq c\), then \(\text{swapFourSyms} \ a \ b \ c \ d\) would wrongly leave \(a\) in \(b\)'s position. Instead, we use this definition:

\(\text{swapFourSyms} :: \text{"a tree} \Rightarrow \text{"a} \Rightarrow \text{"a} \Rightarrow \text{"a} \Rightarrow \text{"a tree}\)

\(\text{swapFourSyms} \ t \ a \ b \ c \ d = (\text{if} \ a = d \ \text{then} \ \text{swapSyms} \ t \ b \ c\)

\(\text{else if} \ b = c \ \text{then} \ \text{swapSyms} \ t \ a \ d\)

\(\text{else} \ \text{swapSyms} \ (\text{swapSyms} \ t \ a \ c) \ b \ d)\)

Given a symbol \(a\), the \(\text{mergeSibling}\) function transforms the tree

\[
\begin{align*}
\text{mergeSibling} :: \text{"a tree} \Rightarrow \text{"a} \Rightarrow \text{"a tree} \\
\text{mergeSibling} \ (\text{Leaf} \ w \ b) \ _ &= \text{Leaf} \ w \ b \\
\text{mergeSibling} \ (\text{Node} \ w \ (\text{Leaf} \ w \ b) \ (\text{Leaf} \ w \ c)) \ a &= (\text{if} \ a = b \lor a = c \ \text{then} \ \text{Leaf} \ (w_b + w_c) \ a
\end{align*}
\]
else Node w (Leaf w\(_b\) b) (Leaf w\(_c\) c))
mergeSibling (Node w (Node v va vb) t\(_2\)) a
= Node w (mergeSibling (Node v va vb) a) (mergeSibling t\(_2\) a)
mergeSibling (Node w t\(_1\) (Node v va vb)) a
= Node w (mergeSibling t\(_1\) a) (mergeSibling (Node v va vb) a)

The \textit{sibling} function returns the label of the node that is the (left or right) sibling of the node labeled with the given symbol \(a\) in tree \(t\). If \(a\) is not in \(t\)'s alphabet or it occurs in a node with no sibling leaf node, we simply return \(a\). This gives us the nice property that if \(t\) is consistent, then \(\text{sibling t a} \neq a\) if and only if a has a sibling. The definition, which is omitted here, distinguishes the same cases as \(\text{mergeSibling}\).

Using the \textit{sibling} function, we can state that merging two sibling leaf nodes with weights \(w_a\) and \(w_b\) decreases the cost by \(w_a + w_b\):

\textbf{Lemma 24.2.} consistent \(t\) \& sibling \(t\) \(a\) \neq a \implies cost (mergeSibling \(t\) a) + freq \(t\) a + freq (sibling \(t\) a) = cost \(t\)

The \textit{splitLeaf} function undoes the merging performed by \(\text{mergeSibling}\): Given two symbols \(a\), \(b\) and two frequencies \(w_a\), \(w_b\), it transforms

\[
\begin{array}{c}
\text{into}
\end{array}
\]

In the resulting tree, \(a\) has frequency \(w_a\) and \(b\) has frequency \(w_b\). We normally invoke \(\text{splitLeaf}\) with \(w_a\) and \(w_b\) such that \(\text{freq } t\ a = w_a + w_b\). The definition follows:

\[
\begin{align*}
\text{splitLeaf} :: \text{'}a\text{ tree} & \Rightarrow \text{nat} \Rightarrow \text{'}a \Rightarrow \text{nat} \Rightarrow \text{'}a \Rightarrow \text{'}a\text{ tree} \\
\text{splitLeaf} (\text{Leaf } w_c\ c) \ & w_a\ a\ w_b\ b \\
& = (\text{if } c = a\ \text{then } \text{Node } w_c\ (\text{Leaf } w_a\ a)\ (\text{Leaf } w_b\ b)\ \text{else } \text{Leaf } w_c\ c) \\
\text{splitLeaf} (\text{Node } w\ t_1\ t_2) \ & w_a\ a\ w_b\ b \\
& = \text{Node } w\ (\text{splitLeaf } t_1\ w_a\ a\ w_b\ b)\ (\text{splitLeaf } t_2\ w_a\ a\ w_b\ b)
\end{align*}
\]

Splitting a leaf node with weight \(w_a + w_b\) into two sibling leaf nodes with weights \(w_a\) and \(w_b\) increases the cost by \(w_a + w_b\):
Lemma 24.3. \(\text{consistent } t \land a \in \text{alphabet } t \land \text{freq } t a = w_a + w_b \longrightarrow 
\text{cost (splitLeaf } t \ w_a \ a \ w_b \ b) = \text{cost } t + w_a + w_b\)

Finally, the \textit{minima} predicate expresses that two symbols \(a, b\) have the lowest frequencies in the tree \(t\) and that \(\text{freq } t a \leq \text{freq } t b\):

\[
\text{minima :: 'a tree } \Rightarrow \ 'a \Rightarrow 'a \Rightarrow \text{bool} \\
\text{minima } t \ a \ b \\
= (a \in \text{alphabet } t \land b \in \text{alphabet } t \land a \neq b \land \\
(\forall c\in\text{alphabet } t. \\
\quad c \neq a \rightarrow c \neq b \rightarrow \text{freq } t a \leq \text{freq } t c \land \text{freq } t b \leq \text{freq } t c))
\]

24.6 The Key Lemmas and Theorems

It is easy to prove that the tree returned by Huffman’s algorithm preserves the alphabet, consistency and symbol frequencies of the original forest:

\[
ts \neq [] \longrightarrow \text{alphabet } (\text{huffman } ts) = \text{alphabet}_F ts \\
\text{consistent}_F ts \land ts \neq [] \longrightarrow \text{consistent } (\text{huffman } ts) \\
ts \neq [] \longrightarrow \text{freq } (\text{huffman } ts) \ a = \text{freq}_F ts \ a
\]

The main difficulty is to prove the optimality of the tree constructed by Huffman’s algorithm. We need to introduce three lemmas before we can present the optimality theorem.

First, if \(a\) and \(b\) are minima and \(c\) and \(d\) are at the very bottom of the tree, then exchanging \(a\) and \(b\) with \(c\) and \(d\) does not increase the tree’s cost. Graphically, we have

\[
\text{cost } a \ b \ c \ d \ \leq \ \text{cost } a \ b \ c \ d
\]

Lemma 24.4. \(\text{consistent } t \land \text{minima } t \ a \ b \land 
\text{c } \in \text{alphabet } t \land \text{d } \in \text{alphabet } t \land 
\text{depth } t \ c = \text{height } t \land \text{depth } t \ d = \text{height } t \land c \neq d 
\longrightarrow 
\text{cost } (\text{swapFourSyms } t \ a \ b \ c \ d) \leq \text{cost } t\)
Proof by case analysis on \( a = c, a = d, b = c \) and \( b = d \). The cases are easy to prove by expanding the definition of \( \text{swapFourSyms} \) and applying Lemma 24.1.

The tree \( \text{splitLeaf} t \ w_a \ a \ w_b \ b \) is optimum if \( t \) is optimum, under a few assumptions, notably that \( \text{freq} t \ a = w_a + w_b \). Graphically:

\[
\begin{array}{c}
optimum \\
\text{a} \\
\Rightarrow \\
\text{optimum} \\
\text{b} \\
\end{array}
\]

Lemma 24.5. \text{consistent} \ t \land \text{optimum} \ t \land \\
a \in \text{alphabet} \ t \land b \notin \text{alphabet} \ t \land \text{freq} t \ a = w_a + w_b \land \\
(\forall c \in \text{alphabet} \ t. \ w_a \leq \text{freq} t \ c \land w_b \leq \text{freq} t \ c) \rightarrow \\
\text{optimum} (\text{splitLeaf} t \ w_a \ a \ w_b \ b)
\]

Proof. We assume that \( t \)'s cost is less than or equal to that of any other comparable tree \( v \) and show that \( \text{splitLeaf} t \ w_a \ a \ w_b \ b \) has a cost less than or equal to that of any other comparable tree \( u \). For the non-trivial case where \( \text{height} t > 0 \), it is easy to prove that there must be two symbols \( c \) and \( d \) occurring in sibling nodes at the very bottom of \( u \). From \( u \) we construct the tree \( \text{swapFourSyms} u \ a \ b \ c \ d \) in which the minima \( a \) and \( b \) are siblings:

\[
\begin{array}{c}
? \\
a \\
b \\
\end{array}
\]

The question mark reminds us that we hardly know anything about \( u \)'s structure. Merging \( a \) and \( b \) gives a tree comparable with \( t \), which we can use to instantiate \( v \):

\[
\begin{align*}
\text{cost} (\text{splitLeaf} t \ a \ w_a \ b \ w_b) &= \text{cost} t + w_a + w_b & \text{by Lemma 24.3} \\
&\leq \text{cost} (\text{mergeSibling} (\text{swapFourSyms} \ u \ a \ b \ c \ d) \ a) + w_a + w_b & \text{by optimality assumption} \\
&= \text{cost} (\text{swapFourSyms} \ u \ a \ b \ c \ d) & \text{by Lemma 24.2} \\
&\leq \text{cost} u & \text{by Lemma 24.4}
\end{align*}
\]
Once it has combined two lowest-weight trees using `uniteTrees`, Huffman’s algorithm does not visit these trees ever again. This suggests that splitting a leaf node before applying the algorithm should give the same result as applying the algorithm first and splitting the leaf node afterward.

**Lemma 24.6.**

\[
\text{consistent}_{F} ts \land ts \neq [] \land a \in \text{alphabet}_F ts \land \text{freq}_F ts \ a = w_a + w_b \rightarrow \\
\text{splitLeaf} \ (huffman \ ts) \ w_a \ a \ w_b \ b = huffman \ (\text{splitLeaf}_F \ ts \ w_a \ a \ w_b \ b)
\]

The proof is by straightforward induction on the length of the forest \(ts\).

As a consequence of this commutativity lemma, applying Huffman’s algorithm on a forest of the form

\[
\begin{array}{c}
\frac{c}{w_c} \\
\frac{a}{w_a} \ rac{b}{w_b}
\end{array}
\begin{array}{c}
\frac{d}{w_d} \\
\frac{z}{w_z}
\end{array}
\]
gives the same result as applying the algorithm on the “flat” forest

\[
\begin{array}{c}
\frac{c}{w_c} \\
\frac{a}{w_a + w_b}
\end{array}
\begin{array}{c}
\frac{d}{w_d}
\end{array}
\begin{array}{c}
\frac{z}{w_z}
\end{array}
\]

followed by splitting the leaf node \(a\) into two nodes \(a\) and \(b\) with frequencies \(w_a, w_b\).

The lemma provides a way to flatten the forest at each step of the algorithm.

This leads us to our main result.

**Theorem 24.7.**

\[
\text{consistent}_{F} ts \land \text{height}_F ts = 0 \land \text{sortedByWeight} ts \land ts \neq [] \rightarrow \\
\text{optimum} \ (huffman \ ts)
\]

**Proof** by induction on the length of \(ts\). The assumptions ensure that \(ts\) is of the form

\[
\begin{array}{c}
\frac{a}{w_a} \\
\frac{b}{w_b} \\
\frac{c}{w_c} \\
\frac{d}{w_d}
\end{array}
\begin{array}{c}
\frac{z}{w_z}
\end{array}
\]

with \(w_a \leq w_b \leq w_c \leq w_d \leq \cdots \leq w_z\). If \(ts\) consists of a single node, the node has cost 0 and is therefore optimum. If \(ts\) has length 2 or more, the first step of the algorithm leaves us with a term such as
In the diagram, we put the newly created tree at position 2 in the forest; in general, it could be anywhere. By Lemma 24.6, the above tree equals

\[
\text{splitLeaf} \left( \text{huffman} \begin{array}{ccc}
  & c & \\
  w_c & & w_d \\
 a & & b \\
 & w_a & w_b \\

d & & z \\
 & w_d & w_z \\
\end{array} \right) w_a \ a \ w_b \ b
\]

To prove that this tree is optimum, it suffices by Lemma 24.5 to show that

\[
\text{huffman} \begin{array}{ccc}
  & c & \\
  w_c & & w_d \\
 a & & b \\
 & w_a & w_b \\

d & & z \\
 & w_d & w_z \\
\end{array}
\]

is optimum, which follows from the induction hypothesis. \qed

In summary, we have established that the \text{huffman} program, which constitutes a functional implementation of Huffman's algorithm, constructs a binary tree that represents an optimal binary code for the specified alphabet and frequencies.

### 24.7 Chapter Notes

The sorted list of trees constitutes a simple priority queue (Part III). The time complexity of Huffman's algorithm is quadratic in the size \( n \) of this queue. By using a binary search to implement \text{insortTree}, we can obtain an \( O(n \log n) \) imperative implementation. An \( O(n) \) implementation is possible by maintaining two queues, one containing the unprocessed leaf nodes and the other containing the combined trees [Knuth 1997].

Huffman's algorithm was invented by Huffman [1952]. The proof above was inspired by Knuth's informal argument [Knuth 1997]. This chapter's text is based on a published article [Blanchette 2009], with the publisher's permission. An alternative formal proof, developed using Coq, is due to Théry [2004].

Knuth [1982] presented an alternative, more abstract view of Huffman's algorithm as a “Huffman algebra.” Could his approach help simplify our proof? The most tedious steps above concerned splitting nodes, merging siblings and swapping symbols. These steps would still be necessary, as the algebraic approach seems restricted to abstracting over the arithmetic reasoning, which is not very difficult in the first place. On the other hand, with Knuth's approach, perhaps the proof would gain in elegance.
This chapter is about searching the best possible move in a game tree. Alpha-beta pruning is a technique for decreasing the number of nodes that need to be examined by discarding whole subtrees during the search. There are many variations on this theme and we progress from the simple to the more sophisticated. We start by introducing the notion of a game tree.

### 25.1 Game Trees and Their Evaluation

A game tree represents a two-player game, such as tic-tac-toe or chess. Each node in the tree represents a possible position in the game. Each move is represented by an edge from one position to a child node, the successor position. There may be any number of successor positions and thus children. An example game tree is shown in Figure 25.1. In a two-player game, the players take turns. Thus each level in the tree...

![Tic-tac-toe game tree](image)
is associated with one of the two players, the one who is about to move, and this alternates from level to level. Leaf nodes in a game tree are terminal positions. The rules of the game must determine the outcome at a leaf, i.e. who has won or if it is a draw. More generally, what the value of that leaf is, because the game might involve, for example, money that one player loses and the other wins.

We model game trees by the following datatype:

```
datatype 'a tree = Lf 'a | Nd ('a tree list)
```

The interpretation: 'a is the type of values, Lf v is a leaf of value v and Nd ts is a node with a list of successor nodes ts. In an induction on trees, the induction step needs to prove \( P(Nd \ ts) \) under the IH that \( P \) is true for all \( t \) in \( ts \): \( \forall t \in set \ ts. \ P t. \)

Usually the type of values is fixed to be some numeric type extended with \( \infty \) and \( - \infty \), e.g. the extended real numbers (type `ereal` in Isabelle). Instead, we will only assume that 'a is a linear order with least and greatest elements \( \bot \) and \( \top \):

\[
\bot \leq a \quad a \leq \top
\]

This is a bounded linear order. Until further notice we assume that 'a is a bounded linear order. For concreteness, the reader is welcome to think in terms of some extended numeric type.

Type tree is an abstraction of an actual game tree (as in Figure 25.1) because the positions are not part of the tree. This is justified because we will only be interested in the value of a game tree, not the positions within it. Given a game tree, we want to find the best move for the start player, i.e. which of its successor nodes it should move to. Essentially equivalent is the question of the value of the game tree. This is the highest value of all leafs that the start player can reach, no matter what the opponent does, who will try to thwart those efforts as best as it can. Formally, there is a maximizing and a minimizing player. Thus the value of a game tree depends who is about to move. Function maxmin maximizes and minmax minimizes:

```
maxmin :: 'a tree ⇒ 'a
maxmin (Lf x) = x
maxmin (Nd ts) = mazs (map minmax ts)
```

```
minmax :: 'a tree ⇒ 'a
minmax (Lf x) = x
minmax (Nd ts) = mins (map maxmin ts)
```
The two evaluation functions `maxmin` and `minmax` should be considered the (executable) specification of what this chapter is about, namely more efficient evaluation functions that do not always examine the whole tree.

Figure 25.2 shows a game tree where each node is labeled with its value. The final level are the leaves. The squares are maximizing nodes, the circles are minimizing nodes. The value 3 at the root shows that the maximizer can reach a leaf of value at least 3, no matter which moves the minimizer chooses.

It is usually impossible to build a complete game tree because it is too large. Therefore the tree is typically only built up to some (possibly variable) depth. For simplicity we do not model this building process but start from the generated game tree where the leafs are not necessarily terminal positions (whose value would be determined by the rules of the game) but arbitrary ones where the tree building has stopped (e.g. due to some depth limit) and the value is give by some heuristic evaluation function. However, by starting with a game tree we abstract from all of these issues.

### 25.2 Alpha-Beta Pruning

The idea underlying alpha-beta pruning is, in the simplest case, this: if the maximizer finds a move that leads to a definite win, the remaining moves need not be considered
anymore. More generally, if the maximizer finds that some move from some (non-root) node $B$ leads to a higher value than the value computed for a previously evaluated sibling $A$ of $B$, it can stop exploring the successors of $B$ because the minimizer would always move to $A$ rather than $B$ to force the lower outcome. This situation is exemplified in Figure 25.3 (the tree with the by now familiar leaf sequence, but evaluated with alpha-beta pruning; ignore the $a,b$ labels for now) when looking at the subtree with root 5. Conversely, the minimizer can stop exploring successors of a (non-root) node $B$ if it finds a move that leads to a lower value than the value of some sibling of $B$. This is what happened at node 1.

![Figure 25.3 Alpha-beta pruning](image)

Alpha-beta pruning is more general still by keeping track of two bounds. It is parameterized by two values $a$ and $b$ (or $\alpha$ and $\beta$) such that $a$ is the maximum value that the maximizer is already assured of and $b$ is the minimum value that the minimizer is already assured of (by the search so far, assuming optimal play by both players). The maximizer searches its successor positions and increases $a$ accordingly. Once $a \geq b$, the search at this level can stop: if $a > b$, the minimizer would never allow the maximizer to reach the parent node because the minimizer can already enforce $b$ elsewhere; if $a = b$, the minimizer will only allow the maximizer to reach the parent node if the remaining successor positions do not yield a value $> a$. In summary, the open interval from $a$ to $b$ is the window in which alpha-beta pruning searches for nodes that increase $a$ until the interval becomes empty. Dually for the minimizer. This is the actual code:

```
ab_max :: 'a => 'a => 'a tree => 'a
ab_max _ _ (Lf x) = x
ab_max a b (Nd ts) = ab_maxs a b ts

ab_maxs :: 'a => 'a => 'a tree list => 'a
```
\[\begin{align*}
\text{ab\_maxs} & \ a \ [ ] \ = \ a \\
\text{ab\_maxs} & \ ab \ (t \ \# \ ts) \\
& = (\text{let} \ a' = \max \ a \ (\text{ab\_min} \ a \ b \ t) \ \text{in} \ \text{if} \ b \ \leq \ a' \ \text{then} \ a' \ \text{else} \ \text{ab\_maxs} \ a' \ b \ ts)
\end{align*}\]

\[\begin{align*}
\text{ab\_min} & \ : \ 'a \ \Rightarrow \ 'a \ \Rightarrow \ 'a \ \text{tree} \ \Rightarrow \ 'a \\
\text{ab\_min} _ _ (Lf \ x) & = x \\
\text{ab\_min} \ a \ b (Nd \ ts) & = \text{ab\_mins} \ a \ b \ ts
\end{align*}\]

\[\begin{align*}
\text{ab\_mins} & \ : \ 'a \ \Rightarrow \ 'a \ \Rightarrow \ 'a \ \text{tree list} \ \Rightarrow \ 'a \\
\text{ab\_mins} _ b [ ] & = b \\
\text{ab\_mins} & \ ab \ (t \ \# \ ts) \\
& = (\text{let} \ b' = \min \ b \ (\text{ab\_max} \ a \ b \ t) \ \text{in} \ \text{if} \ b' \ \leq \ a \ \text{then} \ b' \ \text{else} \ \text{ab\_mins} \ a \ b' \ ts)
\end{align*}\]

There are more compact ways to formulate these functions (see Exercises 25.6 and 25.7) but the explicitness of the above code leads to more elementary proofs where the \text{min} cases are completely dual to the \text{max} cases. If we only consider one of the two cases in a definition, a lemma or a proof, the other one is completely dual. An example is this simple inductive property of \text{ab\_maxs}

\[a \ \leq \ \text{ab\_maxs} \ ab \ ts \quad (25.1)\]

where we leave the dual property of \text{ab\_mins} unstated.

Alpha-beta pruning implicitly assumes \(a < b\) and many of its properties only hold under that assumption, property (25.1) being an exception.

### 25.2.1 Correctness and Proof

This is the top-level correctness property we want in the end:

\[\text{ab\_max} \ \bot \ t = \max\min \ t \quad (25.2)\]

Of course, a proof will require a generalization from \(\bot\) and \(\top\) to arbitrary \(a\) and \(b\). Unsurprisingly, \text{ab\_max} \text{a b t = max\min t} does not hold in general. For example, \text{ab\_max} \text{1 2 (Nd [Lf 0]) = 1 but max\min (Nd [Lf 0]) = 0. Thus we first need to find a suitable generalization of (25.2).}\\n
The following relations between \text{ab\_max} and \text{max\min} state that \text{ab\_max} coincides with \text{max\min} for values inside the \((a, b)\) interval and that \text{ab\_max} bounds \text{max\min} outside that interval:

\[\begin{align*}
\text{ab\_max} \ ab \ b \ t & \ \leq \ a \rightarrow \ \max\min \ t \ \leq \ \text{ab\_max} \ ab \ b \ t \\
 a & \ < \ ab\_max \ ab \ b \ t \ < \ b \rightarrow \ \text{ab\_max} \ ab \ b \ t = \max\min \ t \\
\text{ab\_max} \ ab \ b \ t & \ \geq \ b \rightarrow \ \max\min \ t \ \geq \ \text{ab\_max} \ ab \ b \ t
\end{align*}\]
These properties do not specify \texttt{ab\_max} uniquely but they are strong enough to imply (as we see below) the key correctness property (25.2).

To facilitate the further discussion, we define the following abbreviation (due to Fishburn [1983]):

\[
\text{fishburn } a \ b \ v \ ab \\
(ab \leq a \rightarrow v \leq ab) \land \ \\
(a < ab \land ab < b \rightarrow ab = v) \land \ \\
(b \leq ab \rightarrow ab \leq v)
\]

The conjunction of (25.3)–(25.5) is \texttt{fishburn } a \ b \ (\texttt{maxmin } t) \ (\texttt{ab\_max } a \ b \ t).

Although \texttt{fishburn} is a relation, it can also be read as a function that tells us in which of the three intervals (not lists!) \([\bot,ab],[ab,ab]\) or \([b,\top]\) \(v\) is located, depending on where \(ab\) lies w.r.t. \(a\) and \(b\).

Correctness can now be shown simultaneously for all four functions:

\textbf{Theorem 25.1.}
\begin{align*}
a < b & \rightarrow \text{fishburn } a \ b \ (\texttt{maxmin } t) \ (\texttt{ab\_max } a \ b \ t) \\
a < b & \rightarrow \text{fishburn } a \ b \ (\texttt{maxmin } (Nd \ ts)) \ (\texttt{ab\_maxs } a \ b \ ts) \\
a < b & \rightarrow \text{fishburn } a \ b \ (\texttt{minmax } t) \ (\texttt{ab\_min } a \ b \ t) \\
a < b & \rightarrow \text{fishburn } a \ b \ (\texttt{minmax } (Nd \ ts)) \ (\texttt{ab\_mins } a \ b \ ts)
\end{align*}

\textit{Proof} by simultaneous induction on the computation of \texttt{ab\_max} and friends. The only two nontrivial cases are the ones stemming from the recursion equations for \texttt{ab\_maxs} and \texttt{ab\_mins}. We concentrate on \texttt{ab\_maxs}. For succinctness we introduce the following abbreviations:

\[
\text{abt } \equiv \text{ab\_min } a \ b \ t \quad \text{abts } \equiv \text{ab\_maxs } a \ b \ ts \quad \text{a’ } \equiv \text{max } a \ abt \\
\text{vt } \equiv \text{minmax } t \quad \text{vts } \equiv \text{maxmin } (Nd \ ts)
\]

The two IHs are

\[
\text{fishburn } a \ b \ vt \ abt \quad \text{IH1} \\
a’ < b \rightarrow \text{fishburn } a’ \ b \ vts \ abts \quad \text{IH2}
\]

and we need to prove \texttt{fishburn } a \ b \ vts \ abts where

\[
\text{abts } \equiv \text{ab\_maxs } a \ b \ (t \ # \ ts) \quad \text{vts } \equiv \text{maxmin } (Nd \ (t \ # \ ts)) = \text{max } vt \ vts
\]

We focus on the most complex part of \texttt{fishburn } a \ b \ vts \ abts, conjunct 2. That is, we assume \(a < abts < b\) and prove \(abts = vts\) by case analysis. The case \(b \leq a’\) is impossible because it would imply \(a’ = abts\), which, combined with the assumption
\( \text{abtts} < b \), would imply \( b < b \). Hence we can assume \( a' < b \) and thus \( \text{abtts} = \text{abts} \) and \( a < \text{abts} < b \). Hence we now need to prove
\[
\text{abts} = \max vt \upsilon ts
\]

For the following detailed arguments we display and name the relevant conjuncts of IH1 and IH2 (where the premise \( a' < b \) is now assumed):
\[
\begin{align*}
\text{abt} \leq a & \quad \rightarrow \quad vt \leq \text{abt} \quad \text{(IH11)} \\
\text{a} < \text{abt} < b & \quad \rightarrow \quad \text{abt} = vt \quad \text{(IH12)} \\
\text{abts} \leq a' & \quad \rightarrow \quad \upsilon ts \leq \text{abts} \quad \text{(IH21)} \\
\text{a}' < \text{abts} < b & \quad \rightarrow \quad \text{abts} = \upsilon ts \quad \text{(IH22)}
\end{align*}
\]

The proof continues with a case analysis. First assume \( \text{abt} \leq a \). Hence \( a' = a \) and thus IH22 and \( a < \text{abts} < b \) yield \( \text{abts} = \upsilon ts \). Moreover, \( vt \leq \upsilon ts \) follows from IH11, \( \text{abt} \leq a \), \( a < \text{abts} \) and \( \text{abts} = \upsilon ts \). Together this proves \( \text{abts} = \max vt \upsilon ts \).

Now assume \( a < \text{abt} \). This implies \( a' = \text{abt} \), \( \text{abt} = vt \) (using IH12) and \( \text{abt} < b \) (using \( a' < b \)). From (25.1) we obtain \( a' \leq \text{abts} \) and perform another case analysis. First assume \( a' < \text{abts} \). Because \( \text{abts} < b \), IH22 yields \( \text{abts} = \upsilon ts \). Assumption \( a' < \text{abts} \) implies \( \text{tab} < \text{abts} \) and thus \( vt < \upsilon ts \) which proves \( \text{abts} = \max vt \upsilon ts \). Now assume \( a' = \text{abts} \). IH21 implies \( \upsilon ts \leq \text{abts} \). Moreover, \( \text{abts} = a' = \text{abt} = vt \). Together this implies \( \text{abts} = \max vt \upsilon ts \).

The top-level correctness property \( \text{ab} \_\text{max} \perp \top \ t = \text{maxmin} \ t \) (25.2) is a consequence of (25.6) where \( a = \perp \) and \( b = \top \). Let us first deal with the standard case that \( \perp < \top \). Then (25.6) yields \( \text{fishburn} \ a \ b \ (\text{maxmin} \ t) \ (\text{ab} \_\text{max} \ a \ b \ t) \). The claim \( \text{ab} \_\text{max} \perp \top \ t = \text{maxmin} \ t \) follows from this general property of \( \text{fishburn} \)
\[
\text{fishburn} \perp \top \ x \ y \rightarrow x = y
\]

which is easy to prove: If \( \perp < y < \top \), the definition of \( \text{fishburn} \) yields the result directly. If \( y \leq \perp \) then the definition of \( \text{fishburn} \) implies \( x \leq y \leq \perp \) and uniqueness of \( \perp \) yields \( y = x = \perp \). The case \( y \geq \top \) is dual.

Now consider the corner case which does not arise for numeric types, namely
\( \neg \perp < \top \) In that case, everything collapses and (25.2) trivially holds:
\[
\neg \perp < \top \rightarrow x = y
\]

The proof is left as an exercise.

25.2.2 Fail-Soft

Function \( \text{ab} \_\text{maxs} \) is less precise than it could be: \( \text{ab} \_\text{maxs} \ a \ b \ ts = a \) even if \( \text{ab} \_\text{min} \ a \ b \ t < a \) for all \( t \in \text{set} \ ts \). But in this case \( \text{maxmin} (N\!d \ ts) < a \) and
Chapter 25  Alpha-Beta Pruning

ab_maxs could have produced a better bound for maxmin (Nd ts) if it did not return a but ⊥ at the end of the list. These are the improved ab_max functions:

\[
\begin{align*}
ab_max' :: \text{'a} \Rightarrow \text{'a} \\
ab_max' \_ \_ (Lf x) &= x \\
ab_max' a b (Nd ts) &= ab_maxs' a b \bot ts \\
\end{align*}
\]

In the literature, ab_maxs is called the fail-hard variant (because it brutally cuts off at a) and ab_maxs' the fail-soft variant (because it "fails" more gracefully).

For a start we have that ab_max' bounds maxmin (and is thus correct w.r.t. maxmin):

**Theorem 25.2.** \(a < b \rightarrow \text{fishburn a b (maxmin t) (ab_max' a b t)}\)

\[\max m a < b \rightarrow \text{fishburn (max m a) b (maxmin (Nd ts)) (ab_maxs' a b m ts)}\]

This is similar to the correctness theorem for ab_max but slightly more involved because of the additional parameter of ab_max'. The proof is also similar, including the need for the lemmas \(m \leq ab_maxs' a b m ts\) and \(ab_mins' a b m ts \leq m\).

Moreover, ab_max bounds ab_max':

**Theorem 25.3.** \(a < b \rightarrow \text{fishburn a b (ab_max' a b t) (ab_max a b t)}\)

\[\max m a < b \rightarrow \text{fishburn a b (ab_maxs' a b m ts) (ab_maxs (max m a) b ts)}\]

The proof is similar to that of the previous theorem but requires no lemmas.

In summary, we now know that ab_max' bounds maxmin at least as precisely as ab_max does. In fact, it can be more precise, as the following example shows:

\[ab_max' 0 1 (Nd []) = \text{maxmin (Nd [])} = \bot \text{ but } ab_max 0 1 (Nd []) = 0 > \bot.\]

Both variants search the same part of the trees. To verify this, we define functions that return the part of the trees that ab_max(') and ab_maxs(') traverse.

\[
\begin{align*}
\text{abt_max :: 'a} \Rightarrow 'a & \Rightarrow 'a \text{ tree} \Rightarrow 'a \\
\text{abt_max} \_ \_ (Lf x) &= Lf x \\
\text{abt_max} a b (Nd ts) &= Nd (ab_maxs a b ts) \\
\end{align*}
\]
25.2 Alpha-Beta Pruning

Indeed, they search the same part of the trees:

**Theorem 25.4.** \( a \lt b \implies abt_{\max'} a b t = abt_{\max} a b t \)
\( \max m a \lt b \implies abt_{\max's} a b m t s = abt_{\maxs} (\max m a) b t s \)

The proof is the usual simultaneous induction and relies on Theorem 25.3.

The following section answers the question how the improved precision of the soft variant can be exploited to optimize the search further.

### 25.2.3 From Trees to Graphs

Game trees are in fact graphs, because different paths may lead to the same position. Moreover, positions have symmetries, and different positions may be equivalent, for example by rotating or reflecting the board. For efficiency reasons it is vital to factor in these symmetries when searching the graph. This is usually taken care of by a so-called transposition table, which is a cache for storing evaluations of previously seen positions (modulo symmetries). However, evaluations of the same position from different parts of the graph typically come with different a,b windows. Nevertheless, the result of a previous evaluation can help to narrow the a,b window in later evaluations of the same position. In the following little lemma, we assume that \( abf :: 'a \Rightarrow 'a \Rightarrow 'a \ tree \ list \Rightarrow 'a \ tree \ list \)

\[ \forall a b. \ \text{fishburn} a b (\maxmin t) (ab a b t) \]  

\[ (*) \]
If in a previous call $b \leq ab a b t$, then (*) implies $abf a b t \leq maxmin t$. Thus $abf a b t$ can be used as a lower bound for future $abf$ calls. That is, in a call $abf a' b' t$ we can replace $a'$ by $max a'$ ($abf a b t$), provided this does not push us above $b'$ (in which case there is no need to call $abf$ again):

$$b \leq ab a b t \land max a' (ab a b t) < b' \rightarrow \text{fishburn } a' b' (maxmin t) (ab (max a' (ab a b t)) b' t)$$

Similarly, if $ab a b t \leq a$, then $abf a b t$ can be used as an upper bound for future $abf$ calls, i.e. we can replace $b'$ by $min b'$ ($abf a b t$). Hence $ab_max'$ has the edge over $ab_max$ in this scenario: it can lead to smaller search windows.

Of course, if $a < abf a b t < b$, then $abf a b t = maxmin t$ and we can return the exact value right away.

The advantage of narrowing the $a,b$ window is that the search space decreases. The intuitive reason is clear: as $b$ decreases, $a$ will reach it more quickly (and conversely). More precisely, the search space with a smaller window is a prefix of that with the larger window in the following sense:

```
prefix :: 'a tree ⇒ 'a tree ⇒ bool
prefix (Lf x) (Lf y) = (x = y)
prefix (Nd ts) (Nd us) = prefixes ts us
prefix _ _ = False

prefixs :: 'a tree list ⇒ 'a tree list ⇒ bool
prefixs [] _ = True
prefixs (t # ts) (u # us) = (prefix t u ∧ prefixes ts us)
prefixs (_ # _) [] = False
```

Now we can employ the $ab$ functions to obtain the searched space:

**Theorem 25.5.**

$$a < b \land a' \leq a \land b \leq b' \rightarrow prefix (abt_max' a b t) (abt_max' a' b' t)$$

$$max m a < b \land a' \leq a \land b' \land m' \leq m \rightarrow$$

$$prefixs (abt_max's a b m ts) (abt_max's a' b' m' ts)$$

The proof is by the usual computation induction but also requires a lemma. It expresses that when we narrow the search window, the result becomes less precise:

**Lemma 25.6.**

$$a < b \land a' \leq a \land b \leq b' \rightarrow \text{fishburn } a b (ab_max' a' b' t) (ab_max' a b t)$$
max m a < b ∧ a' ≤ a ∧ b ≤ b' ∧ m' ≤ m →
fishburn (max m a) b (ab_maxs' a' b' m' ts) (ab_maxs' a b m ts)

This lemma can be proved directly, i.e. without requiring further lemmas.

### 25.2.4 Exercises

Exercise 25.1. We can get away without ⊥ and ⊤ if we require that the list of successor positions, i.e. the arguments of \(N_d\), are nonempty. Formalize this requirement as a predicate \(invar :: 'a tree ⇒ \text{bool}\), define new versions of \(maxs\), \(mins\), \(maxmin\) and \(minmax\) (without using \(⊥\) and \(⊤\)) and prove \(invar t → maxmin_1 t = maxmin t\) (where the new versions are distinguished by an appended 1).

Exercise 25.2. Prove that \(fishburn\) can be expressed by a chain of inequations:

\[
a < b → fishburn a b v ab ←→ min v b ≤ ab ∧ ab ≤ max v a
\]

Exercise 25.3. Consider this slightly weaker version of \(fishburn\) due to Knuth and Moore [1975]:

\[
knuth a b x y ≡ (y ≤ a → x ≤ a) ∧ (a < y ∧ y < b → y = x) ∧ (b ≤ y → b ≤ x)
\]

Similar to \(fishburn\) we have \(knuth\) \(⊥ \top x y → x = y\). Prove

\[
a < b → knuth a b (maxmin t) (ab_max a b t)
\]

following the proof of Theorem 25.1. Do not simply employ that \(fishburn a b x y\) implies \(knuth a b x y\).

Exercise 25.4. Consider the operation \(max a (min x b)\) that squashes \(x\) into the closed interval \([a,b]\) (assuming \(a ≤ b\)) by returning \(a\) if \(x < a\) and \(b\) if \(x > b\) and leaving \(x\) unchanged otherwise. Note that if \(a ≤ b\), then the order of \(max\) and \(min\) is irrelevant: \(a ≤ b → max a (min x b) = min b (max x a)\).

Prove that with the help of this operation, \(knuth\) (see Exercise 25.3) can be expressed purely equationally:

\[
a < b → max a (min x b) = max a (min y b) ←→ knuth a b y x
\]

Because the \(max/min\) equation is symmetric in \(x\) and \(y\), it follows that \(knuth\) is symmetric as well: \(a < b → knuth a b x y ←→ knuth a b y x\).

Exercise 25.5. Consider the \(max a (min x b)\) operation from Exercise 25.4 and modify \(ab_max(s)\) (and analogously \(ab_min(s)\)) as follows:
Chapter 25  Alpha-Beta Pruning

\(\text{ab\_max} \colon 'a \Rightarrow 'a \Rightarrow 'a\)  
\(\text{ab\_max} \ a \ b \ (Lf \ x) = \text{max} \ a \ (\text{min} \ x \ b)\)  
\(\text{ab\_max} \ a \ b \ (N\_d \ t\_s) = \text{ab\_max\_s} \ a \ b \ t\_s\)

\(\text{ab\_max\_s} \colon 'a \Rightarrow 'a \Rightarrow 'a\)  
\(\text{ab\_max\_s} \ a \ [] = a\)  
\(\text{ab\_max\_s} \ a \ b \ (t \ # \ t\_s) = (\text{let} \ a' = \text{ab\_min} \ a \ b \ t \ \text{in} \ \text{if} \ a' = b \ \text{then} \ a' \ \text{else} \ \text{ab\_max\_s} \ a' \ b \ t\_s)\)

Both \(\text{max}\) and \(\text{min}\) have moved to the \(Lf\) cases, thus assuring that the result of all \(\text{ab}\) functions lies in the closed interval \([a, b]\). Prove the following correctness theorem

\[ a \leq b \implies \text{ab\_max} \ a \ b \ t = \text{max} \ a \ (\text{min} \ (\text{max\_min} \ t) \ b)\]

The corollary \(\text{ab\_max} \bot \ T \ t = \text{max\_min} \ t\) is immediate.

The following exercises are concerned with more compact definitions exploiting the symmetries between maximizer and minimizer.

Exercise 25.6. The functions \(\text{ab\_max}\) and \(\text{ab\_min}\) and the functions \(\text{ab\_max\_s}\) \(\text{ab\_min\_s}\) are completely dual to each other: exchange \(\text{min}\) and \(\text{max}\), \((\leq)\) and \((\geq)\) and which parameter (\(a\) or \(b\)) is modified in the recursive call. All of this can be captured uniformly by making \((\leq)\) a parameter, expressing \(\text{max}/\text{min}\) by means of \((\leq)\) and by exchanging \(a\) and \(b\) when switching between maximizer and minimizer.

Define two functions

\(\text{ab\_le} \colon ('a \Rightarrow 'a \Rightarrow \text{bool}) \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a\)
\(\text{ab\_les} :: ('a \Rightarrow 'a \Rightarrow \text{bool}) \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a\)

and prove

\(\text{ab\_max} \ a \ b \ t = \text{ab\_le} \ (\leq) \ a \ b \ t\)
\(\text{ab\_max\_s} \ a \ b \ t\_s = \text{ab\_les} \ (\leq) \ a \ b \ t\_s\)
\(\text{ab\_min} \ a \ b \ t = \text{ab\_le} \ (\lambda x \ y. \ y \leq x) \ a \ b \ t\)
\(\text{ab\_min\_s} \ a \ b \ t\_s = \text{ab\_les} \ (\lambda x \ y. \ y \leq x) \ a \ b \ t\_s\)

Exercise 25.7. Functions \(\text{max}\_\text{min}\) and \(\text{min}\_\text{max}\) (and friends) exhibit the same symmetries as \(\text{ab\_max}\) and friends. Define a single function \(\text{max}\_\text{min\_le}\) that takes a comparison operation \(\text{le}\) (and maybe more) and behaves like \(\text{max}\_\text{min}\) or \(\text{min}\_\text{max}\), depending on the parameter \(\text{le}\). Prove \(\text{max}\_\text{min\_le} \ldots (\leq) \ t = \text{max}\_\text{min} \ t\) (and more). Follow the approach for \(\text{ab\_le}\) in Exercise 25.6.
25.3 Negative Values

In this section we examine a popular approach to exploiting the symmetries between maximizer and minimizer. As a result, we only need two instead of four functions, both for game tree evaluation and alpha-beta pruning. It can be seen as another variation of the approaches sketched in Exercises 25.6 and 25.7. This time we exploit the symmetries between positive and negative values. A value $v$ for one player can be viewed as a value $-v$ for the other player: one player’s gain is the other player’s loss. This seems to work only for numeric value types, but it turns out that the following properties are sufficient to make it work more generally:

\[
- \min x y = \max (-x) (-y) \\
- (-x) = x
\] (25.7)

We call a bounded linear order satisfying the above two properties a de Morgan order because of the first de-Morgan-like property. For the rest of this section, we assume that ‘a is a de Morgan order. For concreteness you may think of the extended reals. Of course de Morgan orders satisfy many other properties that follow easily, in particular the dual de Morgan property

\[
- \max x y = \min (-x) (-y)
\]

We will not list them here because they are all familiar from extended numeric types.

25.3.1 Game Tree Evaluation

With the help of negation we can unify the evaluation functions $maxmin$ and $minmax$ into a single function $negmax$:

\[
\begin{align*}
\text{negmax} & : 'a \text{ tree} \Rightarrow 'a \\
\text{negmax} (Lf x) & = x \\
\text{negmax} (Nd ts) & = \maxs (\text{map} (\lambda t. - \text{negmax} t) ts)
\end{align*}
\]

Figure 25.4 shows the evaluation of the same tree as in Figure 25.2 but with $negmax$. We have to negate the leaves because they belong to the minimizer but the root (which we evaluate) to the maximizer.

Function $negate \ b \ t$ performs the negation of the minimizer leaves of $t$, where $b = True$ iff the root of $t$ is a minimizer level.

\[
\begin{align*}
negate & : \text{bool} \Rightarrow 'a \text{ tree} \Rightarrow 'a \text{ tree}
\end{align*}
\]
Chapter 25  Alpha-Beta Pruning

Figure 25.4  Game tree evaluation with *negmax*

\[ \text{negate } b \ (L f \ x) = L f \ (\text{if } b \ \text{then } -x \ \text{else } x) \]
\[ \text{negate } b \ (N d \ ts) = N d \ (\text{map } (\text{negate } (\neg)) \ ts) \]

Now we can express that *negmax* correctly mimics the behaviour of *maxmin* and *minmax*:

\[ \text{maxmin } t = \text{negmax } (\text{negate } \text{False } t) \quad (25.8) \]
\[ \text{minmax } t = -\text{negmax } (\text{negate } \text{True } t) \quad (25.9) \]

The proof is by simultaneous induction on the computations of *maxmin* and *minmax*. We focus on the induction step. By IH the equation holds for all \( t \in \text{set } ts \). The IH will be combined with the following general congruence property for *map*:

\[ (\forall x \in \text{set } xs. \ f x = g x) \rightarrow \text{map } f \ xs = \text{map } g \ xs \quad (25.10) \]

The proof of (25.8) follows:

\[
\begin{align*}
\text{maxmin } (N d \ ts) &= \text{maxs } (\text{map } \text{minmax } ts) \\
&= \text{maxs } (\text{map } (\lambda t. \text{negmax } (\text{negate } \text{True } t)) \ ts) \ \text{by (25.10) and IH} \\
&= \text{maxs } (\text{map } ((\lambda t. \text{negmax } t) \circ \text{negate } \text{True}) \ ts) \\
&= \text{maxs } (\text{map } (\lambda t. \text{negmax } t) (\text{map } (\text{negate } \text{True}) \ ts)) \\
&= \text{negmax } (N d (\text{map } (\text{negate } \text{True}) \ ts)) \\
&= \text{negmax } (\text{negate } \text{False } (N d \ ts))
\end{align*}
\]

The proof of (25.9) is almost dual but also uses a generalization of (25.7) to lists, which follows easily by induction:

\[-\text{mins } (\text{map } f \ xs) = \text{maxs } (\lambda x. -f x) \ xs\]
25.3.2 Alpha-Beta Pruning

Alpha-beta pruning for de Morgan orders is easily derived from the $ab\_max/min$ functions using negation ("\(") and swapping $a$ and $b$ when switching between players:

\[
\begin{align*}
\text{ab\_negmax} &:: 'a \Rightarrow 'a \Rightarrow 'a \text{ tree} \Rightarrow 'a \\
\text{ab\_negmax} _ _ (Lf x) & = x \\
\text{ab\_negmax} a b (Nd ts) & = \text{ab\_negmaxs} a b ts \\
\text{ab\_negmaxs} &:: 'a \Rightarrow 'a \Rightarrow 'a \text{ tree list} \Rightarrow 'a \\
\text{ab\_negmaxs} a _ [] & = a \\
\text{ab\_negmaxs} a b (t \# ts) & = (\text{let } a' = \text{max} a (- \text{ab\_negmax} (- b) (- a) t) \\
& \text{ in if } b \leq a' \text{ then } a' \text{ else } \text{ab\_negmaxs} a' b ts)
\end{align*}
\]

Correctness can be proved easily by simultaneous induction

\[
\begin{align*}
\text{fishburn} a b (\text{negmax } t) (\text{ab\_negmax } a b t) & \text{ if } a \leq b \\
\text{fishburn} a b (\text{negmax } (Nd ts)) (\text{ab\_negmaxs } a b ts) & \text{ if } a \leq b
\end{align*}
\]

using this simple inductive fact: $a \leq \text{ab\_negmaxs } a b ts$.

25.3.3 Exercises

Exercise 25.8. It is straightforward to connect $\text{ab\_negmax}$ and $\text{ab\_max}$

\[
\text{ab\_max } a b t = \text{ab\_negmax } a b (\text{negate False } t)
\]

by simultaneous computation induction involving a further three analogous equations connecting pairs of alpha-beta functions.

Exercises 25.5 and ?? carry over to negative values, \textit{mutatis mutandis}.

25.4 Alpha-Beta Pruning for Distributive Lattices

Although alpha-beta pruning is customarily presented for linear orderings, it also works for the more general domain of distributive lattices. This has applications to games with incomplete information such as many card games because distributive lattices can represent sets of possible situations. For games of complete information such as chess, distributive lattices have applications too. They support heuristic evaluations with multiple components (e.g. material, mobility, etc.) without being forced to combine them into a single value or order them linearly because tuples of numbers form a distributive lattice.
25.4.1 Lattices

A lattice on some type \('a\) is a partial order (\(\leq\)) such that any two elements have a greatest lower and a least upper bound. These two operations are denoted by the following constants and are also called infimum and supremum:

\[
\begin{align*}
(\sqcap) &:: 'a \Rightarrow 'a \Rightarrow 'a \\
(\sqcup) &:: 'a \Rightarrow 'a \Rightarrow 'a
\end{align*}
\]

They fulfill these properties:

\[
\begin{align*}
x \sqcap y \leq x &\quad x \sqcap y \leq y &\quad x \leq y \land x \leq z \rightarrow x \leq y \sqcap z \\
x \leq x \sqcup y &\quad y \leq x \sqcup y &\quad y \leq x \land z \leq x \rightarrow y \sqcup z \leq x
\end{align*}
\]

That is, \(\sqcap\) is the greatest lower and \(\sqcup\) the least upper bound. Note that \(\sqcap\) has a higher precedence than \(\sqcup\): \(x \sqcup y \sqcap z\) means \(x \sqcup (y \sqcap z)\). Just like \(\land/\lor\) and \(\sqcap/\sqcup\).

Any linear order is a lattice where \(\sqcap = \text{min}\) and \(\sqcup = \text{max}\). An example of a lattice that is not a linear order is the type of sets where \(\sqcap = \emptyset\) and \(\sqcup = \bigcup\).

It turns out that \(\sqcap\) and \(\sqcup\) have very nice algebraic properties: both are associative and commutative and enjoy these absorption properties:

\[
\begin{align*}
x \sqcap x &= x &\quad x \sqcap (x \sqcup y) &= x \\
x \sqcup x &= x &\quad x \sqcup x \sqcap y &= x
\end{align*}
\]

A distributive lattice is a lattice where \(\sqcap\) and \(\sqcup\) distribute over each other:

\[
\begin{align*}
x \sqcup y \sqcap z &= (x \sqcup y) \sqcap (x \sqcap z) \\
x \sqcap (y \sqcup z) &= x \sqcap y \sqcup x \sqcap z
\end{align*}
\]

Clearly, linear orders and sets form distributive lattices. Moreover, the Cartesian product of distributive lattices is again a distributive lattice.

In the rest of this section we work in a distributive lattice. Often we also assume that the lattice is bounded, i.e. has a least and a greatest element \(\bot\) and \(\top\). Of course bounded lattices satisfy the obvious properties \(\bot \sqcap x = \bot, \top \sqcap x = x, \bot \sqcup x = x\) and \(\top \sqcup x = \top\).

In the sequel, we rarely enlarge on parts of a proof that follow by distributive lattice laws alone; we take those for granted. For concreteness the reader may think in terms of sets rather than distributive lattices and will not be mislead.

25.4.2 Alpha-Beta Pruning

Both game tree evaluation and alpha-beta pruning are completely analogous to before, except that \(\text{min}\) and \(\text{max}\) are generalized to \(\sqcap\) and \(\sqcup\). The result is shown in Figure 25.5.

We will prove \(ab_{\sup} \sqcap \top \sqcup t = \sup inf t\), but we cannot proceed via the following naive generalization of Theorem 25.1
\begin{verbatim}
supinf :: 'a tree \Rightarrow 'a
supinf (Lf x) = x
supinf (Nd ts) = sups (map infsup ts)

infsup :: 'a tree \Rightarrow 'a
infsup (Lf x) = x
infsup (Nd ts) = infs (map supinf ts)

sups :: 'a list \Rightarrow 'a
sups [] = ⊥
sups (x # xs) = x \sqcap sups xs

infs :: 'a list \Rightarrow 'a
infs [] = ⊤
infs (x # xs) = x \sqcap infs xs

ab_sup :: 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a tree \Rightarrow 'a
ab_sup _ _ (Lf x) = x
ab_sup a b (Nd ts) = ab_sups a b ts

ab_sups :: 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a tree list \Rightarrow 'a
ab_sups a b [] = a
ab_sups a b (t # ts) = (let a' = a \sqcup ab_inf a b t in if b \leq a' then a' else ab_sups a' b ts)

ab_inf :: 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a tree \Rightarrow 'a
ab_inf _ _ (Lf x) = x
ab_inf a b (Nd ts) = ab_infs a b ts

ab_infs :: 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a tree list \Rightarrow 'a
ab_infs _ b [] = b
ab_infs a b (t # ts) = (let b' = b \sqcap ab_sup a b t in if b' \leq a then b' else ab_infs a b' ts)
\end{verbatim}

Figure 25.5  Game tree evaluation and alpha-beta pruning for lattices
Chapter 25  Alpha-Beta Pruning

\[ a < b \rightarrow \text{fishburn } a \ b \ (\supinf \ t) \ (ab\_sup \ a \ b \ t) \]  
(25.11)
because it does not hold.

25.4.2.1 Counterexamples

Property (25.11) does not hold in general as the following counterexample for the distributive lattice bool set shows. Let \( a = \{\text{False}\}, b = \{\text{False, True}\} \) (\( a < b \)) and \( t = \text{Nd} [\text{Lf} \ \{\text{True}\}] \). Then \( \supinf \ t = \{\text{True}\} =: v \) and \( ab\_sup \ a \ b \ t = \{\text{False, True}\} =: ab \) But although \( ab \geq b \), we don’t have \( v \geq b \) as knuth and fishburn would require.

More generally, the definition of \( \text{fishburn } a \ b \ v \ ab \) implicitly assumes that \( ab \) is the result of alpha-beta pruning, satisfies one of the three alternatives \( ab \leq a \), \( a < ab \ < b \) or \( b < ab \). In a distributive lattice this may no longer be the case. Take \( a = \{\}, b = \{\text{True}\} \) and \( t = \text{Nd} [\text{Lf} \ \{\text{False}\}] \). Then \( \supinf \ t = \{\text{False}\} =: v \) and \( ab\_sup \ a \ b \ t = \{\text{True}\} =: ab \). But now all three comparisons \( ab \leq a \), \( a < ab \wedge ab < b \) and \( b < ab \) are false. Thus we cannot draw any conclusion about \( v \) from \( ab \).

In summary, for distributive lattices, \( \text{fishburn} \) is unsuitable for relating the result of alpha-beta pruning to the true tree value.

25.4.3 Correctness and Proof

We will phrase correctness by means of the operation \( a \sqcup x \sqcap b \) that squashes \( x \) into the closed interval \([a,b]\), assuming \( a \leq b \):

\[ a < b \rightarrow a \leq a \sqcup x \sqcap b \leq b \]

If \( a \leq x \leq b \) then \( a \sqcup x \sqcap b = x \). Note also that if \( a \leq b \), then the order of \( \sqcup \) and \( \sqcap \) is irrelevant: \( a \leq b \rightarrow a \sqcup x \sqcap b = (a \sqcup x) \sqcap b \).

Although \( a \sqcup x \sqcap b \) has particularly nice properties if \( a \leq b \), it can be manipulated algebraically even in the absence of \( a \leq b \). As an example we have this weak form of the preceding associativity property:

\[ a \sqcup x \sqcap b = a \sqcup y \sqcap b \iff (a \sqcup x) \sqcap b = (a \sqcup y) \sqcap b \]  
(25.12)

Let \( v \) be the value of tree \( t \) and let \( ab \) be the result of alpha-beta pruning of \( t \). We can express the correctness of \( ab \) w.r.t. \( v \) as saying that they are the same modulo “squashing”: \( a \sqcup ab \sqcap b = a \sqcup v \sqcap b \). Correctness can be shown simultaneously for all four functions:

**Theorem 25.7.**

\[ (a \sqcup ab\_sup \ a \ b \ t) \sqcap b = (a \sqcup \supinf \ t) \sqcap b \]
\[ (a \sqcup ab\_sup \ a \ b \ t \ s) \sqcap b = (a \sqcup \supinf \ (\text{Nd} \ ts)) \sqcap b \]
\[ a \sqcup ab\_inf \ a \ b \ t \sqcap b = a \sqcup \infsup \ t \sqcap b \]
\[ a \sqcup ab\_inf \ a \ b \ t \ s \sqcap b = a \sqcup \infsup \ (\text{Nd} \ ts) \sqcap b \]
Proof by simultaneous computation induction. The only two nontrivial cases are the ones stemming from the recursion equations for \(ab_{sups}\) and \(ab_{infs}\). We concentrate on \(ab_{sups}\). For succinctness we introduce the following abbreviations:

\[
\begin{align*}
abt & \equiv ab_{inf} a b t \\
vts & \equiv supinf (Nd ts)
\end{align*}
\]

The two IHs are

\[
\begin{align*}
& a \uplus abt \cap b = a \uplus vt \cap b \tag{IH1} \\
\neg b \leq a \uplus abt \rightarrow (a \uplus abt \uplus abts) \cap b = (a \uplus abt \uplus vts) \cap b \tag{IH2}
\end{align*}
\]

and we need to prove

\[
(a \uplus ab_{sups} a b (t \# ts)) \cap b = (a \uplus supinf (Nd (t \# ts))) \cap b.
\]

The proof is by cases.

First we assume \(b \leq a \uplus abt\). Using (25.12) we can transform IH1 into

\[
(a \uplus abt) \cap b = (a \uplus vt) \cap b \tag{IH1'}
\]

With \(b \leq a \uplus abt\) this implies \(b = (a \uplus vt) \cap b\) (*). Now we prove the main equation:

\[
\begin{align*}
(a \uplus ab_{sups} a b (t \# ts)) \cap b & = (a \uplus (a \uplus abt)) \cap b \text{ because } b \leq a \uplus abt \\
& = (a \uplus abt) \cap b \\
& = (a \uplus vt) \cap b \quad \text{by IH1'} \\
& = (a \uplus vt \uplus vts) \cap (a \uplus vt) \cap b \\
& = (a \uplus vt \uplus vts) \cap b \quad \text{by (*)} \\
& = (a \uplus supinf (Nd (t \# ts))) \cap b
\end{align*}
\]

Now we assume \(\neg b \leq a \uplus abt\). In this case we need the following simple inductive property of \(ab_{sups}\): \(x \leq ab_{sups} x y ts\). With the help of this property and \(\neg b \leq a \uplus abt\), IH2 yields

\[
(a \uplus abts) \cap b = (a \uplus abt \uplus vts) \cap b \tag{IH2'}
\]

Now we prove the main equation:

\[
\begin{align*}
(a \uplus ab_{sups} a b (t \# ts)) \cap b & = (a \uplus abts) \cap b \text{ because } \neg b \leq a \uplus abt \\
& = (a \uplus abt \uplus vts) \cap b \quad \text{by IH2'} \\
& = a \cap b \uplus abt \cap b \uplus vts \cap b \\
& = (a \uplus abt \cap b) \cap b \uplus vts \cap b \\
& = (a \uplus vt \cap b) \cap b \uplus vts \cap b \quad \text{by IH1} \\
& = (a \uplus vt \uplus vts) \cap b \\
& = (a \uplus supinf (Nd (t \# ts))) \cap b \quad \square
\end{align*}
\]

Corollary 25.8. \(ab_{sup} \perp \top t = supinf t\) (25.13)
25.4.4 Negative Values

We can deal with negative values in the context of bounded distributive lattices by requiring the same properties of as for de Morgan orders, but with \((\cap)\) instead of \(\min\):

\[
- (x \cap y) = - x \cup - y \\
- (- x) = x
\]

The resulting structure is called a de Morgan algebra. Just as in Section 25.3 we can define game tree evaluation

\[
\text{negsup} :: 'a \Rightarrow 'a \\
\text{negsup} (Lf x) = x \\
\text{negsup} (Nd ts) = \text{sup} (\map (\lambda t. - \text{negsup} t) ts)
\]

and alpha-beta pruning for de Morgan algebras:

\[
\text{ab_negsup} :: 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a \\
\text{ab_negsup} _ _ (Lf x) = x \\
\text{ab_negsup} a b (Nd ts) = \text{ab_negsups} a b ts
\]

\[
\text{ab_negsups} :: 'a \Rightarrow 'a \Rightarrow 'a \Rightarrow 'a \\
\text{ab_negsups} a [] = a \\
\text{ab_negsups} a b (t \# ts) = (\let a' = a \cup - \text{ab_negsup} (- b) (- a) t \\
\text{in if } b \leq a' \text{ then } a' \text{ else } a' \text{ ab_negsups } a' \text{ b ts})
\]

We can also relate the ordinary and the negated versions by simultaneous computation induction

\[
\text{ab_sup} a b t = \text{ab_negsup} a b (\text{negate False} t) \\
\text{ab_sup} a b ts = \text{ab_negsup} a b (\map (\text{negate True}) ts) \\
\text{ab_inf} a b t = - \text{ab_negsup} (- b) (- a) (\text{negate True} t) \\
\text{ab_inf} a b ts = - \text{ab_negsups} (- b) (- a) (\map (\text{negate False}) ts)
\]

and conclude

\[
\text{ab_negsup} \perp \top t = \text{supinf} (\text{negate False} t)
\]

with the help of \((25.13)\) (and the inductive lemma \(\text{negate f} (\text{negate f} t) = t\)).
25.4.5 Exercises

Exercise 25.9. In Exercise 25.2 we considered a reformulation of fishburn. This reformulation (but not the equivalence!) generalizes to lattices in the standard manner:

\[
\text{bounded } a \ b \ v \ ab \equiv b \land v \leq ab \land ab \leq a \lor v
\]

It turns out that this is a suitable correctness notion for alpha-beta pruning in distributive lattices. Give a detailed proof of this generalization of Theorem 25.7:

\[
\text{bounded } a \ b \ (\supinf t) \ (\text{ab}_\text{sup} \ a \ b \ t)
\]

Obviously \(\text{ab}_\text{sup} \top \top t = \supinf t\) follows immediately.

Give a detailed proof of \(\text{bounded } a \ b \ v \ ab \not\equiv a \top v \land b\) and a counterexample to the reverse implication.

Exercise 25.10. There is also a fail-soft version of alpha-beta pruning for distributive lattices:

\[
\text{ab}_\text{sup}' :: \ 'a \Rightarrow \ 'a \Rightarrow \ 'a \Rightarrow \ 'a
\]

\[
\text{ab}_\text{sup}': \ (Lf x) = x
\]

\[
\text{ab}_\text{sup}' a \ b \ (Nd \ ts) = \text{ab}_\text{sup}' a \ b \ ⊥ \ ts
\]

\[
\text{ab}_\text{sups}' :: \ 'a \Rightarrow \ 'a \Rightarrow \ 'a \Rightarrow \ 'a \Rightarrow \ 'a \Rightarrow \ 'a
\]

\[
\text{ab}_\text{sups}' a \ b \ m \ (t \# ts)
\]

\[
= (\text{let } m' = m \land \text{ab}_\text{inf}' (m \lor a) \ b \ t
\]

\[
\text{in } \text{if } b \leq m' \text{ then } m' \text{ else } \text{ab}_\text{sups}' a \ b \ m' \ ts)
\]

Prove its correctness (for \(\text{bounded}\) see Exercise 25.9):

\[
\text{bounded } a \ b \ (\supinf t) \ (\text{ab}_\text{sup}' a \ b \ t)
\]

\[
\text{bounded } (a \land m) \ b \ (\supinf (Nd \ ts)) \ (\text{ab}_\text{sups}' a \ b \ m \ ts)
\]

Exercise 25.11. Based on the definition of \(\text{bounded}\) in Exercise 25.9, prove

\[
\text{bounded } a \ b \ (\text{negsup } t) \ (\text{ab}_\text{negsup} a \ b \ t)
\]

\[
\text{bounded } a \ b \ (\text{negsup } (Nd \ ts)) \ (\text{ab}_\text{negsups} a \ b \ ts)
\]

directly, i.e. without going back to the non-negative relatives. You may need the lemma \(a \leq \text{ab}_\text{negsups} a \ b \ ts\).

Exercise 25.12. The algorithm considered in Exercises 25.5 carries over to distributive lattices, mutatis mutandis. Prove

\[
a \leq b \rightarrow \text{ab}_\text{sup}2 a \ b \ t = a \lor \supinf t \land b
\]

Obviously \(\text{ab}_\text{sup}2 \bot \top t = \supinf t\) follows immediately.
Chapter Notes

Variants of alpha-beta pruning have a long history in the literature about computer chess. It appears that the first reasonably precise correctness proof was given by Knuth and Moore [1975]. The improvement from fail-hard to fail-soft was proposed by Fishburn [1983] with the suggestion of using it to narrow the $a, b$ window in future searches of the same position. Marsland [1986] spells out the details of the code. Surprisingly, Fishburn [1983] only attributes the weak bounding property \textit{knuth} to the fail-hard variant and the bounding property \textit{fishburn} to the fail-soft variant. Although the former is true, he does not seem to have realized that even the fail-hard variant satisfies \textit{fishburn} and that the distinguishing property is that fail-hard bounds fail-soft (Theorem 25.3).

Hughes [1989] derives a version of alpha-beta pruning for numbers from the definition of maxmin. However, he ends up with shallow pruning only, i.e. function $F_1$ by Knuth and Moore [1975], not $F_2$, the real alpha-beta pruning. In their historic survey, Knuth and Moore [1975, pp.303-304] point out that this mistake has been made frequently, including by Knuth himself.

The fact that alpha-beta pruning extends to distributed lattices was discovered twice. First by Bird and Hughes [1987], who (like Hughes [1989]) derive an algorithm from the definition of maxmin. Confusingly they talk about Boolean algebras although they merely work in a distributive lattice. Their version of alpha-beta pruning could be classified as fail-extremely-hard because it always returns a result in the interval $[a, b]$ (see Exercise 25.12). Ginsberg and Jaffray [2002] rediscovered that alpha-beta pruning also works in distributed lattices. Li et al. [2022] extend alpha-beta pruning in distributive lattices to fail-soft on a game graph using a cache. They employ the squashing operation $a \sqcup x \sqcap b$ introduced by Bird and Hughes [1987] to state correctness. Both Ginsberg and Jaffray [2002] and Li et al. [2022] are unaware of the work by Bird and Hughes [1987] who in turn are unaware of the work by Knuth and Moore [1975].

De Morgan algebras were introduced and studied by Moisil [1936, p. 91] (without the assumption of boundedness). The term “de Morgan order” is not standard and was coined by the author in analogy with de Morgan algebras. Pearl [1980, 1982] provided the definitive quantitative analysis of alpha-beta pruning and showed that, for random game trees, alpha-beta pruning is optimal.
Part VI

Appendix
List Library

The following functions on lists are predefined:

\[
\text{length} :: \text{`a list} \Rightarrow \text{nat} \\
|[]| = 0 \\
|\text{x} \# \text{xs}| = |\text{xs}| + 1
\]

\[
(\@) :: \text{`a list} \Rightarrow \text{`a list} \Rightarrow \text{`a list} \\
[] @ \text{ys} = \text{ys} \\
(\text{x} \# \text{xs}) @ \text{ys} = \text{x} \# \text{xs} @ \text{ys}
\]

\[
\text{set} :: \text{`a list} \Rightarrow \text{`a set} \\
\text{set} [] = \{\} \\
\text{set} (\text{x} \# \text{xs}) = \{\text{x}\} \cup \text{set} \text{xs}
\]

\[
\text{map} :: (\text{`a} \Rightarrow \text{`b}) \Rightarrow \text{`a list} \Rightarrow \text{`b list} \\
\text{map} \text{f} [] = [] \\
\text{map} \text{f} (\text{x} \# \text{xs}) = \text{f} \text{x} \# \text{map} \text{f} \text{xs}
\]

\[
\text{filter} :: (\text{`a} \Rightarrow \text{bool}) \Rightarrow \text{`a list} \Rightarrow \text{`a list} \\
\text{filter} \text{p} [] = [] \\
\text{filter} \text{p} (\text{x} \# \text{xs}) = \text{(if } \text{p} \text{x} \text{then } \text{x} \# \text{filter} \text{p} \text{xs} \text{else } \text{filter} \text{p} \text{xs})
\]

\[
\text{concat} :: \text{`a list list} \Rightarrow \text{`a list} \\
\text{concat} [] = [] \\
\text{concat} (\text{x} \# \text{xs}) = \text{x} @ \text{concat} \text{xs}
\]

\[
\text{take} :: \text{nat} \Rightarrow \text{`a list} \Rightarrow \text{`a list} \\
\text{take} _ [] = [] \\
\text{take} \text{n} (\text{x} \# \text{xs}) = \text{(case } \text{n} \text{of } 0 \Rightarrow [] | m + 1 \Rightarrow \text{x} \# \text{take} \text{m} \text{xs})
\]

\[
\text{drop} :: \text{nat} \Rightarrow \text{`a list} \Rightarrow \text{`a list}
\]
\[
\begin{align*}
\text{drop} _ [] &= [] \\
\text{drop} n (x \# xs) &= (\text{case } n \text{ of } 0 \Rightarrow x \# xs | m + 1 \Rightarrow \text{drop } m xs)
\end{align*}
\]

\[
hd :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\textquotesingle}a
\]

\[
hd (x \# xs) = x
\]

\[
tl :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
tl [] = [] \\
\text{\texttt{\textquotesingle}a list }\]

\[
tl (x \# xs) = xs
\]

\[
\text{butlast} :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
\text{butlast} [] = [] \\
\text{\texttt{\textquotesingle}a list }\]

\[
\text{butlast} (x \# xs) = (\text{if } xs = [] \text{ then } [] \text{ else } x \# \text{butlast } xs)
\]

\[
\text{rev} :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
\text{rev} [] = [] \\
\text{\texttt{\textquotesingle}a list }\]

\[
\text{rev} (x \# xs) = \text{rev } xs \circ [x]
\]

\[
(\text{!}) :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
x \# xs) ! n = (\text{case } n \text{ of } 0 \Rightarrow x | k + 1 \Rightarrow xs ! k)
\]

\[
\text{list_update} :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
[[\_ := \_]] = [] \\
(x \# xs)[i := v] = (\text{case } i \text{ of } 0 \Rightarrow v \# xs | j + 1 \Rightarrow x \# xs[j := v])
\]

\[
\text{upt} :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
[\_..<0] = [] \\
[i..<j + 1] = (\text{if } i \leq j \text{ then } [i..<j] \circ [j] \text{ else } [])
\]

\[
\text{replicate} :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
\text{replicate } 0 \_ = [] \\
\text{\texttt{\textquotesingle}a list }\]

\[
\text{replicate} (n + 1) x = x \# \text{replicate } n x
\]

\[
\text{sum_list} :: \text{\texttt{\textquotesingle}a list } \Rightarrow \text{\texttt{\textquotesingle}a list}
\]

\[
\text{sum_list} [] = 0
\]

\[
\text{sum_list} [x] = x
\]
\[
\text{sum\_list}\ (x \#\ xs) = x + \text{sum\_list}\ xs
\]

\[
\text{min\_list} :: \ 'a\ list \Rightarrow\ 'a
\]
\[
\text{min\_list}\ (x \#\ xs) = (\text{case}\ xs\ \text{of}\ [] \Rightarrow x | _ \# \_ \Rightarrow \text{min}\ x\ (\text{min\_list}\ xs))
\]

\[
\text{sorted\_wrt} :: (\ 'a \Rightarrow\ 'a \Rightarrow\ \text{bool}) \Rightarrow\ 'a\ list \Rightarrow\ \text{bool}
\]
\[
\text{sorted\_wrt}\ P\ [] = \text{True}
\]
\[
\text{sorted\_wrt}\ P\ (x \#\ ys) = ((\forall y \in\ \text{set}\ ys.\ P\ x\ y) \land\ \text{sorted\_wrt}\ P\ ys)
\]
Time Functions

Time functions that are 0 by definition have already been simplified away.

B.1 Lists

\[ T_{\text{append}} :: 'a \text{ list} \Rightarrow 'a \text{ list} \Rightarrow \text{nat} \]
\[ T_{\text{append}} [\_ , \_] = 1 \]
\[ T_{\text{append}} (\_ \# xs) \ ys = T_{\text{append}} \ xs \ ys + 1 \]

\[ T_{\text{length}} :: 'a \text{ list} \Rightarrow \text{nat} \]
\[ T_{\text{length}} [\_] = 1 \]
\[ T_{\text{length}} (\_ \# xs) = T_{\text{length}} \ xs + 1 \]

\[ T_{\text{map}} :: ('a \Rightarrow \text{nat}) \Rightarrow 'a \text{ list} \Rightarrow \text{nat} \]
\[ T_{\text{map}} [\_] = 1 \]
\[ T_{\text{map}} \ Tf (x \# xs) = Tf \ x + T_{\text{map}} \ Tf \ xs + 1 \]

\[ T_{\text{filter}} :: ('a \Rightarrow \text{nat}) \Rightarrow 'a \text{ list} \Rightarrow \text{nat} \]
\[ T_{\text{filter}} [\_] = 1 \]
\[ T_{\text{filter}} \Tp (x \# xs) = Tp \ x + T_{\text{filter}} \ Tp \ xs + 1 \]

\[ T_{\text{take}} :: \text{nat} \Rightarrow 'a \text{ list} \Rightarrow \text{nat} \]
\[ T_{\text{take}} [\_] = 1 \]
\[ T_{\text{take}} n (\_ \# xs) = (\text{case } n \text{ of } 0 \Rightarrow 0 \mid m + 1 \Rightarrow T_{\text{take}} m \ xs) + 1 \]

\[ T_{\text{drop}} :: \text{nat} \Rightarrow 'a \text{ list} \Rightarrow \text{nat} \]
\[ T_{\text{drop}} [\_] = 1 \]
\[ T_{\text{drop}} n (\_ \# xs) = (\text{case } n \text{ of } 0 \Rightarrow 0 \mid m + 1 \Rightarrow T_{\text{drop}} m \ xs) + 1 \]

\[ T_{\text{nth}} :: 'a \text{ list} \Rightarrow \text{nat} \Rightarrow \text{nat} \]
\[ T_{\text{nth}} (\_ \# xs) n = (\text{case } n \text{ of } 0 \Rightarrow 0 \mid x + 1 \Rightarrow T_{\text{nth}} \ xs \ x) + 1 \]
Simple properties:

\[ T_{\text{append}} \ x s \ y s = |x s| + 1 \]
\[ T_{\text{length}} \ x s = |x s| + 1 \]
\[ T_{\text{map}} \ T_f \ x s = \left( \sum_{x \in x s} T_f \ x \right) + |x s| + 1 \]
\[ T_{\text{filter}} \ T_p \ x s = \left( \sum_{x \in x s} T_p \ x \right) + |x s| + 1 \]
\[ T_{\text{take}} \ n \ x s = \text{min} \ n \ |x s| + 1 \]
\[ T_{\text{drop}} \ n \ x s = \text{min} \ n \ |x s| + 1 \]
\[ n < |x s| \quad \rightarrow \quad T_{\text{nth}} \ x s \ n = n + 1 \]

### B.2 Selection

**`T_{\text{chop}} :: \text{nat} \Rightarrow 'a \ list \Rightarrow \text{nat}`**

\[ T_{\text{chop}} \ 0 \ _ = 1 \]
\[ T_{\text{chop}} \ _ \ [] = 1 \]
\[ T_{\text{chop}} \ n \ x s = T_{\text{take}} \ n \ x s + T_{\text{drop}} \ n \ x s + T_{\text{chop}} \ n \ (\text{drop} \ n \ x s) + 1 \]

**`T_{\text{partition3}} :: 'a \Rightarrow 'a \ list \Rightarrow \text{nat}`**

\[ T_{\text{partition3}} \ _ \ [] = 1 \]
\[ T_{\text{partition3}} \ x \ (_ \ \# \ y s) = T_{\text{partition3}} \ x \ y s + 1 \]

**`T_{\text{slow\_select}} :: \text{nat} \Rightarrow 'a \ list \Rightarrow \text{nat}`**

\[ T_{\text{slow\_select}} \ k \ x s = T_{\text{insort}} \ x s + T_{\text{nth}} \ (\text{insort} \ x s) \ k \]

**`T_{\text{slow\_median}} :: 'a \ list \Rightarrow \text{nat}`**

\[ T_{\text{slow\_median}} \ x s = T_{\text{length}} \ x s + T_{\text{slow\_select}} \ ((|x s| - 1) \ \text{div} \ 2) \ x s \]

Simple properties:

\[ T_{\text{chop}} \ d \ x s \leq 5 \cdot |x s| + 1 \]
\[ T_{\text{partition3}} \ x \ x s = |x s| + 1 \]
\[ k < |x s| \quad \rightarrow \quad T_{\text{slow\_select}} \ k \ x s \leq |x s|^2 + 3 \cdot |x s| + 1 \]
\[ x s \neq [] \quad \rightarrow \quad T_{\text{slow\_median}} \ x s \leq |x s|^2 + 4 \cdot |x s| + 2 \]

### B.3 2-3 Trees
### B.4 Arrays via Braun Trees

**T_{\text{join\_adj}}** :: 'a tree23s \(\Rightarrow\) nat

\[
T_{\text{join\_adj}}(TTs \_ \_ (T \_)) = 1
\]
\[
T_{\text{join\_adj}}(TTs \_ \_ (TTs \_ \_ (T \_))) = 1
\]
\[
T_{\text{join\_adj}}(TTs \_ \_ (TTs \_ \_ ts)) = T_{\text{join\_adj}} ts + 1
\]

**T_{\text{join\_all}}** :: 'a tree23s \(\Rightarrow\) nat

\[
T_{\text{join\_all}}(T \_ \_ \_ (T \_)) = 1
\]
\[
T_{\text{join\_all}} ts = T_{\text{join\_adj}} ts + T_{\text{join\_all}}(\text{join\_adj} ts) + 1
\]

**T_{\text{leaves}}** :: 'a list \(\Rightarrow\) nat

\[
T_{\text{leaves}}[] = 1
\]
\[
T_{\text{leaves}}(\_ \# as) = T_{\text{leaves}} as + 1
\]

**T_{\text{tree23\_of\_list}}** :: 'a list \(\Rightarrow\) nat

\[
T_{\text{tree23\_of\_list}} as = T_{\text{leaves}} as + T_{\text{join\_all}}(\text{leaves as})
\]

### B.5 Leftist Heaps

**T_{\text{nodes}}** :: 'a tree list \(\Rightarrow\) 'a list \(\Rightarrow\) 'a tree list \(\Rightarrow\) nat

\[
T_{\text{nodes}}(\_ \# ls) (\_ \# xs) (\_ \# rs) = T_{\text{nodes}} ls xs rs + 1
\]
\[
T_{\text{nodes}}(\_ \# ls) (\_ \# xs) [] = T_{\text{nodes}} ls xs [] + 1
\]
\[
T_{\text{nodes}}[] (\_ \# xs) (\_ \# rs) = T_{\text{nodes}} [] xs rs + 1
\]
\[
T_{\text{nodes}}[] (\_ \# xs) [] = T_{\text{nodes}} [] xs [] + 1
\]
\[
T_{\text{nodes}} [] [] = 1
\]

**T_{\text{merge}}** :: ('a \times nat) tree \(\Rightarrow\) ('a \times nat) tree \(\Rightarrow\) nat

\[
T_{\text{merge}}() = 1
\]
\[
T_{\text{merge}}() = 1
\]
\[
T_{\text{merge}}((l_1, (a_1, n_1), r_1)) = T_{\text{merge}} ((l_2, (a_2, n_2), r_2)) = t_1 (if a_1 \leq a_2 then T_{\text{merge}} r_1 t_2 else T_{\text{merge}} t_1 r_2) + 1
\]

**T_{\text{insert}}** :: 'a \(\Rightarrow\) ('a \times nat) tree \(\Rightarrow\) nat
Appendix B  Time Functions

\[ T_{\text{insert}} \ x \ t = T_{\text{merge}} \ (\langle \rangle, (x, 1), \langle \rangle) \ t \]

\[ T_{\text{del\_min}} :: ('a \times \text{nat}) \ \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{del\_min}} \ (\langle \rangle) = 0 \]
\[ T_{\text{del\_min}} \ (l, _ _, r) = T_{\text{merge}} \ l \ r \]

\[ T_{\text{merge\_all}} :: ('a \times \text{nat}) \ \text{tree} \ \text{list} \Rightarrow \text{nat} \]
\[ T_{\text{merge\_all}} \ [] = 0 \]
\[ T_{\text{merge\_all}} \ [\_ ] = 0 \]
\[ T_{\text{merge\_all}} \ ts = T_{\text{merge\_all}} \ (\text{merge\_adj} \ ts) + T_{\text{merge\_adj}} \ ts \]

B.6 Priority Queues Based on Braun Trees

\[ T_{\text{insert}} :: 'a \Rightarrow 'a \ \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{insert}} \ (\langle \rangle) = 1 \]
\[ T_{\text{insert}} \ a \ (l, x, r) = (\text{if} \ a < x \ \text{then} \ T_{\text{insert}} \ x \ r \ \text{else} \ T_{\text{insert}} \ a \ r) + 1 \]

\[ T_{\text{del\_min}} :: 'a \ \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{del\_min}} \ (\langle \rangle) = 0 \]
\[ T_{\text{del\_min}} \ (l, _ _, _ ) = 0 \]
\[ T_{\text{del\_min}} \ (l, _ _, r) = T_{\text{del\_left}} \ l + (\text{let} \ (y, l') = \text{del\_left} \ l \ \text{in} \ T_{\text{sift\_down}} \ r \ y \ l') \]

\[ T_{\text{del\_left}} :: 'a \ \text{tree} \Rightarrow 'a \ \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{del\_left}} \ (\langle \rangle, _ _, _ ) = 1 \]
\[ T_{\text{del\_left}} \ (l, _ _, _ ) = T_{\text{del\_left}} \ l + 1 \]

\[ T_{\text{sift\_down}} :: 'a \ \text{tree} \Rightarrow 'a \ \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{sift\_down}} \ (\langle \rangle, _ _, _ ) = 1 \]
\[ T_{\text{sift\_down}} \ (l, x_1, r_1) \ a \ (l_2, x_2, r_2) = (\text{if} \ a \leq x_1 \ \& \ a \leq x_2 \ \text{then} \ 0 \ \text{else if} \ x_1 \leq x_2 \ \text{then} \ T_{\text{sift\_down}} \ l_1 \ a \ r_1 \ \text{else} \ T_{\text{sift\_down}} \ l_2 \ a \ r_2) + 1 \]
B.7 Binomial Heaps

The functions \( T_{\text{link}}, T_{\text{rank}}, T_{\text{root}} \) and \( T_{\text{min}} \) are 0 everywhere and have been eliminated from the following definitions.

\[
T_{\text{ins\_tree}} :: 'a\ \text{tree} \Rightarrow 'a\ \text{tree\ list} \Rightarrow \text{nat}
\]
\[
T_{\text{ins\_tree}}\ [] = 1
\]
\[
T_{\text{ins\_tree}}\ t_1\ (t_2\ \#\ ts) = (\text{if } \text{rank}\ t_1 < \text{rank}\ t_2 \text{ then } 0 \text{ else } T_{\text{ins\_tree}}\ (\text{link}\ t_1\ t_2)\ ts) + 1
\]

\[
T_{\text{insert}} :: 'a \Rightarrow 'a\ \text{tree\ list} \Rightarrow \text{nat}
\]
\[
T_{\text{insert}}\ x\ ts = T_{\text{ins\_tree}}\ (\text{Node}\ 0\ x\ [])\ ts
\]

\[
T_{\text{merge}} :: 'a\ \text{tree\ list} \Rightarrow 'a\ \text{tree\ list} \Rightarrow \text{nat}
\]
\[
T_{\text{merge}}\ [] = 1
\]
\[
T_{\text{merge}}\ (t_1\ \#\ ts_1 =: h_1)\ (t_2\ \#\ ts_2 =: h_2) = (\text{if } \text{rank}\ t_1 < \text{rank}\ t_2 \text{ then } T_{\text{merge}}\ ts_1\ h_2
\]
\[\text{else if } \text{rank}\ t_2 < \text{rank}\ t_1 \text{ then } T_{\text{merge}}\ h_1\ ts_2
\]
\[\text{else } T_{\text{merge}}\ ts_1\ ts_2 + T_{\text{ins\_tree}}\ (\text{link}\ t_1\ t_2)\ (\text{merge}\ ts_1\ ts_2) + 1
\]

\[
T_{\text{get\_min}} :: 'a\ \text{tree\ list} \Rightarrow \text{nat}
\]
\[
T_{\text{get\_min}}\ [] = 1
\]
\[
T_{\text{get\_min}}\ (_\ \#\ ts) = T_{\text{get\_min}}\ ts + 1
\]

\[
T_{\text{get\_min\_rest}} :: 'a\ \text{tree\ list} \Rightarrow \text{nat}
\]
\[
T_{\text{get\_min\_rest}}\ [] = 1
\]
\[
T_{\text{get\_min\_rest}}\ (_\ \#\ ts) = T_{\text{get\_min\_rest}}\ ts + 1
\]

\[
T_{\text{rev}} :: 'a\ \text{list} \Rightarrow \text{nat}
\]
\[
T_{\text{rev}}\ xs = |xs| + 1
\]

\[
T_{\text{del\_min}} :: 'a\ \text{tree\ list} \Rightarrow \text{nat}
\]
\[
T_{\text{del\_min}}\ ts = T_{\text{get\_min\_rest}}\ ts +
\]
\[\text{(case } \text{get\_min\_rest}\ ts \text{ of}
\]
\[\text{(Node} _\ _\ ts_1,\ ts_2) \Rightarrow T_{\text{rev}}\ ts_1 + T_{\text{merge}}\ (\text{rev}\ ts_1)\ ts_2)
\]
### B.8 Queues

\[ T_{\text{norm}} : 'a \, \text{list} \times 'a \, \text{list} \Rightarrow \text{nat} \]
\[ T_{\text{norm}} (fs, rs) = (\text{if} \, fs = [] \, \text{then} \, T_{\text{strev}} \, rs \, [] \, \text{else} \, 0) \]

\[ T_{\text{enq}} : 'a \Rightarrow 'a \, \text{list} \times 'a \, \text{list} \Rightarrow \text{nat} \]
\[ T_{\text{enq}} \, a \, (fs, \, rs) = T_{\text{norm}} \, (fs, \, a \, \# \, rs) \]

\[ T_{\text{deq}} : 'a \, \text{list} \times 'a \, \text{list} \Rightarrow \text{nat} \]
\[ T_{\text{deq}} \, (fs, \, rs) = (\text{if} \, fs = [] \, \text{then} \, 0 \, \text{else} \, T_{\text{norm}} \, (\text{tl} \, fs, \, rs)) \]

### B.9 Splay Trees

\[ T_{\text{splay}} : 'a \Rightarrow 'a \, \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{splay}} \, \langle \rangle = 1 \]
\[ T_{\text{splay}} \, x \, \langle AB, \, b, \, CD \rangle\]
\[ = (\text{case} \, \text{cmp} \, x \, b \, \text{of} \]
\[ \text{LT} \Rightarrow \text{case} \, AB \, \text{of} \]
\[ \langle \rangle \Rightarrow 0 \mid \]
\[ \langle A, \, a, \, B \rangle \Rightarrow (\text{case} \, \text{cmp} \, x \, a \, \text{of} \]
\[ \text{LT} \Rightarrow \text{if} \, A = \langle \rangle \, \text{then} \, 0 \, \text{else} \, T_{\text{splay}} \, x \, A \mid \]
\[ \text{EQ} \Rightarrow 0 \mid \]
\[ \text{GT} \Rightarrow \text{if} \, B = \langle \rangle \, \text{then} \, 0 \, \text{else} \, T_{\text{splay}} \, x \, B \mid \]
\[ \text{EQ} \Rightarrow 0 \mid \]
\[ \text{GT} \Rightarrow \text{case} \, CD \, \text{of} \]
\[ \langle \rangle \Rightarrow 0 \mid \]
\[ \langle C, \, c, \, D \rangle \Rightarrow (\text{case} \, \text{cmp} \, x \, c \, \text{of} \]
\[ \text{LT} \Rightarrow \text{if} \, C = \langle \rangle \, \text{then} \, 0 \, \text{else} \, T_{\text{splay}} \, x \, C \mid \]
\[ \text{EQ} \Rightarrow 0 \mid \]
\[ \text{GT} \Rightarrow \text{if} \, D = \langle \rangle \, \text{then} \, 0 \, \text{else} \, T_{\text{splay}} \, x \, D \rangle + 1 \]

\[ T_{\text{splay\_max}} : 'a \, \text{tree} \Rightarrow \text{nat} \]
\[ T_{\text{splay\_max}} \, \langle \rangle = 1 \]
\[ T_{\text{splay\_max}} \, \langle \_, \_, \langle \rangle \rangle = 1 \]
\[ T_{\text{splay\_max}} \, \langle \_, \_, \langle \_, \_, \_, CD \rangle \rangle \]
\[ T = (\text{if } CD = \langle \rangle \text{ then } 0 \text{ else } T_{\text{splay max}} CD + (\text{case } splay_{\text{max}} CD \text{ of } \langle \_, \_, \_ \rangle \Rightarrow 0)) + 1 \]

\[ T_{\text{insert}} :: 'a \Rightarrow 'a \text{ tree } \Rightarrow \text{nat} \]
\[ T_{\text{insert}} x t = (\text{if } t = \langle \rangle \text{ then } 0 \text{ else } T_{\text{splay}} x t) \]

\[ T_{\text{delete}} :: 'a \Rightarrow 'a \text{ tree } \Rightarrow \text{nat} \]
\[ T_{\text{delete}} x t = (\text{if } t = \langle \rangle \text{ then } 0 \text{ else } T_{\text{splay}} x t + (\text{case } splay x t \text{ of } \langle l, a, \_ \rangle \Rightarrow \text{if } x \neq a \text{ then } 0 \text{ else if } l = \langle \rangle \text{ then } 0 \text{ else } T_{\text{splay max}} l + (\text{case } splay_{\text{max}} l \text{ of } \langle \_, \_, \_ \rangle \Rightarrow 0))) \]

B.10 Skew Heaps

\[ T_{\text{merge}} :: 'a \text{ tree } \Rightarrow 'a \text{ tree } \Rightarrow \text{nat} \]
\[ T_{\text{merge}} \langle \rangle \_ = 1 \]
\[ T_{\text{merge}} \_ \langle \rangle = 1 \]
\[ T_{\text{merge}} \langle l_1, a_1, r_1 \rangle \langle l_2, a_2, r_2 \rangle = (\text{if } a_1 \leq a_2 \text{ then } T_{\text{merge}} \langle l_2, a_2, r_2 \rangle r_1 \text{ else } T_{\text{merge}} \langle l_1, a_1, r_1 \rangle r_2) + 1 \]

\[ T_{\text{insert}} :: 'a \Rightarrow 'a \text{ tree } \Rightarrow \text{int} \]
\[ T_{\text{insert}} a t = T_{\text{merge}} \langle \langle \rangle, a, \langle \rangle \rangle t \]

\[ T_{\text{del min}} :: 'a \text{ tree } \Rightarrow \text{int} \]
\[ T_{\text{del min}} t = (\text{case } t \text{ of } \langle \rangle \Rightarrow 0 \text{ | } \langle t_1, \_, t_2 \rangle \Rightarrow T_{\text{merge}} t_1 t_2) \]

B.11 Pairing Heaps

\[ T_{\text{insert}} :: 'a \Rightarrow 'a \text{ tree } \Rightarrow \text{nat} \]
\( T_{\text{insert}} ~_~ = 1 \)

\( T_{\text{merge}} : 'a \text{ tree} \Rightarrow 'a \text{ tree} \Rightarrow \text{nat} \)
\( T_{\text{merge}} ~_~ = 1 \)

\( T_{\text{del_min}} : 'a \text{ tree} \Rightarrow \text{nat} \)
\( T_{\text{del_min}} \langle \rangle = 1 \)
\( T_{\text{del_min}} \langle \text{hs}, ~_~, ~_~ \rangle = T_{\text{pass2}}(\text{pass}_1 \hs) + T_{\text{pass1}} \hs + 1 \)

\( T_{\text{pass1}} : 'a \text{ tree} \Rightarrow \text{nat} \)
\( T_{\text{pass1}} \langle \_, \_, \langle \_, \_, \hs' \rangle \rangle = T_{\text{pass1}} \hs' + 1 \)
\( T_{\text{pass1}} \langle \rangle = 1 \)
\( T_{\text{pass1}} \langle \_, \_, \rangle \rangle = 1 \)

\( T_{\text{pass2}} : 'a \text{ tree} \Rightarrow \text{nat} \)
\( T_{\text{pass2}} \langle \rangle = 1 \)
\( T_{\text{pass2}} \langle \_, \_, \hs \rangle = T_{\text{pass2}} \hs + 1 \)
## Symbol Table

The following table gives an overview of all the special symbols used in this book and how to enter them into Isabelle. The second column shows the full internal name of the symbol; the third column shows additional ASCII abbreviations. Either of these can be used to input the character using the auto-completion popup.

<table>
<thead>
<tr>
<th>Code</th>
<th>ASCII abbrev.</th>
<th>Comment</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\lambda$</td>
<td>&lt;lambda&gt;</td>
<td>function abstraction</td>
</tr>
<tr>
<td>$\equiv$</td>
<td>&lt;equiv&gt;</td>
<td>meta equality</td>
</tr>
<tr>
<td>$\neq$</td>
<td>&lt;noteq&gt;</td>
<td></td>
</tr>
<tr>
<td>$\land$</td>
<td>&lt;And&gt;</td>
<td></td>
</tr>
<tr>
<td>$\forall$</td>
<td>&lt;forall&gt;</td>
<td></td>
</tr>
<tr>
<td>$\exists$</td>
<td>&lt;exists&gt;</td>
<td></td>
</tr>
<tr>
<td>$\Longrightarrow$</td>
<td>&lt;Longrightarrow&gt;</td>
<td>meta implication</td>
</tr>
<tr>
<td>$\rightarrow$</td>
<td>&lt;longrightarrow&gt;</td>
<td>HOL implication</td>
</tr>
<tr>
<td>$\iff$</td>
<td>&lt;longleftrightarrow&gt;</td>
<td>$\leftrightarrow$ or $\longleftrightarrow$</td>
</tr>
<tr>
<td>$\Rightarrow$</td>
<td>&lt;Rightarrow&gt;</td>
<td>arrow in function types</td>
</tr>
<tr>
<td>$\leftarrow$</td>
<td>&lt;leftarrow&gt;</td>
<td></td>
</tr>
<tr>
<td>$\neg$</td>
<td>&lt;not&gt;</td>
<td></td>
</tr>
<tr>
<td>$\land$</td>
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<td>$\lor$</td>
<td>&lt;or&gt;</td>
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<td>&lt;in&gt;</td>
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<td>$\notin$</td>
<td>&lt;notin&gt;</td>
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<td>&lt;inter&gt;</td>
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<td>&lt;Union&gt;</td>
<td>Union or UN</td>
</tr>
<tr>
<td>$\bigcap$</td>
<td>&lt;Inter&gt;</td>
<td>Inter or INT</td>
</tr>
<tr>
<td>$\subseteq$</td>
<td>&lt;subseteq&gt;</td>
<td></td>
</tr>
<tr>
<td>$\subset$</td>
<td>&lt;subset&gt;</td>
<td></td>
</tr>
<tr>
<td>$\leq$</td>
<td>&lt;le&gt;</td>
<td></td>
</tr>
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</table>


### Notation

<table>
<thead>
<tr>
<th>Code</th>
<th>ASCII abbrev.</th>
<th>Comment</th>
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<td>( \geq )</td>
<td>( \geq )</td>
</tr>
<tr>
<td>( \circ )</td>
<td>( \circ )</td>
<td>function composition</td>
</tr>
<tr>
<td>( \times )</td>
<td>( \times )</td>
<td>cartesian prod., prod. type</td>
</tr>
<tr>
<td>( \mid )</td>
<td>( \mid )</td>
<td>absolute value</td>
</tr>
<tr>
<td>( \lfloor )</td>
<td>( \lfloor )</td>
<td>floor</td>
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<td>( \rfloor )</td>
<td>( \rfloor )</td>
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</tr>
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<td>ceiling</td>
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<tr>
<td>( \rceil )</td>
<td>( \rceil )</td>
<td></td>
</tr>
<tr>
<td>( \sum )</td>
<td>( \sum )</td>
<td>SUM</td>
</tr>
<tr>
<td>( \prod )</td>
<td>( \prod )</td>
<td>PROD</td>
</tr>
</tbody>
</table>

Note that the symbols “\{” and “\}” that is used for multiset notation in the book do not exist Isabelle; instead, the ASCII notation \{# and #} are used (cf. Section C.3).

#### C.2 Subscripts and Superscripts

In addition to this, subscripts and superscripts with a single symbol can be rendered using two special symbols, \(^\text{sub}\) and \(^\text{sup}\). The term \(x_{0}\) for instance can be input as \(x^{\text{sub}0}\).

Longer subscripts and superscripts can be written using the symbols \(^\text{sub}\ldots\text{sub}\) and \(^\text{sup}\ldots\text{sup}\), but this is only rendered in the somewhat visually displeasing form \(\ldots\) by Isabelle/jEdit.
C.3 Syntactic Sugar

The following table lists relevant syntactic sugar that is used in the book or its supplementary material. In some cases, the book notation deviates slightly from the Isabelle notation for better readability.

The last column gives the formal meaning of the notation (i.e. what it expands to). In most cases, this is not important for the user to know, but it can occasionally be useful to find relevant lemmas, or to understand that e.g. if one encounters the term $\sum x \in A. f \ x$, this is just the $\eta$-contracted form of $\sum x \in A. f \ x$.

The variables in the table follow the following convention:
- $x$ and $y$ are of arbitrary type
- $m$ and $n$ are natural numbers
- $P$ and $Q$ are Boolean values or predicates
- $xs$ is a list
- $A$ is a set
- $M$ is a multiset

<table>
<thead>
<tr>
<th>Book notation</th>
<th>Isabelle notation</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>$x \cdot y$</td>
<td>$x \times y$</td>
<td>times $x \ y$</td>
</tr>
<tr>
<td>$x \div y$ or $\frac{x}{y}$</td>
<td>$x \div y$</td>
<td>divide $x \ y$ (for type real)</td>
</tr>
<tr>
<td>$x \div y$</td>
<td>$x \div y$</td>
<td>divide $x \ y$ (for type nat or int)</td>
</tr>
<tr>
<td>$</td>
<td>x</td>
<td>$</td>
</tr>
<tr>
<td>$[x]$</td>
<td>$[x]$</td>
<td>floor $x$</td>
</tr>
<tr>
<td>$\lceil x \rceil$</td>
<td>$\lceil x \rceil$</td>
<td>ceiling $x$</td>
</tr>
<tr>
<td>$x^n$</td>
<td>$x \ ^\ n$</td>
<td>power $x \ n$</td>
</tr>
</tbody>
</table>
### Appendix C Notation

#### Book notation | Isabelle notation | Meaning
--- | --- | ---

**Lists**

<p>| | | |</p>
<table>
<thead>
<tr>
<th></th>
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</thead>
<tbody>
<tr>
<td>$</td>
<td>xs</td>
<td>$</td>
</tr>
<tr>
<td>$[]$</td>
<td>$[]$</td>
<td>$Nil$</td>
</tr>
<tr>
<td>$x # xs$</td>
<td>$x # xs$</td>
<td>$Cons ; x ; xs$</td>
</tr>
<tr>
<td>$[x, y]$</td>
<td>$[x, y]$</td>
<td>$x # y # []$</td>
</tr>
<tr>
<td>$[m..&lt;n]$</td>
<td>$[m..&lt;n]$</td>
<td>$upt ; m ; n$</td>
</tr>
<tr>
<td>$xs;!;n$</td>
<td>$xs;!;n$</td>
<td>$nth; xs; n$</td>
</tr>
<tr>
<td>$xs[n:=; y]$</td>
<td>$xs[n:=; y]$</td>
<td>$list_update; xs; n; y$</td>
</tr>
</tbody>
</table>

**Sets**

<p>| | | |</p>
<table>
<thead>
<tr>
<th></th>
<th></th>
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</tr>
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<tbody>
<tr>
<td>$\emptyset$</td>
<td>$\emptyset$</td>
<td>$empty$</td>
</tr>
<tr>
<td>${x, y}$</td>
<td>${x, y}$</td>
<td>$insert ; x ; (insert ; y ; {})$</td>
</tr>
<tr>
<td>$x \in A$</td>
<td>$x \in A$</td>
<td>$Set_member ; x ; A$</td>
</tr>
<tr>
<td>$x \notin A$</td>
<td>$x \notin A$</td>
<td>$\neg(x \in A)$</td>
</tr>
<tr>
<td>$A \cup B$</td>
<td>$A \cup B$</td>
<td>$union ; A ; B$</td>
</tr>
<tr>
<td>$A \cap B$</td>
<td>$A \cap B$</td>
<td>$inter ; A ; B$</td>
</tr>
<tr>
<td>$A \subseteq B$</td>
<td>$A \subseteq B$</td>
<td>$subset_eq ; A ; B$</td>
</tr>
<tr>
<td>$A \subset B$</td>
<td>$A \subset B$</td>
<td>$subset ; A ; B$</td>
</tr>
<tr>
<td>$f ; ' ; A$</td>
<td>$f ; ' ; A$</td>
<td>$image ; f ; A$</td>
</tr>
<tr>
<td>$f ; -; ' ; A$</td>
<td>$f ; -; ' ; A$</td>
<td>$vimage ; f ; A$</td>
</tr>
<tr>
<td>${x \mid P ; x}$</td>
<td>${x \mid P ; x}$</td>
<td>$Collect ; P$</td>
</tr>
<tr>
<td>${x \in A \mid P ; x}$</td>
<td>${x \in A \mid P ; x}$</td>
<td>${x \mid P ; x \land x \in A}$</td>
</tr>
<tr>
<td>${f ; y \mid P ; x ; y}$</td>
<td>${f ; x \mid f ; y \mid P ; x ; y}$</td>
<td>${z \mid \exists ; x ; y. ; z = f ; x ; y \land P ; x ; y}$</td>
</tr>
<tr>
<td>$\bigcup_{x \in A} f ; x$</td>
<td>$\bigcup_{x \in A} f ; x$</td>
<td>$\bigcup(f ; ' ; A)$</td>
</tr>
<tr>
<td>$\forall x \in A. ; P ; x$</td>
<td>$\forall x \in A. ; P ; x$</td>
<td>$Ball ; A ; P$</td>
</tr>
<tr>
<td>$\exists x \in A. ; P ; x$</td>
<td>$\exists x \in A. ; P ; x$</td>
<td>$Bex ; A ; P$</td>
</tr>
</tbody>
</table>
### C.3 Syntactic Sugar

#### Multisets

<table>
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<tr>
<th>Book notation</th>
<th>Isabelle notation</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>(</td>
<td>M</td>
<td>)</td>
</tr>
<tr>
<td>(\emptyset)</td>
<td>{#}</td>
<td>empty_mset</td>
</tr>
<tr>
<td>({x} + M)</td>
<td>add_mset x M</td>
<td></td>
</tr>
<tr>
<td>({x, y})</td>
<td>add_mset x (add_mset y {#})</td>
<td></td>
</tr>
<tr>
<td>(x \in_s M)</td>
<td>(x \in # M)</td>
<td>(x \in \text{set_mset} M)</td>
</tr>
<tr>
<td>(x \notin_s M)</td>
<td>(x \notin # M)</td>
<td>(\neg (x \in # M))</td>
</tr>
<tr>
<td>({x \in_s M \mid P x})</td>
<td>filter_mset P M</td>
<td></td>
</tr>
<tr>
<td>({f x \mid x \in_s M})</td>
<td>image_mset f M</td>
<td></td>
</tr>
<tr>
<td>(\forall x \in_s M. P x)</td>
<td>(\forall x \in # M. P x)</td>
<td>(\forall x \in \text{set_mset} M. P x)</td>
</tr>
<tr>
<td>(\exists x \in_s M. P x)</td>
<td>(\exists x \in # M. P x)</td>
<td>(\exists x \in \text{set_mset} M. P x)</td>
</tr>
<tr>
<td>(M \subseteq_s M')</td>
<td>(M \subseteq # M')</td>
<td>subseteq_mset M M'</td>
</tr>
</tbody>
</table>

#### Sums

| \(\sum A\) | \(\sum A\) | sum \((\lambda x. x) A\) |
| \(\sum_{x \in A} f x\) | \(\sum x \in A. f x\) | sum f A |
| \(\sum_{k=i}^j f k\) | \(\sum k=i..j. f k\) | sum f \{i..j\} |
| \(\sum_M\) | \(\sum \# M\) | sum_mset M |
| \(\sum_{x \in s} f x\) | \(\sum x \in \# M. f x\) | sum_mset (image_mset f M) |
| \(\sum_{x \leftarrow xs} f x\) | \(\sum x \leftarrow xs. f x\) | sum_list (map f xs) |

(analogous for products)

#### Intervals (for ordered types)

| \(\{x..\}\) | \(\{x..\}\) | atLeast x |
| \(\{..y\}\) | \(\{..y\}\) | atMost y |
| \(\{x..y\}\) | \(\{x..y\}\) | atLeastAtMost x y |
| \(\{x.<y\}\) | \(\{x.<y\}\) | atLeastLessThan x y |
| \(\{x..<y\}\) | \(\{x..<y\}\) | greaterThanAtMost x y |
| \(\{x..<y\}\) | \(\{x..<y\}\) | greaterThanLessThan x y |
Bibliography


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# Index

<table>
<thead>
<tr>
<th>Symbol</th>
<th>Page</th>
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</thead>
<tbody>
<tr>
<td>()</td>
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<td>[]</td>
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<td>⇒,</td>
<td>2</td>
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<td>≡,</td>
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<td>;</td>
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<tr>
<td>=:</td>
<td>3</td>
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<tr>
<td>(Θ),</td>
<td>305</td>
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<tr>
<td>(!),</td>
<td>306</td>
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<td>⟨⟩,</td>
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<tr>
<td>⟨_, _, _⟩</td>
<td>47</td>
</tr>
</tbody>
</table>

abstraction function, 59, 78
*ac*omplete, 51
Akra–Bazzi Theorem, 10, 41
algebra
de Morgan, 300
almost complete tree, 51
alpha-beta pruning, 281
fail-hard, 288
fail-soft, 288
amortized running time, 214
array, 127
augmented tree, 55
AVL tree, 105
balance factor, 112
balanced search trees, 85
Bellman-Ford algorithm, 202
binary search trees, 59
black height, 96
`bool`, 2
Braun tree, 128, 129
`bst`, 59
`butlast`, 306
child, 47
cmp, 60
coercion, 3
*com*plete, 50
complete tree, 50
`concat`, 305
*delete*
BST, 61
deletion by joining, 62
deletion by replacing, 61
depth, 47
deq, 219
difference array, 256
difference lists, 22
divide-and-conquer
galgorithm, 16, 17, 34, 120
recurrence, 10, 41
dom, 198
drop, 306
dynamic programming, 191
dynamic table, 215

enq, 219

False, 2
Fibonacci tree, 111
filter, 305
filter_mset, 4
first, 219
fst, 2

game tree
  value, 282

h, 49
hd, 306
head, 2
heap, 164
  binomial, 179, 180
  leftist, 167
height, 49
height-balanced tree, 111
Huffman’s algorithm, 267

IH, 5
image_mset, 4
inclusion, 3
induction, 5
  computation, 5
  strong, 42
  structural, 5
infimum, 296
inorder, 48
insert
  BST, 61
insertion sort, 14
int, 3
interval tree, 71
invariant, 63, 78
  preservation, 63
isin
  BST, 61
itrev, 8

join approach, 118

lattice, 296
  bounded, 296
  distributive, 296
Leaf, 47
length, 305
level, 47
lg, 3
lnorder, 13, 26
list (queue abstraction), 219
list (type), 2
list extensionality, 133
list-form, 67
locales, 78
loop invariant, 259

Map, 81
map, 305
map_of, 83
map_tree, 48

Master Theorem, see Akra–Bazzi Theorem
match, 256
median, 35–41
  of medians, 36–41
merge sort, 17
mh, 49
Min, 163
min_height, 49
min_list, 307
Min_mset, 163
move, 281
mset, 4
multiset, 3
multiset, 4
nat, 3
Node, 47
None, 2
observer functions, 214
optimal binary search tree, 206
option, 2
order
  de Morgan, 293
  linear, 13, 26
  bounded, 282
pair, 2
pairing heap, 243
Patricia trie, 154
pivot, 16, 34–38
position, 69, 281
preorder, 48
Priority.Queue, 163
priority queue, 163
mergeable, 163
Priority.Queue.Merge, 164
Queue, 219
queue, 219
  Hood-Melville, 221
quickselect, see selection, quickselect, 35
quicksort, 16, 34
real, 3
recurrence relation, 10, 41, 106
red-black tree, 95
replicate, 306
rev, 306
right-heavy, 240
root, 47
rotation, 67
rotation distance, 76
running time, 6
running time function, 6
runs, 21
selection, 31
  in worst-case linear time, 38
  introselect, 44
  specification, 31
  uniqueness, 31
Set, 78
set, 305
set (type), 2
set_mset, 4
set_tree, 47
skew heap, 239
snd, 2
Some, 2
sorted, 13, 64
sorted_wrt, 307
sorting, 13
  uniqueness, 25, 32
specification, 63
spine, 47
splay tree, 231
split_min, 61
stability (sorting), 26
stable, 26
subtrees, 49
sum_list, 306
supremum, 296
T, 6
tail, 2
take, 305
time function, see running time function
tl, 306
transposition table, 289
tree, 47
  2-3, 85
  binary, 47
  binomial, 179, 180
game, 281
leftist, 167
rose, 50
tree, 47
trie, 149
    binary, 151
    ternary, 158
True, 2
tuple, 2
type class, 13
type of interest, 77
type variable, 2

uniqueness
    of sorting, see sorting, uniqueness,
        see selection, uniqueness
unit, 2

weakening, 257