Lecture notes

Advanced Programming

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Prof. Dr. Michael Hanus Priv.Doz. Dr. Frank Huch

Programming Languages and Compiler Construction Department of Computer Science Kiel University

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Pre-Preface

These lecture notes are a translation of the german lecture notes associated with the "Advanced Programming" lectures held by Prof. Michael Hanus. Moreover, these notes do not translate the entire lecture by Prof. Hanus but only the chapters relevant according to the newly established pre-masters conditions for international students in Computer Science at the University of Kiel.

Preface

This lecture introduces advanced programming concepts which go beyond the first semesters of programming. The following is presented on the basis of various programming languages that represent the most important programming paradigms. Modern functional programming techniques are developed using the language Haskell. Logical and constraint-oriented programming is shown in the language Prolog. Concepts for concurrent and distributed programming are presented using the language Java.

This script is a revised version of a transcript, originally written and typeset in LATEX by Nick Prühs in SS 2009. My thanks go to Nick Prühs for the first LATEX template as well as Björn Peemöller and Lars Noelle for corrections.

One more important note: This script is only intended to give an overview of what is covered in the lecture. It does not replace attending the lecture, which is important for understanding the concepts and techniques of advanced programming. For in-depth self-study, a look at the textbooks and references given in the lecture is recommended.

Kiel, February 2017

Michael Hanus

PS: Anyone who did not find any errors while reading the notes probably has not paid enough attention. I am thankful for all hints on errors I receive, both in person as well as in writing or by e-mail.

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1 Java Generics

Since version 5.0 (released in 2004) Java supports generic programming: Classes and methods can be parameterized with types. This opens up similar possibilities as templates in C++. As an easy example we consider a very simple container class which can store either a value or no value. In Java, this could be defined as follows.

```
public class Optional {
    private Object value;
    private boolean present;

public Optional() {
        present = false;
}

public Optional(Object v) {
        value = v;
        present = true;
}

public boolean isPresent() {
        return present;
}

public Object get() {
        if (present) {
            return value;
        }
        throw new NoSuchElementException();
}
```

We use the fact that NoSuchElementException is derived from RuntimeException, that is, it is a so-called *unchecked exception* and does not need to be declared.

The class could be used as follows.

```
Optional opt = new Optional(new Integer(42));
if (opt.isPresent()) {
    Integer n = (Integer) opt.get();
        System.out.println(n);
}
```

Each time the stored object of the container class is accessed, an explicit type cast is

done. There is *no type security*: In case of a false cast a ClassCastException is thrown at runtime.

Keep in mind that the definition of the class Optional is independent of the type of the stored value; the type of value is Object. This allows using arbitrary types that are represented in an abstract way in the definition of the class: Passing a type as a parameter is called *parametric polymorphism*.

Type parameters are marked by angle brackets and are used in place of Object. The adapted class definition is.

```
public class Optional<T> {
           private T value;
           private boolean present;
           public Optional() {
                    present = false;
           public Optional(T v) {
                    value = v;
                    present = true;
           public boolean isPresent() {
                    return present;
           public T get() {
                    if (present) {
                            return value;
                    throw new NoSuchElementException();
           }
   }
The class is now used like this.
   Optional<Integer> opt = new Optional<Integer>(new Integer(42));
   if (opt.isPresent()) {
           Integer n = opt.get();
       System.out.println(n);
   }
Explicit type casts are not necessary and an expression like
   opt = new Optional<Integer>(opt);
returns a type error at compile time.
Naturally, multiple type parameters are also possible.
   public class Pair<A, B> {
```

```
private A first;
private B second;

public Pair(A first, B second) {
                this.first = first;
                 this.second = second;
}

public A first() {
                return first;
}

public B second() {
                return second;
}
```

1.1 Interaction with Inheritance

It is also possible to restrict type parameters so that only classes can be used that provide certain methods. As an example we will extend the class Optional so that it implements the interface Comparable.

```
public class Optional<T extends Comparable<T>>
implements Comparable<Optional<T>> {
          ...
          @Override
    public int compareTo(Optional<T> o) {
            if (present) {
                return o.isPresent() ? value.compareTo(o.get()) : 1;
            } else {
                return o.isPresent() ? -1 : 0;
            }
        }
}
```

For interfaces as well as inheritance the keyword extends is used. It is also possible to list multiple restrictions of type variables. For three type parameters (T, S and U) this would look like the following.

```
<T extends A<T>, S, U extends B<T,S>>
```

In this example, T has to provide the methods of A<T> and U has to provide the methods of B<T,S>. A and B need to be classes or interfaces, not type variables.

1.2 Wildcards

The class Integer is a sub class of the class Number. Therefore, the following code should be valid.

```
Optional<Integer> oi = new Optional<Integer>(new Integer(42));
Optional<Number> on = oi;
```

However, the type system does not allow this because Optional<Integer> is **no** subtype of Optional<Number>! The problem becomes more obvious when we add a setter method to the class Optional.

```
public void set(T v) {
    present = true;
    value = v;
}
```

In consequence, this would allow the following.

```
on.set(new Float(42));
Integer i = oi.get();
```

Since oi and on are names for the same objects, executing this code would lead to a type error. For this reason the type system does not allow the above code.

Nevertheless, sometimes a supertype of polymorphic classes, that is, a type that includes all other types, is needed. An Optional of unknown type is described like this.

```
Optional<?> ox = oi;
```

The symbol "?" is known as a *wildcard* type. Optional<?> represents all other instances of Optional, for example Optional<Integer>, Optional<String> and Optional<Optional<Object>>.

One downside of this approach is that only methods, which fit every type, work.

```
Object o = ox.get();
```

We can use the method set with ox but we need a value that belongs to every type: null.

```
ox.set(null);
```

This is the only possible set-call.

Unfortunately, the wildcard type "?" is often not sufficient. For example, when multiple subtypes like GUI elements are stored in a collection. This can be solved by using *bounded wildcards*.

```
<? extends A> includes all subtypes of A (covariance)
```

```
<? super A> includes all supertypes of A (contra variance)
```

Keep in mind that a type is sub- and supertype of itself, that is, for all types T the properties <T extends T> and <T super T> hold.

In the following, a few examples of using bounded wildcards are shown.

Expression	valid?	Reasoning						
Optional extends Number on = oi;								
<pre>Integer i = on.get();</pre>	no!	? could also be Float						
<pre>Number n = on.get();</pre>	yes							
on.set(n);	no!	? coulde be more specific than Number						
<pre>on.set(new Integer(42));</pre>	no!	? could also be Float						
<pre>on.set(null);</pre>	yes							
<pre>Optional<? super Integer> ox = oi;</pre>								
<pre>Integer i = ox.get();</pre>	no!	? could also be Number or Object						
<pre>Number n = ox.get();</pre>	no!	? could also be Object						
<pre>Object o = ox.get();</pre>	yes							
<pre>ox.set(o);</pre>	no!	? could also be Number						
<pre>ox.set(n);</pre>	no!	n could also have the type Float						
<pre>ox.set(i);</pre>	yes							

Figure 1.1: Expressions with wildcards

This means that we can extract objects of type A from an Optional<? extends A> but only insert null, while we can insert objects of the type A into an Optional<? super A> but only extract objects of type Object.

1.3 Type Inference

When initializing a variable with a generic type parameter, the parameter is stated twice.

Since version 7, the Java compiler is able to infer the generic type when initializing an object with new in most cases. Therefore, the type only needs to be stated in the variable declaration. When calling the new operator, the type is calculated by the diamond operator <>.

```
Optional<Integer> mv = new Optional<>(new Integer(42));
List<Optional<Integer>> 1 = new ArrayList<>(mv);
```

Only the whole type within the angle brackets can be omitted, types like <Optional<>> are not allowed. If the compiler is not able to infer the type, it still needs to be stated manually.

Besides the shorter code, the diamond operator also enables creating generic singleton objects, which was not possible before.

```
Optional<?> empty = new Optional<>();
```

This is especially useful when only one instance of immutable objects should be kept in memory.

2 Concurrent Programming in Java

An application in computer science is described as *concurrent* if it does not have a strictly sequential flow, but when the application has several activities which run concurrently, that is, (almost) in parallel. These activities are often referred to as tasks, threads, or processes. These tasks are sequentially running programs themselves. In a concurrent program, there are therefore not only one program counter which determines the next instruction to be processed but many program counters. We will see that the development of concurrent software is associated with many pitfalls. In this chapter we will discuss how to develop concurrent software that is as reliable as possible. Building on concurrent concepts, we will later also consider distributed software. We will use Java as programming language, but the concepts can also be applied to other languages.

Why do we need concurrent programming? We would often like one application to take over several tasks. At the same time the *reactivity* of the application should be preserved. Examples of such applications are listed in the following.

- GUIs
- operating system routines
- distributed applications (web servers, chat, ...)

Solution We achieve this by means of *Concurrency*). By the use of threads or processes, individual tasks of an application can be programmed and executed independently of other tasks.

Parallelism The parallel execution of several processes intends to achieve faster execution (high-performance computing).

Distributed system Several components in a network work on a problem together. Usually there is a distributed task, sometimes distributed systems are used for parallelization.

When we speak of *multitasking*, we mean that the processor time is distributed by a scheduler among concurrent threads or processes. We distinguish between two types of multitasking.

Cooperative multitasking A thread continues to calculate until it returns control (for example with yield()) or until it waits for messages (suspend()). In Java we find this in so-called *green threads*.

Preemptive multitasking The scheduler can take control from tasks. Here we often enjoy more programming comfort because we do not need to think about where we should give up control.

2.1 Synchronization

2.1.1 Interprocess Communication and Synchronization

In addition to the generation of threads or processes, communication between them is also important. This is usually done via shared memory or variables. We consider the following example in pseudo code.

```
int i = 0;
par
   { i = i + 1; }
   { i = i * 2; }
end par;
print(i);
```

Concurrency makes programs non-deterministic, that is, different results can be obtained depending on the scheduling. Thus, the above program can generate outputs 1 or 2, depending on how the scheduler executes the two concurrent processes. If the result of a program run depends on the order of the scheduling, this is called a *race condition*.

In addition to the two results 1 and 2, another result, namely 0, is also possible. This is due to the fact that it has not yet been clearly specified which actions are really executed atomically. By translating the program into byte or machine code, the following instructions can result.

```
1. i = i + 1; \rightarrow LOAD i; INC; STORE i;
2. i = i * 2; \rightarrow LOAD i; SHIFTL; STORE i;
```

Then the following sequence leads to output 0.

```
(2) LOAD i;
(1) LOAD i;
(1) INC;
(1) STORE i;
(2) SHIFTL;
(2) STORE i;
```

So we need synchronization to ensure the atomic execution of certain sections of code that work concurrently on the same resources. We call such code sections *critical sections*.

2.1.2 Synchronization with Semaphores

A well-known concept for synchronizing concurrent threads or processes goes back to Dijkstra from 1968. Dijkstra developed an abstract data type with the aim of synchronizing the atomic (uninterrupted) execution of certain program sections. These *semaphores* provide two atomic operations.

```
p(s) {
  if s >= 1
  then s = s - 1;
  else add executing thread to queue for s and suspend it;
}

v(s) {
  if waiting list for s not empty
  then wake first process in queue
  else s = s + 1;
}
```

p(s) stands for pass or passeer, v(s) stands for leave or verlaat. Now we can prevent the output 0 of our above program as follows.

```
int i = 0;
Semaphore s = 1;
par
    { p(s); i = i + 1; v(s); }
    { p(s); i = i * 2; v(s); }
end par;
```

The initial value of the semaphore determines the maximum number of processes in the critical area. Usually we find the value 1 here. We also call such semaphores binary semaphore.

Another application of semaphores is the producer-consumer problem: n producers produce goods that are consumed by m consumers. One simple solution to this problem uses an unrestricted buffer.

```
Buffer buffer = ...
Semaphore num = 0;
The producer's code looks like this.
while (true) {
    newproduct = produce();
    push(newproduct, buffer);
    v(num);
}
The consumer's code looks like this.
while (true) {
    p(num);
    prod = pull(buffer);
    consume(prod);
```

What is still missing is the synchronization to **buffer**. The synchronization can be realized by adding another semaphore.

```
Buffer buffer = ...
Semaphore num = 0;
```

```
Semaphore bufferAccess = 1;
```

Below is the adapted code of the producer.

```
while (true) {
  newproduct = produce();
  p(bufferAccess);
  push(newproduct, buffer);
  v(bufferAccess);
  v(num);
}
```

Below is the adapted code of the consumer.

```
while (true) {
  p(num);
  p(bufferAccess);
  prod = pull(buffer);
  v(bufferAccess);
  consume(prod);
}
```

However, the use of semaphores also has some disadvantages. The code with semaphores quickly looks unstructured and confusing. In addition, we cannot use semaphores compositionally: the simple code p(s); p(s); can already generate a deadlock on a binary semaphore s.

The concept of *monitors* which may be familiar from lectures like "Operating and Communication Systems" offers an improvement here. In fact, Java uses a mechanism similar to these monitors for synchronization.

2.1.3 Dining Philosophers

The $dining\ philosophers$ problem with n philosophers can be modelled as follows using semaphores.

```
p(stick1);
p(stickr);
eat();

v(stick1);
v(stickr);
}
```

However, a deadlock can occur if all philosophers take their left stick at the same time. We can avoid this deadlock by putting it back.

```
while (true) {
  think();

p(stickl);

if (lookup(stickr) == 0) {  # look up integer value of stick
  v(stickl);
} else {
  p(stickr);

  eat();

  v(stickl);
  v(stickl);
  v(stickr);
}
```

Here lookup(s) denotes a lookup function of the abstract data type semaphore that returns the integer value of a semaphore s.

The program now has a livelock, that is, individual philosophers can starve to death. We do not want to discuss this problem further here.

2.2 Threads in Java

2.2.1 The Class Thread

The API of Java offers a class Thread in the package java.lang. Own threads can be derived from this. The code to be executed in parallel is written to the run() method. After we have created a new thread with the help of its constructor, we can start it for concurrent execution with the method start().

We consider the following simple thread as an example.

```
public class ConcurrentPrint extends Thread {
  private String s;
```

```
public ConcurrentPrint(String s) {
   this.s = s;
}

public void run() {
   while (true) {
     System.out.print(s + " ");
   }
}

public static void main(String[] args) {
   new ConcurrentPrint("a").start();
   new ConcurrentPrint("b").start();
}
```

The above program flow can lead to many possible outputs:

```
a a b b a a b b ...
a a a b b ...
a b a a a b a a b b ...
a a a a a a a a a a ...
```

The latter is guaranteed if cooperative scheduling is used.

2.2.2 The Interface Runnable

Java does not offer inheriting from multiple classes. Therefore, an extension of the class Thread is often unfavorable. An alternative is the Interface Runnable:

```
public class ConcurrentPrint implements Runnable {
   private String s;

public ConcurrentPrint(String s) {
    this.s = s;
}

public void run() {
   while (true) {
     System.out.print(s + " ");
   }
}

public static void main(String[] args) {
   Runnable aThread = new ConcurrentPrint("a");
   Runnable bThread = new ConcurrentPrint("b");

   new Thread(aThread).start();
   new Thread(bThread).start();
}
```

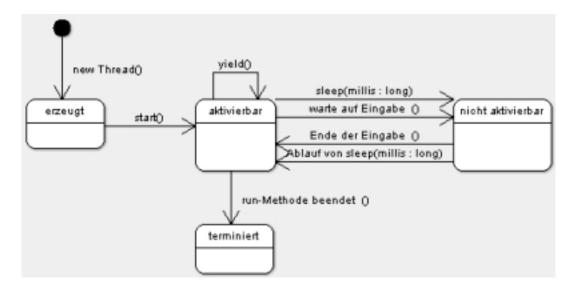


Figure 2.1: States of threads

Note that within the above implementation of ConcurrentPrint, this no longer returns an object of type Thread. The current thread object can instead be reached via the static method Thread.currentThread().

2.2.3 Properties of Thread Objects

Every thread object in Java has a number of properties.

Name Each thread has a name, such as "main-Thread", "Thread-0" or "Thread-1". Access to the name of a thread is defined using the methods getName() and setName(String). You can use these for example as names for debugging.

State Each thread is always in a certain state. An overview of these states and the state transitions is shown in figure 2.1. A thread object remains in the *terminated* state until all references to it have been discarded.

Demon A thread can be created as a background thread by calling setDaemon(true) before calling the start() method. The JVM terminates when only daemon threads are running. Examples of such threads are AWT threads or the garbage collector.

Priority Each thread in Java has a specific priority that is platform-specific.

Thread group Threads can also be divided into groups for simultaneous management.

Method sleep(long) Lets the thread sleep for the specified time. A call to this method can throw a InterruptedException which must be caught.

2.2.4 Synchronization of Threads

Java offers a monitor-like concept for thread synchronization that allows to set and release locks on objects.

The methods of a thread in Java can be declared as synchronized. In all synchronized methods of an object there may be a maximum of one thread at a time. This also includes calculations that are called in a synchronized method and unsynchronized methods of the same object. A synchronized method is not left by calling sleep(long) or yield().

Furthermore, each object has its own *lock*. When attempting to execute a method declared as synchronized, we distinguish three cases.

- 1. If the lock is released, the thread takes it.
- 2. If the thread already has the lock, it continues.
- 3. Otherwise, the thread is suspended.

The lock is released again if the method is exited in which it was acquired. Compared to semaphores, the Java approach seems more structured, you can't forget to unlock. Yet it is less flexible.

2.2.5 The Example Class Account

A simple example is intended to illustrate the use of synchronized methods. We are looking at an implementation of a class for a bank account.

```
public class Account {
    private int balance;

public Account(int initialDeposit) {
    balance = initialDeposit;
    }

public synchronized int getBalance() {
    return balance;
    }

public synchronized void deposit(int amount) {
    balance += amount;
    }
}

We now want to use the class as follows.
Account a = new Account(300);
...
a.deposit(100); // concurrent, first thread
...
a.deposit(100); // concurrent, second thread
...
System.out.println(a.getBalance());
```

The calls of the method deposit(int) should be made concurrently from different threads. Without the keyword synchronized, the output would be 500. The output 400 would also be possible (see section 2.1.1), Interprocess communication and synchronization). With synchronized, an output of 500 is guaranteed.

2.2.6 A Closer Look at synchronized

Inherited synchronized methods do not necessarily have to be synchronized again. If you overwrite such methods, you can omit the keyword synchronized. This is called *refined implementation*. The method of the superclass remains synchronized. On the other hand, unsynchronized methods can also be overwritten by synchronized ones.

Class methods declared as synchronized (static synchronized) have no interaction with synchronized object methods. The class therefore has its own lock.

You can also synchronize individual statements.

```
synchronized (expr) block
```

The expression expr must evaluate an object, whose lock is then used for synchronization. Strictly speaking, synchronized methods are only syntactic sugar: The method declaration

```
synchronized A m(args) block
is expanded to
    A m(args) {
        synchronized (this) block
}
```

Synchronizing individual statements is useful to reduce the amount of code that needs to be synchronized or sequentialized.

```
private double state;

public void calc() {
   double res;
   // do some really expensive computation
   ...
   // save the result to an instance variable
   synchronized (this) {
    state = res;
   }
}
```

Synchronizing individual statements is also useful to synchronize on other objects. We consider a simple implementation of a synchronized collection as an example.

```
class Store {
  public synchronized boolean hasSpace() {
    ...
  }
  public synchronized void insert(int i)
```

```
throws NoSpaceAvailableException {
    ...
}
```

We now want to use this collection as follows.

```
Store s = new Store();
...
if (s.hasSpace()) {
   s.insert(42);
}
```

But this leads to problems, because we cannot exclude that a re-schedule happens between the calls of hasSpace() and insert(int). Since defining special methods for such cases often turns out to be impracticable, we better use the collection like this.

```
synchronized(s) {
  if (s.hasSpace()) {
    s.insert(42);
  }
}
```

2.2.7 Differentiating Synchronization in the Context of OO

We call synchronized methods and synchronized instructions in object methods serverside synchronization. Synchronization of the calls of an object is called *client-side syn*chronisation.

For efficiency reasons, Java API objects, especially collections, are no longer synchronized. For Collections, however, there exist synchronized versions via wrappers such as synchronizedCollection, synchronizedSet, synchronizedSet, synchronizedList, synchronizedMap or synchronizedSortedMap.

Safely copying a list into an array is possible in two different ways now. First, we create an instance of a synchronized list.

```
List<Integer> unsyncList = new List<Integer>();
... // fill the list
List<Integer> list = Collections.synchronizedList(unsyncList);
Now we copy this list to an array with either a single line
    Integer[] a = list.toArray(new Integer[0]);
or with
    Integer[] b;
synchronized (list) {
    b = new Integer[list.size()];
    list.toArray(b);
}
```

With the second, two-line variant, synchronization to the list is indispensable: We access

the collection in both lines and cannot guarantee that another thread will not change the collection in the meantime. This is a classic example of client-side synchronization.

2.2.8 Communication Between Threads

Threads communicate with each other via shared objects. How to find we find out when a variable contains a value? For this there are several possible solutions.

The first option is to display the modification of a component of the object, for example, by setting a flag (boolean). However, this has the disadvantage that checking the flag leads to busy waiting. Busy waiting means that a thread is waiting for the occurrence of an event, thereby continuing to calculate and thus consuming resources like processor time. Therefore, a thread is suspended using the method wait() of the object and woken up later by using notify() or notifyAll().

```
public class C {
   private int state = 0;

public synchronized void printNewState()
   throws InterruptedException {
   wait();
   System.out.println(state);
}

public synchronized void setValue(int v) {
   state = v;
   notify();
   System.out.println("value set");
}
```

Two thread call the methods printNewState() and setValue(42) concurrently. Now the only possible output is

```
value set
```

If the call of wait() only comes after the method setValue(int) has already been left by the first thread, this leads to the output value set.

The methods wait(), notify() and notifyAll() may only be used within synchronized methods or blocks and are methods of the locked object with the following semantics.

wait() puts the executing thread to sleep and releases the lock of the object again.

notify() awakens one sleeping thread of the object and continues with its own calculation. The awakened thread now applies for the lock. If no thread sleeps, the notify() is lost.

notifyAll() does the same as notify(), only for all threads that were laid to sleep with wait() for this object.

Please note that these three methods may only be called on objects whose lock has been received before. The call must therefore be made in a synchronized method or in a synchronized block, otherwise a IllegalMonitorStateException is thrown at runtime.

We would like to write a program that outputs all changes of the state.

```
private boolean modified = false; // signals state changes
...

public synchronized void printNewState()
  throws InterruptedException {
  while (true) {
    if (!modified) {
      wait();
    }

    System.out.println(state);
    modified = false;
  }
}

public synchronized void setValue(int v) {
  state = v;
  notify();
  modified = true;
  System.out.println("value set");
}
```

One thread now executes printNewState(), other threads change the state using setValue(int). This leads to a problem: With several setting threads, the output of individual intermediate states can be lost. So setValue(int) also has to wait and wake up if necessary.

```
public synchronized void printNewState()
  throws InterruptedException {
    while (true) {
        if (!modified) {
            wait();
        }

        System.out.println(state);
        modified = false;
        notify();
    }
}

public synchronized void setValue(int v)
    throws InterruptedException {
    if (modified) {
        wait();
    }
}
```

```
state = v;
notify();
modified = true;
System.out.println("value set");
}
```

But now it is not guaranteed that the call of notify() in the method setValue(int) wakes up the printNewState thread! In Java we solve this problem with notifyAll() and accept a little busy waiting.

```
public synchronized void printNewState()
  throws InterruptedException {
  while (true) {
    while (!modified) {
      wait();
    System.out.println(state);
    modified = false;
   notify();
public synchronized void setValue(int v)
  throws InterruptedException {
  while (modified) {
   wait();
  state = v;
 notifyAll();
 modified = true;
  System.out.println("value set");
```

The wait() method is also overloaded several times in Java:

wait(long) interrupts execution for the specified number of milliseconds.

wait(long, int) interrupts the execution for the specified number of milli- and nanoseconds.

Note: It is strongly discouraged to base the correctness of the program on these overloads! The calls wait(0), wait(0, 0) and wait() all cause the thread to wait until it wakes up again.

2.2.9 Case Study: Single-element Buffer

A single-element buffer is convenient for communication between threads. Since the buffer is single-element, it can only be empty or full. A value can be written into an empty buffer via a method put from a full buffer. The value can be removed using take.

take suspends on an empty buffer, put suspends on a full buffer.

```
public class Buffer1<T> {
  private T content;
 private boolean empty;
 public Buffer1() {
    empty = true;
 public Buffer1(T content) {
   this.content = content;
    empty = false;
  }
  public synchronized T take() throws InterruptedException {
    while (empty) {
      wait();
    empty = true;
   notifyAll();
   return content;
  public synchronized void put(T o) throws InterruptedException {
    while (!empty) {
      wait();
    }
    empty = false;
   notifyAll();
   content = o;
 public synchronized boolean isEmpty() {
   return empty;
```

What is unfortunate about the above solution is that too many threads are awakened, that is, notifyAll() always awakens all reading threads as well as all writing threads, most of which are immediately put to sleep again. Can we wake up threads in a targeted way? Yes! We use special objects to synchronize the take and put threads.

```
public class Buffer1<T> {
  private T content;
  private boolean empty;
```

```
private Object r = new Object();
private Object w = new Object();
public Buffer1() {
  empty = true;
public Buffer1(T content) {
  this.content = content;
  empty = false;
public T take() throws InterruptedException {
  synchronized (r) {
    while (empty) {
      r.wait();
    synchronized (w) {
      empty = true;
      w.notify();
      return content;
    }
  }
}
public void put(T o) throws InterruptedException {
  synchronized(w) {
    while (!empty) {
      w.wait();
    synchronized (r) {
      empty = false;
      r.notify();
      content = o;
 }
}
public boolean isEmpty() {
 return empty;
```

Here the while is very important! Another thread entering the method from the outside could overtake a waiting (and just awakened) thread!

2.2.10 Exiting and Interrupting Threads

Java offers several ways to terminate threads:

- 1. terminating the run() method
- 2. aborting the run() method
- 3. calling the destroy() method (deprecated, partly no longer implemented)
- 4. demon thread and program end

With 1 and 2 all locks are released. With 3, locks are not released which makes this method uncontrollable. For this reason, this method should not be used. With 4, the locks do not matter.

Java also provides a way to interrupt threads via *interrupts* . Each thread has a flag indicating interrupts.

The Thread method interrupt() sends an interrupt to a thread, the flag is set. If the thread is sleeping due to a call to sleep() or wait(), it is awakened and a InterruptedException is thrown.

```
synchronized (o) {
    ...
    try {
        ...
        o.wait();
        ...
    } catch (InterruptedException e) {
        ...
}
```

In an interrupt after calling wait(), the catch block is not entered until the thread has recovered the lock on the o object of the surrounding synchronized block!

In contrast, in the suspension with synchronized the thread is not awakened, but only the flag is set.

The method public boolean isInterrupted() tests if a thread has received interrupts. public static boolean interrupted() tests the current thread for an interrupt and clears the interrupted flag.

Handling interrupts in a synchronized method is possible as follows:

```
synchronized void m(...) {
    ...
    if (Thread.currentThread().isInterrupted()) {
        throw new InterruptedException();
    }
}
```

If a InterruptedException is caught, the flag is also cleared. Then the flag needs to be set again!

2.3 Distributed Programming in Java

As an abstraction of network communication, Java offers the $Remote\ Method\ Invocation\ (RMI)$. This allows remote objects to be used on other computers as if they were local objects. In order for the data to be sent over a network, it must be converted (arguments and results of method calls) into byte sequences, usually referred to as serialization.

2.3.1 Serialization/Deserialization of Data

The serialization of an object o returns a byte sequence, the deserialization of the byte sequence returns a new object o1. Both objects should be the same with regard to their behavior, but have different object identities, i.e. o1 is a copy of o.

The (de-)serialization takes place recursively. Thus, contained objects must also be (de)serialized. To specify that an object can be serialized, Java offers the interface Serializable.

```
public class C implements java.io.Serializable { ... }
```

This interface does not contain any methods and is therefore only a "marker interface" that specifies that objects of this class can be (de-)serialized.

During serialization, certain time- or security-critical parts of an object can be hidden using the keyword transient:

```
protected transient String password;
```

Transient values should be set explicitly after descrialization (for example a timer) or should not be used (for example passwords).

For serializing you use the class ObjectOutputStream, for deserializing the class ObjectInputStream. Their constructors are a OutputStream and InputStream respectively. Objects can be written

```
public void writeObject(Object o)
using and read by means of
  public final Object readObject()
and a cast of appropriate type.
```

This could be used to "manually" transfer objects from one machine to another (for example by using socket connections) and then work with them on the other machine. You can avoid this manual work, however, if you only want to use certain functionalities of an object somewhere else. This is made possible in Java by Remote Method Invocation.

2.3.2 Remote Method Invocation (RMI)

In OO-programming we have a client-server view of objects. When a method is called, the calling object is seen as the client and the called object as the server.

In a distributed context, messages become real messages on the Internet (transmitted via TCP). Processes that communicate with each other are

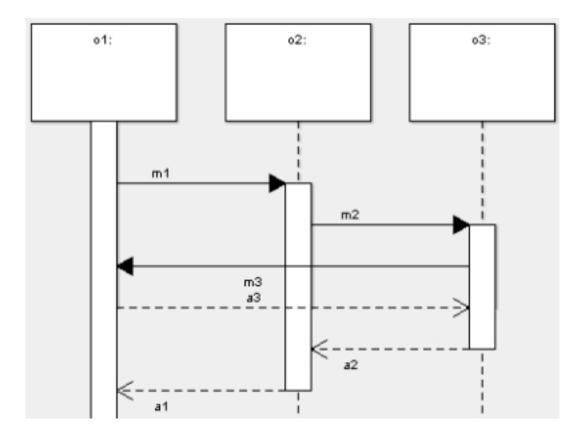


Figure 2.2: Remote Method Invocation in Java

- servers that provide information.
- clients that request information.

Any communication pattern (for example peer-to-peer) can be represented. The idea of RMI goes back to *Remote Procedure Call (RPC)*, which was developed for C.

First, an RMI client needs a reference to the remote object. The RMI registration serves this purpose. It requests the reference using a URL:

rmi://hostname:port/servicename

hostname can be a computer name or an IP address, servicename is a string describing an object. The default port of RMI is 1099.

There is also a second way to access a remote object in a program: as the argument of a method call. Usually you can use the above registration only for the "'first contact", then (remote) objects are exchanged and used transparently (like local objects).

Such objects can be distributed on any computer. Method calls for remote objects are implemented by network communication.

The interface for RMI is divided into stub and skeleton. Since Java 5 these are no longer visible, but are generated implicitly at runtime.

The network communication takes place via TCP/IP, but this is also not visible for the application programmer.

The parameters and the return value of a remotely called method must of course be converted into bytes and be transferred. The following rules apply.

- For Remote objects only a reference of the object is transferred.
- Serializable objects are converted to byte[]. On the other side a copy of the object will be created.
- Primitive values are copied.

To use objects from remote nodes, an RMI Service Interface must first be defined for this object. This interface describes the methods that can be called from another computer. Consider a simple "Flip" server as an example, that is, an object containing a Boolean state which can be changed by a flip message. The RMI service interface can then be defined as follows.

```
import java.rmi.*;

public interface FlipServer extends Remote {
   public void flip() throws RemoteException;
   public boolean getState() throws RemoteException;
}
```

An implementation of the RMI interface on the server side could look like the following.

In older Java versions it was necessary to generate the stub and skeleton classes with the RMI compiler rmic - this is no longer the case. The class UnicastRemoteObject represents the simplest way of realizing remote objects. All necessary translation steps are done automatically. In some cases, it may be necessary to intervene in the (de-)serialization process yourself to define properties of the RemoteObjects. For this purpose,

other interfaces exist.

2.3.3 The RMI Registry

To access a server object from another host, Java provides an RMI registry. To access a remote object, an RMI registry server must be initialized on the host on which the server object is executed. This RMI registry server can be started explicitly using the command rmiregistry (for example, in a UNIX shell). Its use for object registration is shown in the following example.

```
import java.rmi.*;
import java.net.*;

public class Server {
    public static void main(String[] args) {
        try {
            String host = args.length >= 1 ? args[0] : "localhost";
            String uri = "rmi://" + host + "/FlipServer";

            FlipServerImpl server = new FlipServerImpl();
            Naming.rebind(uri, server);
        } catch (MalformedURLException|RemoteException e) { ... }
    }
}
```

rebind registers the name of the server object. On the client side, the RMI registry must be contacted to get a reference to the remote object so that its flip method can be called.

```
import java.rmi.*;
import java.net.*;

public class Client {
   public static void main(String[] args) {
      try {
       String host = args.length >= 1 ? args[0] : "localhost";
       String uri = "rmi://" + host + "/FlipServer";

      FlipServer s = (FlipServer) Naming.lookup(uri);
      s.flip();
      System.out.println("State: " + s.getState());
      } catch (MalformedURLException|NotBoundException|RemoteException e) {
        ...
    }
   }
}
```

However, remind that the RMI registry is only used for a "'first contact" und usually only on server object is registered. After that, other remote objects are passed as prameters or return values and nodes in the distributed system get aware of all relevant objects of

the whole system.

Thus, RMI represents a continuation of the sequential programming on distributed objects. This means that several distributed processes can access an object "'simultaneously". So you also have to consider the problem of concurrent synchronization! However, synchronization can be more difficult here, as the following example shows.

As we have seen above, *client-side synchronization* is to guarantee the atomic execution of several methods which can already be synchronized. For this purpose we consider the simple Flip-server once again and a client that uses the flip method twice without interrupt calls.

```
FlipServer s = (FlipServer) Naming.lookup(uri);
synchronized(s) {
    System.out.println("State1: " + s.getState());
    s.flip();
    Thread.sleep(2000);
    s.flip();
    System.out.println("State2: " + s.getState());
}
```

Since the client synchronizes both calls on the flip server s, no one else should be able to change the state of the flip server during this time, so that the results of both outputs are always the same. This would also be the case in the purely concurrent context, but in the distributed context this no longer applies, since synchronization does not take place on the remote objects, but on the local stub objects!

Dynamic loading, security concepts or distributed memory cleanup (garbage collection) are other aspects of Java RMI that we do not consider here.

3 Functional Programming

One focus of this lecture is this chapter on functional programming. The practical relevance of functional programming is emphasized, for example, by Thomas Ball and Benjamin Zorn of Microsoft in the article [?], which bears the meaningful subtitle "Industry is ready and waiting for more graduates educated in the principles of programming languages".

Second, would-be programmers (CS majors or non-majors) should be exposed as early as possible to functional programming languages to gain experience in the declarative programming paradigm. The value of functional/declarative language abstractions is clear: they allow programmers to do more with less and enable compilation to more efficient code across a wide range of runtime targets.

Another interesting recommendation of the authors is the following.

First, computer science majors, many of whom will be the designers and implementers of next-generation systems, should get a grounding in logic, its application in design formalisms, and experience the creation and debugging of formal specifications with automated tools...

Therefore, the study of mathematical foundations and logic is an important aspect of a computer science degree, but it is not part of this lecture.

Functional programming offers a number of advantages over classical imperative programming.

- High level of abstraction, no manipulation of memory cells
- No side effects, therefore easier code optimization and better comprehensibility
- Programming via properties, not via the execution sequence
- Implicit memory management
- Simpler proof of correctness and verification
- Compact source programs, therefore shorter development time, more readable programs, better maintainability
- Modular program structure, polymorphism, higher-order functions, reusability of code

For practical programming, knowledge of functional programming is important, as functional programming techniques and language constructs lead to better structured programs, as explained in the article [?]. Therefore, functional concepts can also be found in

many modern programming languages in a limited form. However, we will first introduce purely functional programming.

In functional programs, a *variable* represents an unknown value. A *program* is a set of function definitions. The memory is not explicitly usable, but is automatically managed and cleared. A *program flow* consists of the reduction of expressions. This goes back to the mathematical theory of the λ calculus from Church [?]. In the following we introduce the purely functional programming using the Haskell programming language [?].

3.1 Expressions and Functions

In mathematics, a variable represents unknown (arbitrary) values, so that we often use expressions like

$$x^2 - 4x + 4 = 0 \Leftrightarrow x = 2$$

In imperative languages, on the other hand, we often see expressions such as the following.

$$x = x + 1$$
 or $x := x + 1$

This represents a contradiction to mathematics. In functional programming, as in mathematics, variables are interpreted as unknown values (and not as names for memory cells)!

While functions in mathematics are used for calculation, we use procedures or functions in programming languages for structuring. There, however, is no real connection due to side effects. In functional programming languages, however, there are no side effects, so every function call with the same arguments produces the same result.

Functions can be defined in Haskell as follows.

```
f x1 \dots xn = e
```

f is the function name, x1 to xn are formal parameters or variables, and e is the body, an expression over x1 to xn.

Expressions are constructed in Haskell by combining elements from the (incomplete) list below. gebildet werden:

- 1. Numbers: 3, 3.14159
- 2. Basic operations: 3 + 4, 5 * 7
- 3. Function application: (f e1 ... en). Parentheses can be omitted if the context allows.
- 4. Conditional expressions: (if b then e1 else e2)

Haskell almost looks like a scripting language. In contrast to script languages like PHP, Ruby or Python, Haskell has all elements to realize even large software systems. In particular, unlike scripting languages, Haskell is *strictly typed*, that is, all values and expressions have a type that is checked by the Haskell system *before* the program is executed. We will discuss the different data types in more detail in chapter 3.2. Here,

we want to annotate the type for all functions to make clear what they are used for.¹

A basic data type is Int, the set of integers (or a finite subset of it). The type of a function is annotated by "::", where several argument types and the result type are separated by " \rightarrow ". Furthermore, the intuitive meaning of functions should be explained by a *comment* before the function definition, where comments are introduced by two minus signs and reach until the end of the line.

As an example, consider a function for calculating squares. We can define this in Haskell as follows.

```
-- Computes the square of a number.
square :: Int -> Int
square x = x * x
```

If this definition is stored in the file Square.hs, then you can use the Haskell interpreter ghci, which belongs to the Glasgow Haskell Compiler (GHC), as follows to use this function.

A function for calculating the minimum of two numbers can be defined in Haskell in this way.

```
-- Computes the mininum of two numbers.
min :: Int -> Int -> Int
min x y = if x <= y then x else y
```

Next, we have look at the factorial function. It is defined mathematically as follows.

$$n! = \begin{cases} 1, & \text{falls } n = 0\\ n \cdot (n-1)!, & \text{otherwise} \end{cases}$$

The Haskell implementation looks like this.

```
-- Computes the factorial of a non-negative number. fac :: Int \rightarrow Int fac n = if n == 0 then 1 else n * fac (n - 1)
```

¹Haskell can, unlike many other languages, infer function types, so that it is not necessary write the type down. Nevertheless, it is a better programming style to add function types for program documentation.

3.1.1 Evaluation

The evaluation of function definitions in Haskell is done by oriented calculation from left to right: First the current parameters are bound, that is, the formal parameters are replaced by the current ones. Then the left side is replaced by the right side.

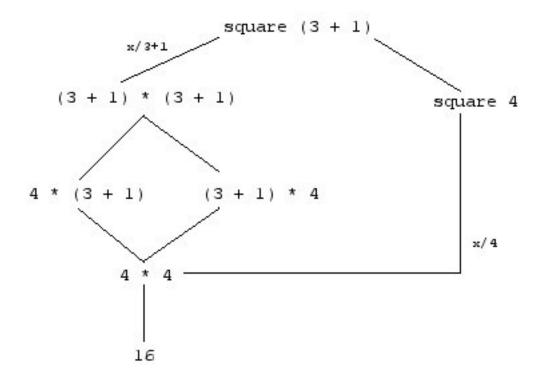


Figure 3.1: Ways to evaluate a function

Figure 3.1 shows how functions can be evaluated. The right branch shows how a function is evaluated in Java, the left branch resembles the evaluation in Haskell if double calculations of 3 + 1 are omitted.

Another example is the evaluation of a call to our function fac.

We now want to develop an efficient function for calculating Fibonacci numbers. Our first version is directly motivated by the mathematical definition.

However, this variant is extremely inefficient: its execution time is $O(2^n)$.

How can we improve our first version? We calculate the Fibonacci numbers from the bottom: The numbers are enumerated from 0 until the nth number is reached: 0 1 1 2 3 ... fib(n).

This programming technique is known by the name of accumulator technique. To do this, we must always keep the two previous numbers as parameters.

Here, fibn is the nth Fibonacci number and fibnp1 the (n+1)th. Thus we achieve a linear runtime.

From a software-technical point of view, our second variant is unattractive: We want to avoid other external calls to fib2'. How this works is described in the next section.

3.1.2 Local Definitions

Haskell offers several ways to define functions locally. One possibility is the keyword where:

```
fib2 :: Int -> Int
fib2 n = fib2' 0 1 n
  where fib2' fibn fibnp1 n =
        if n == 0
        then fibn
        else fib2' fibnp1 (fibn + fibnp1) (n - 1)
```

where definitions are visible in the previous equation, outside they are invisible.

Alternatively, the keyword let can be used.

```
fib2 n =
  let fib2' fibn fibnp1 n =
      if n == 0
      then fibn
      else fib2' fibnp1 (fibn + fibnp1) (n - 1)
  in fib2' 0 1 n
```

In contrast to where, let is an expression. The fib2' defined by let is only visible within the let expression.

let ... in ... can occur as an arbitrary expression.

```
(let x = 3

y = 1

in x + y) + 2
```

The above expression evaluates to 6.

The syntax of Haskell does not require parentheses and no separation of the individual definitions (for example by a semicolon) when defining such blocks. In Haskell, the *off-side rule* applies: The next symbol after where or let that is not a whitespace defines a *block*.

- If the next line starts to the right of the block, it belongs to the same definition.
- If the next line starts at the edge of the block, a new definition in the block starts here.
- However, if the next line starts to the left of the block, the block before it ends here.

Local definitions offer a number of advantages.

- Name conflicts can be prevented
- Wrong usage of helper functions can be prevented
- Better readability
- Redundant computations can be avoided

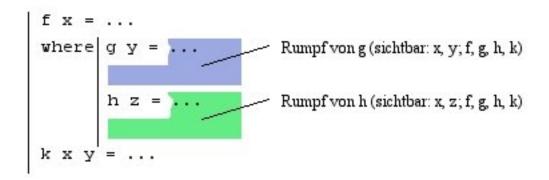


Figure 3.2: Off-side rule in Haskell

• Helper functions with less parameters

Let us look at an example to avoid multiple calculations. Instead of the confusing function definition

By using the local declaration constructs let and where it is also possible to save parameters for helper functions. This is illustrated by the following example.

The predicate isPrim is supposed to check if the given number is a prime number. In Haskell we can express this as follows.

Here /= expresses inequality, && stands for a conjunction, || for the logical or and div for integer division.

In the last line we do not need further parentheses, because && binds stronger than ||.

The n is visible in checkDiv, because the latter is locally defined.

3.2 Data Types

3.2.1 Basic Data Types

Integers

We have already used a basic data type of Haskell before: integers. In fact, Haskell distinguishes between two types of integers:

```
Int: Values -2^{31} ... 2^{31} - 1
Integer: arbitrary size (limited by memory)
Arithmetic operators: + - * div mod
Comparison operators: < > <= >= == /=
```

Booleans

Booleans are another basic data type.

```
Bool: True False

Operators: && || not == /=

== means equivalence, /= exclusive or (XOR).
```

Floats

Floating point numbers are a basic data type as well.

```
Float: 0.3 -1.5e-2

Operators: similar to Int, but / instead of div, no mod
```

Chars

Unicode characters are a basic data type, too.

```
Char: 'a' '\n' '\NUL' '\214'
```

The operators are defined in the library Data. Char and can be imported into the program by adding "import Data. Char" to the beginning of the Haskell program.

```
chr :: Int -> Char
ord :: Char -> Int
```

The operator "::" is used for optional type annotations of expressions, as explained below.

3.2.2 Type Annotations

As already mentioned, Haskell is a strictly typed programming language, that is, all values and expressions in Haskell have a type which can also be annotated.

```
3 :: Int
```

In this example, 3 is a value or expression, and Int is a type expression. Other examples of type annotations are as follows.

```
3 :: Integer
(3 == 4) || True :: Bool
(3 == (4 :: Int)) || True :: Bool
```

We can also specify type annotations for functions. These are written in a separate line:

```
square :: Int -> Int
square x = x * x
```

But what is the type of min (see section ??), considering that the function has two arguments?

```
min :: Int -> Int -> Int
```

A function arrow is therefore also used between the argument types written. We will see later why this is necessary.

3.2.3 Algebraic Data Structures

Own data structures can be defined as new data types. Values are built using *constructors*. Constructors are freely interpreted functions and therefore cannot be reduced.

Defining an Algebraic Data Type

Algebraic data types are defined in Haskell as follows.

```
data \tau = C_1 \tau_{11} ... \tau_{1n_1} | ... | C_k \tau_{k1} ... \tau_{kn_k}
```

where

- τ is the newly defined type,
- $C_1, ..., C_k$ are the constructors and
- $\tau_{i1}, ..., \tau_{in_i}$ are the argument types of the constructor C_i . Therefore $C_i :: \tau_{i1} \rightarrow \cdots \rightarrow \tau_{in_i} \rightarrow \tau$

holds.

Note: Both type and constructor names must begin with a capital letter in Haskell!

Examples

1. Enumerated types (only nullary constructors):

```
data Color where
  Red :: Color
  Blue :: Color
  Yellow :: Color
```

This data type defines three values of the type Color. Bool is also an enumerated type, predefined as

```
data Bool where
    True :: Bool
    False :: Bool
```

2. Types with only one constructor:

```
data Complex where
    Complex :: Float -> Float -> Complex
Complex 3.0 4.0 :: Complex
```

This is allowed because Haskell works with separate namespaces for types and constructors.

How do we select individual components? In Haskell we use *pattern matching* instead of explicit selection functions (this possibility will be explained later):

```
-- Add two complex numbers:
addC :: Complex -> Complex -> Complex
addC (Complex r1 i1) (Complex r2 i2) = Complex (r1+r2) (i1+i2)
```

3. Lists (mixed types): For now, we consider only lists with elements of type Int:

```
data List where
  Nil :: List
  Cons :: Int -> List -> List
```

Nil represents the empty list. The function append allows concatenating two lists.

This is a definition using several equations, where the first suitable one is selected. For example, a function call could be reduced in the following way.

```
append (Cons 1 (Cons 2 Nil)) (Cons 3 Nil)
= Cons 1 (append (Cons 2 Nil) (Cons 3 Nil))
= Cons 1 (Cons 2 (append Nil (Cons 3 Nil)))
= Cons 1 (Cons 2 (Cons 3 Nil))
```

Haskell's predefined lists look like this.

```
data [Int] = [] | Int:[Int]
```

[] corresponds to Nil, ":" is equivalent to Cons and [Int] is List.

The operator ":" is right-associative. Therefore, the following holds.

```
1:(2:(3:[]))
is equal to
1:2:3:[]
which can be written in the more compact way
[1,2,3]
```

Haskell also offers the operator ++ instead of append, predefined in the Prelude.

```
[] ++ ys = ys
(x:xs) ++ ys = x : xs ++ ys
```

Operators are binary functions that are used infix and begin with a special character. By means of parentheses, operators can be used like normal functions.

```
(++) :: [Int] -> [Int] -> [Int]
[1] ++ [2] is equal to (++) [1] [2].
```

The other way around, binary functions can be used infix with single inverted commas '...'.

```
div 4 2
can be written as
4 `div` 2
```

For user-defined data types, it is not automatically possible to compare or output them. For this the keyword deriving can be added after the data type definition.

```
data MyType where
    ...
    deriving (Eq, Show, Ord)
```

3.3 Polymorphism

For the definition of universally usable data structures and operations Haskell supports type polymorphism. To explain this in more detail, we will determine the length of a list as an example:

```
-- Compute the length of a list of integers:

length :: [Int] -> Int

length [] = 0

length (_:xs) = 1 + length xs
```

Of course, this definition also works for other lists, for example of type [Char], [Bool] or [[Int]].

```
Generally, one could say length :: \forall Type \tau. [\tau] -> Int
```

which is expressed in Haskell by means of type variables.

3 Functional Programming

```
length :: [a] -> Int
What is the type of (++)?
  (++) :: [a] -> [a] -> [a]
```

In consequence only lists with the same argument type can be concatenated. We consider another example.

```
-- Compute the last element of a list:
last :: [a] -> a
last [x] = x
last (x:xs) = last xs
```

This works because [a] is a list type, and [x] is a one-element list (corresponds to x:[]). However, we must not swap the two rules, otherwise every call to the function will yield a run-time error!

How can we define polymorphic data types ourselves? Haskell provides *type constructors* for the construction of types.

These data type definitions therefore look similar to the previous ones but here

- K is a type constructor (not a type),
- a_1, \ldots, a_m are type variables and
- τ_{ik} are type expressions like basic types, type variables or a type constructor applied to type expressions.

Functions and constructors are applied to values or expressions and generate values. Similarly, type constructors are applied to types and generate types.

As an example for a polymorphic datatype we want to model partial values in Haskell.

```
data Maybe a where
   Nothing :: Maybe a
   Just :: a -> Maybe a
```

Then "Maybe Int" and also "Maybe (Maybe Int)" is a valid type.

If a type constructor in a type is applied to type variables, then the resulting type is also called *polymorphic*, as in the following example.

```
isNothing :: Maybe a -> Bool
isNothing Nothing = True
isNothing (Just _) = False
```

Another good example of a polymorphic datatype is the binary tree.

```
data Tree a where
   Leaf :: a -> Tree a
   Node :: (Tree a) -> (Tree a) -> (Tree a)
-- Compute the height of a binary tree:
height :: Tree a -> Int
```

```
height (Leaf _) = 1
height (Node tl tr) = 1 + max (height tl) (height tr)
```

In Haskell, polymorphic lists are also predefined as syntactic sugar.

```
data [a] = [] | a : [a]
```

Although this definition is not a syntactically valid Haskell program, it can be understood as follows: The square brackets around [a] are the type constructor for lists, a is the element type, [] is the empty list and : is the list constructor.

For this reason, the following expressions are all the same.

```
(:) 1 ((:) 2 ((:) 3 []))
1 : (2 : (3 : []))
1 : 2 : 3 : []
[1,2,3]
```

According to the above definition, any complex list types such as [Int], [Maybe Bool] or [[Int]] are possible.

Some functions on lists are for example head, tail, last, concat and (!!).

The last function can also be defined as follows.

```
(x:_ ) !! 0 = x
(_:xs) !! n = xs !! (n - 1)
```

Strings are defined in Haskell as lists of characters.

```
type String = [Char]
```

Here, "type" initiates the definition of a *Typsynonyms*, that is, a new name (String) for another type expression ([Char]).

Thus the string "Hello" corresponds to the list 'H':'e':'l':'l':'o':[]. For this reason, all list functions also work for strings.

```
length ("Hello" ++ " folks!")
```

The above expression is evaluated 12.

Other predefined type constructors in Haskell are as follows.

• Sum of two types

```
data Either a b where
  Left :: a -> (Either a b)
  Right :: b -> (Either a b)
```

This can be used, for example, to write values of "'different types"' to a list.

```
[Left 42, Right "Hallo"] :: [Either Int String]
```

Sum types can be processed via patterns, for example.

```
valOrLen :: Either Int String -> Int
valOrLen (Left v) = v
valOrLen (Right s) = length s
```

• Tuple

```
data (,) a b = (,) a b
data (,,) a b c = (,,) a b c
```

Some functions have already been defined for this as well.

In principle unzip is the inversion of zip, that is, if "unzip zs" evaluates to the result (xs,ys), then "zip xs ys" evaluates again to zs. However, conversely unzip (zip xs ys) does not always evaluate to (xs,ys)!

3.4 Pattern Matching

As we have already seen, functions can be defined by several equations using pattern matching.

```
f pat_{11} \dots pat_{1n} = e_1

:

f pat_{k1} \dots pat_{kn} = e_k
```

This results in very concise programs, since you only have to define the right-hand sides for the special cases specified by the patterns. For multiple rules, Haskell selects and applies the textual first rule with a matching left side. Overlapping rules are allowed in principle, but they can lead to programs that are not very readable and should therefore be avoided.

3.4.1 Structure of Patterns

In principle, the patterns are data terms (that is, they contain no defined functions) with variables. The following patterns are possible.

- x (Variable): matches always, the variable is bound to the current value.
- _ (Wildcard): matches always, no binding.
- C $pat_1...pat_k$ where C is a k-ary Constructor: matches, if the same construktor and arguments match with $pat_1, ..., pat_k$
- x@pat (as pattern): matches if pat matches; Additionally, x is bound to the whole matched term.

The as pattern also avoids overlapping patterns.

```
last :: [a] -> a
last [x] = x
last (x : xs@(_:_)) = last xs
```

In addition there are so-called (n+k) patterns for positive integers. We do not them explain here, because they are often not supported and not important.

Patterns can also be used with let and where:

```
unzip :: [(a, b)] -> ([a], [b])
unzip ((x,y) : xys) = (x:xs, y:ys)
where
  (xs,ys) = unzip xys
```

3.4.2 Case Expressions

Sometimes it is also useful to branch expressions using pattern matching.

```
case e of pat_1 -> e_1 \vdots pat_n -> e_n
```

The above defines an expression with type $e_1, ..., e_n$, which must all have the same type. $e, pat_1, ..., pat_n$ must also have the same type. After the keyword of, the off-side rule also applies, that is, the patterns $pat_1, ..., pat_n$ must all begin in the same column.

The result of **e** is matched against pat_1 to pat_n one after the other. If a pattern matches, the whole case expression is replaced by the corresponding e_i .

As an example, consider extracting the lines of a string.

```
-- Breaks a string into a list of lines where a line is terminated at a -- newline character. The resulting lines do not contain newline characters.
```

3.4.3 **Guards**

Each pattern can have an additional boolean condition, also called quard.

otherwise is not a keyword, but a function that always evaluates to True.

By combining guards and case expressions, the first n elements of a list can be extracted.

```
-- Returns prefix of length n.

take :: Int -> [a] -> [a]

take n xs | n <= 0 = []

| otherwise = case xs of

[] -> []

(x:xs) -> x : take (n-1) xs
```

Guards are also allowed for let, where and case.

3.5 Higher Order Functions

In Haskell, functions are not only used to define calculation methods, but functions are also "first class citizens" because they can be used like all other values (for example numbers). This means that functions can appear in data structures and also as parameters or results of other functions. In the latter case one speaks of "'higher-order functions"'... Higher-order functions can used for

- generic programming
- defining program schemes (control structures)

This enables us to achieve a better reusability and a higher modularity of the program code.

3.5.1 Example: Derivative Function

The derivative function is a function that returns a new function for a function. The numerical calculation looks like this.

$$f'(x) = \lim_{dx \to 0} \frac{f(x+dx) - f(x)}{dx}$$

An implementation with a small dx could look like the following.

```
dx :: Float
dx = 0.0001

-- Computes the derivation of a (continuous) function.
derive :: (Float -> Float) -> (Float -> Float)
derive f = f'
  where f' :: Float -> Float
    f' x = (f (x + dx) - f x) / dx
```

Now, (derive sin) 0.0 evaluates to 1.0 and (derive square) 1.0 evaluates to 2.00033.

3.5.2 Anonymous Functions (Lambda Abstraction)

Sometimes one does not want to give every defined function a name (like square), but to write it directly where is is needed, as shown in the example below.

```
derive (\x -> x * x)
```

The argument corresponds to the function $x \mapsto x^2$. Such a function without name is also called *lambda abstraction* or *anonymous function*. Here \ stands for λ , x is a parameter and x * x is an expression (the body of the function).

In general, anonymous functions in Haskell are defined as follows.

```
\setminus p_1 \ldots p_n \rightarrow e
```

where p_1, \ldots, p_n are patterns and e is an expression.

We can also write the following.

```
derive f = \langle x \rangle (f(x + dx) - f(x) / dx
```

When using this function, we can observe that it behaves approximately like the derived function $(y \mapsto 2y)$.

```
(derive (\langle x - \rangle x * x)) 0.0 -> 0.0001
(derive (\langle x - \rangle x * x)) 2.0 -> 4.0001
(derive (\langle x - \rangle x * x)) 4.0 -> 8.0001
```

But derive is not the only function with a functional result. For example, the function add can be defined in three different ways.

```
add :: Int -> Int -> Int
add x y = x + y

or
add = \x y -> x + y

or
add x = \y -> x + y
```

So add can also be seen as a constant that returns a function as a result, or as a function that takes an Int and returns a function that takes another Int and only then returns an Int.

Therefore, the types $\operatorname{Int} \to \operatorname{Int} \to \operatorname{Int}$ and $\operatorname{Int} \to \operatorname{Int}$) must be identical. In fact, the type constructor " \to " is defined as right-associative, so that this binding

always applies if no brackets are written. You should note that $(a \rightarrow b) \rightarrow c$ not the same as $a \rightarrow b \rightarrow c$ or $a \rightarrow (b \rightarrow c)!$

So it would make sense to write the following independently of the definition of add.

```
derive (add 2)
```

If a function is applied to "too few" arguments, this is called *partial application* application, partial. The partial application is made possible syntactically by currying. The name *currying* goes back to *Haskell B. Curry*, who discovered the following isomorphy in the 1940s:

$$[A \times B \to C] \simeq [A \to (B \to C)]$$

This means that a function with two arguments can also be interpreted as a function with one argument, which then returns another function for the second argument.

Actually, this technique was established much earlier by *Moses Schönfinkel* [?] in 1924. But because this article was published in German and "*Schönfinkeling*" cannot be pronounced so well in English, the term "Currying" has prevailed.

A number of functions can now be defined with the help of partial applications.

- take 42 :: [a] -> [a] yields the first up to 42 elements of a list.
- (+) 1 :: Int -> Int is the increment function.

For operators, so-called *sections* offer an additional, shortened notation.

- (1+) is the increment function.
- (2-) is equal to $x \rightarrow 2-x$.
- (/2) is equal to $\x \rightarrow x/2$.
- (-2) is *not* equal to $\x -> x-2$, because the compiler cannot distinguish the minus sign from the unary minus operator.

Therefore, operators can also be partially applied to the second argument.

```
(/b) a = (a/) b = a / b
```

The order of arguments is a design decision because of partial application. The order can be modified with λ -abstractions and the function flip.

```
flip :: (a \rightarrow b \rightarrow c) \rightarrow b \rightarrow a \rightarrow c
flip f = x y \rightarrow f y x
```

The function flip can be used, for example, with take in order to supply the list argument first.

```
(flip take) :: [a] -> Int -> [a]
(flip take) "Hello World!" :: Int -> [Char]
```

3.5.3 Generic Programming

We consider the following functions incList and codeStr.

The expression codeStr "Informatik" evaluates to "Jogpsnbujl". We observe that the definitions of incList and codeStr are nearly identical. Only the function that is applied to the list elements differs.

A generalized version is the function map.

```
map :: (a -> b) -> [a] -> [b]
map _ [] = []
map f (x:xs) = f x : map f xs
```

incList and codeStr can be defined more elegantly by using map.

```
incList = map (+1)
codeStr = map code
```

We look at another two examples: A function that yields the sum of all elements in a list and a function that calculates the sum of a string's Unicode values.

```
sum :: [Int] -> Int
sum [] = 0
sum (x:xs) = x + sum xs

checkSum :: String -> Int
checkSum "" = 1
checkSum (c:cs) = ord c + checkSum cs
```

Is there a common pattern? Indeed, both functions can be defined more easily by means of the powerful function foldr.

```
foldr :: (a -> b -> b) -> b -> [a] -> b
foldr _ e [] = e
foldr f e (x:xs) = f x (foldr f e xs)

sum = foldr (+) 0
checkSum = foldr (\c res -> ord c + res) 1
```

To understand foldr, the following perspective might help. The first argument of foldr of type $(a \rightarrow b \rightarrow b)$ replaces the list constructor (:) in the list. The second argument of type b is the replacement for the empty list []. Note that the type of the supplied function matches the type of (:).

The following expressions are equivalent.

```
foldr f e [1,2,3]
= foldr f e ((:) 1 ((:) 2 ((:) 3 [])))
= (f 1 (f 2 (f 3 e)))
```

The general approach when designing such functions is as follows: Identify a common pattern and implement it with functional parameters.

Another pattern that we know is the function filter that filter elements with certain properties from a list.

We can use filter, for example, to transform a list into a set, that is, removing all duplicates.:

```
nub :: [Int] -> [Int]
nub [] = []
nub (x:xs) = x : nub (filter (/= x) xs)
```

By using filter, we can implement Quicksort to sort lists.

```
qsort :: [Int] -> [Int]
qsort [] = []
qsort (x:xs) =
  qsort (filter (<= x) xs) ++ [x] ++ qsort (filter (> x) xs)
```

filter can also be defined by using foldr.

```
filter p = foldr (\xys -> if p x then x:ys else ys) []
```

The function foldr is a very generic skeleton, it is equivalent to a katamorphism in category theory.

Using foldr can have drawbacks sometimes. The expression

```
foldr (+) 0 [1,2,3] = 1 + (2 + (3 + 0))
```

leads to a large calculation that is first built up on the stack and then evaluated in the end.

An improved version can be implemented by means of the accumulator technique.

```
sum xs = sum' xs 0
where sum' :: [Int] -> Int -> Int
    sum' []    s = s
    sum' (x:xs) s = sum' xs (x + s)
```

The expression sum [1,2,3] is now being reduced to ((0 + 1) + 2) + 3, which can be evaluated immediately.

An equivalent implementation can be achieved by using a different version of fold.

```
foldl :: (a \rightarrow b \rightarrow a) \rightarrow a \rightarrow [b] \rightarrow a
foldl _ e [] = e
foldl f e (x:xs) = foldl f (f e x) xs
```

Hence, a call fold f e $(x_1: x_2: ...: x_n: [])$ is replaced with f ... (f (f e x_1) x_2) ... x_n .

Now sum can be defined even simpler.

```
sum = foldl (+) 0
```

3.5.4 Control Structures

Many control structures that we know from other programming languages can be modeled in Haskell. First, we consider the while loop.

```
x = 1;
while x < 100
do
x = 2*x
od</pre>
```

In general, a while loop consists of the following.

- a state before executing the loop (initial value)
- a condition
- a loop body for changing the state

In Haskell, the while loop looks as follows.

```
while :: (a \rightarrow Bool) \rightarrow (a \rightarrow a) \rightarrow a \rightarrow a
while p f x | p x = while p f (f x)
| otherwise = x
```

For example, the expression while (<100) (2*) 1 evaluates to 128.

Note that this is not a language extension! This control structure is nothing more than a function, a first class citizen.

3.5.5 Functions as Data

What are data structures? From an abstract point of view, data structures are objects with specific operations.

- Constructors (like (:) or [])
- Selectors (like head or tail, and pattern matching)
- Test functions (like null, and pattern matching)
- Conjunctions (like ++)

The important part is the functionality, that is, the interface, not the implementation. Therefore, a data structure corresponds to a set of functions.

As an example, we want to implement arrays with arbitrary elements in Haskell.

The constructs have the following type.

```
emptyArray :: Array a
putIndex :: Array a -> Int -> a -> Array a
```

Now we only need a single selector.

```
getIndex :: Array a -> Int -> a
```

Now we want to implement the interface. We can achieve this simply by not using other data structures, for example, lists or trees, but rather implement the array as a function. The implementation of this approach could look like the following.

The advantage of this implementation is its conceptual clarity because the implementation is the same as the specification. One drawback is the access time that grows with an increasing number of putIndex calls.

3.5.6 Useful Higher Order Functions

One useful higher-order function is the function composition operator (.).

```
(.) :: (b \rightarrow c) \rightarrow (a \rightarrow b) \rightarrow a \rightarrow c
(f . g) x = f (g x)
```

Two more interesting higher-order functions are curry and uncurry. They allow using functions that are defined on tuples with single elements and vice versa.

```
curry :: ((a, b) -> c) -> a -> b -> c
curry f x y = f (x, y)

uncurry :: (a -> b -> c) -> (a, b) -> c
uncurry f (x, y) = f x y
```

Finally we have look at the function const that takes two arguments and returns the first one.

```
const :: a -> b -> a
const x _ = x
```

3.5.7 Higher Order Functions in Imperative Languages

Modern imperative programming languages allow functions as parameters or return values, too. Although partial application – like Haskell offers – is often not supported and the integration is not as seamless, many algorithms can be defined in a more compact way using higher-order functions. Higher-order also allows additional ways of abstraction since functions can not only abstract values but also program behavior.

In the following, we have a look at the usage of functional parameters in Ruby and Java 8. Besides the concrete syntax, we also discuss the problems that arise from the interplay of functions as values and mutable variables, objects or memory cells.

Higher-order functions are part of Ruby's syntax. They are available in the form of "'blocks"'. A block is a sequence of commands that can be parameterized.

The simplest form of blocks are non-parameterized blocks like, for example, loop bodies. Such blocks can have parameters (noted between vertical lines), which makes them equivalent to anonymous functions. Methods and functions can, in addition to normal parameters, also have a block parameter. For some simple examples, we consider the methods each and map for arrays.

```
a = [1,2,3,4,5]
b = a.map do |x| x + 1 end
b.each do |x| puts x end
```

The program outputs the numbers from 2 to 6 on the screen by means of puts. ².

The map-Method applies the supplied unary block to every element of the array. The original array is not modified but instead, an array of the same length is created. The each method also applies the supplied unary block to every element of the array but does not create a new one.

To understand how blocks are used, we want to define our own version of map that mutates the supplied array. In order to not get too deep into specific Ruby details at this time, we define the function not as a method but rather as an independent procedure map! ³ that uses the array as an additional parameter.

```
def map!(a)
  for i in 0..a.size-1 do
    a[i] = yield(a[i])
  end
end
a = [1,2,3,4,5]
map!(a) do |x| x + 1 end
a.each do |x| puts x end
```

A block that is supplied to map! is applied with the keyword yield; in the above example it is applied to the argument (a[i]). The procedure map! is then applied to the array a and given the increment function block as an additional parameter. ⁴

Thus, blocks are equivalent to functional parameters. Multiple blocks as parameters or returning a block is problematic. Therefore, Ruby allows transforming a block into a value of the class Proc by using lambda. The result can be used like any other value, that is, as an ordinary parameter or as part of a data structure.

Besides, the class Proc offers the method call that applies the function, or rather –

²Blocks in Ruby can be noted alternatively by using curly brackets, for example $b = a.map \{|x| x+1\}$.

³The bang in the name of methods and functions is a Ruby custom that indicates that the method is mutating, that is, the supplied object is modified. Many methods exist in both versions.

⁴Ruby also offers a predefined method map!.

3 Functional Programming

the block, to parameters. As an example we define the function foldr with an explicit functional parameter instead of a block.

```
def foldr(f,e,a)
  if a == [] then
   e
  else
    f.call(a[0],foldr(f,e,a[1,a.size-1]))
  end
end
```

The sum of the elements of an array can then be calculated as follows.

```
puts foldr(lambda do |x,y| x+y end, 0, [1,2,3,4,5,6,7,8,9,10])
```

Next, we want to define an array of functions. The i-th position of the array is supposed to be the function that adds the value i to its argument. Intuitively, this can be implement like this.

```
a = [0,1,2,3,4,5,6,7,8,9]

for i in 0 .. a.size-1 do  # iterate over array, begin with 0
   a[i] = lambda {|x| x+i } # Write respective increment function to i-th
end  # position

puts a[3].call(70)  # Apply function at index 3 to the value 70
```

When the program is executed, the result is unexpectedly not 73 but 79. This is because variables correspond to memory cells in Ruby which are modified while the program executes. The program does not add the value that i has in the loop body but instead when the function is executed, which is 9 in the example above. Thus, function bodies in imperative languages should not contain mutable variables. This can be achieved by, for example, creating the array of functions by using map!

```
a = [0,1,2,3,4,5,6,7,8,9]
a.map! { |i| lambda {|x| x + i }}
puts a[3].call(70) # Hier erhalten wir nun 73.
```

Now the variable i is no longer a memory cell that can be modified over time. Instead, the variable is a block parameter that is created each time the block is applied, similar to scoping. The program now returns the expected value 73.

Anonymous functions can also be defined in Java, starting with version 8. Using functional parameters by means of anonymous inner classes was possible before but the new lambda notation increases the code readability drastically and adds to the feeling of functional programming. As an example we define the class Higher that offers a non-mutating map function for lists.

```
import java.util.*;
class Higher {
```

```
interface Fun<A,B> {    // define interface for objects of functional type
    B call (A arg);    // that requires a call method
}

static <A,B> List<B> map (Fun<A,B> f, List<A> xs) {
    List<B> ys = new ArrayList<B> (xs.size());
    for (A x : xs) {
        ys.add(f.call(x));    // the interface is used here
    }
    return ys;
}
```

When using the map method, we now can use the lambda notation to define anonymous implementations of the Fun interface, as shown below.

```
public static void main (String[] args) {
  List<Integer> a = Arrays.asList(1,2,3,4,5);
  a = map(x -> x + 1, a);
  System.out.println(a); // Result: [2,3,4,5,6]
}
```

The lambda function $x \rightarrow x + 1^5$ is an elegant alternative to the definition of an anonymous inner class that implements the interface Fun.

We want to find out next, if the problem of using mutable variables within function bodies (as before in Ruby) is present in Java, too. Therefore, we again implement a loop that returns a list of increment functions.

```
public static void main (String[] args) {
  List<Fun<Integer,Integer>> fs = new ArrayList<>(10);
  for (int i = 0; i<10; i++) {
    fs.add(x -> x + i);
  }
  System.out.println(fs.get(3).call(70));
}
```

At compile time, the following error message appears.

```
local variables referenced from a lambda expression must be final or effectively final fs.add(x \rightarrow x + i);
```

Thus Java recognizes that the variable i is modified in the program and therefore is cannot be used in the body of a lambda expression. The simple solution is creating a final copy of i for every loop iteration.

```
public static void main (String[] args) {
   List<Fun<Integer,Integer>> fs = new ArrayList<Fun<Integer,Integer>>();
   for (int i = 0; i<10; i++) {
      final int j = i;
   }
}</pre>
```

⁵Multiple function arguments are separated by commas within parentheses, for example $(x,y) \rightarrow x + y$. Type annotations like int $x \rightarrow x + 1$ are also possible.

```
fs.add(x -> x + j);
}
System.out.println(fs.get(3).call(70));
}
```

Now the program compiles and returns 73 as expected.

Modern imperative programming languages often offer the possibility to use functional parameters and values, as demonstrated in the examples. The resulting code is often shorter and more comprehensible, especially when predefined functions like map and fold can be used for lists or arrays. The resulting way of programming often differs significantly from the familiar imperative way because control structures are less important and data plus function that manipulate data are moved to the foreground, that is, the core of the implemented algorithms.

We have also seen the major pitfall of using higher-order functions in imperative programming languages: mutable variables in function bodies. The problem can often be mitigated by not modifying variables in this context and programming in a more functional way. Especially in Java, using a final copy of a mutable variable is a solution, too.

3.6 Type Classes and Overloading

We consider the function elem that checks whether an element is contained in a list.

```
elem x [] = False
elem x (y:ys) = x == y \mid\mid elem x ys
```

What are possible type of elem? Some examples are listed below.

```
Int -> [Int] -> Bool
Bool -> [Bool] -> Bool
Char -> String -> Bool
```

Unfortunately, "a -> [a] -> Bool" is not a valid type since an arbitrary type a is too general: a also includes functions, for which defining equality is difficult (there is no correct, general definition in Haskell). Thus, we need a way to restrict types to types for which value equality is defined. This can be written in Haskell as follows.

```
elem :: Eq a => a -> [a] -> Bool
```

"Eq a" is called a *type constraint*. By adding parentheses and commas, multiple type constraints can be used.

The class Eq is defined in Haskell like this.

```
class Eq a where
  (==), (/=) :: a -> a -> Bool
```

A *class* contains one or more functions that need to be defined for all instances of the class, that is, types. In case of Eq, the functions are (==) and (/=).

Types can be defined as *instances* of a class by implementing the functions of the class.

Now (==) and (/=) can be used for the type IntTree and, for example, the above function elem can also be used with lists of elements of the type IntTree.

Class instances can also be defined for polymorphic types. However, this might require type constraints when defining the instance, as shown in the next example.

```
data Tree a where
    Empty :: Tree a
    Node :: (Tree a) -> a -> (Tree a) -> (Tree a)

instance Eq a => Eq (Tree a) where
    ...<as above>...
```

Note that infinitely many types become instances of the class Eq this way.

3.6.1 Predefined Functions of a Class

The definition of (/=) looks like the above version for almost all instances. Therefore, a default definition is often useful for class functions. The default definition can be overwritten (and needs to be in some cases).

```
class Eq a where
  (==), (/=) :: a -> a -> Bool
  x1 == x2 = not (x1 /= x2)
  x1 /= x2 = not (x1 == x2)
```

3.6.2 Predefined Classes

For some types it is useful to define a total order on values of the type. This functionality is provided in Haskell by an extension of Eq. .

```
data Ordering where
  LT :: Ordering
  EQ :: Ordering
  GT :: Ordering

class Eq a => Ord a where
  compare :: a -> a -> Ordering
  (<), (<=), (>=), (>) :: a -> a -> Bool
```

```
max, min :: a -> a -> a
... -- predefined implementations
```

A minimal instance definition needs at least compare or (<=).

Other predefined classes are Num, Show and Read.

- Num represents numbers for calculations. ((+) :: Num a => a -> a -> a)
- Show transforms values into strings. (show :: Show a => a -> String)
- Read constructs values from strings. (read :: Read a => String -> a)

Other predefined classes are presented in the lecture "'Funktionale Programmierung"' (functional programming).

Instanzes can be derived automatically (except for Num) for self-defined data types by appending the keyword deriving and a list of type classes to the data type definition.

```
deriving (\kappa_1, ..., \kappa_n)
```

Exercise: Check the types of all functions which were defined in the lecture for the most general type. This is possible by deleting the type signature for self-defined functions; Haskell will then derive the most general type. A few examples are the following.

```
(+) :: Num a => a -> a -> a

nub :: Eq a => [a] -> [a]

qsort :: Ord a => [a] -> [a]
```

3.6.3 The Class Read

The class Show is used to output data as text. To read data, that is, retrieve a value from a string, the string needs to be *parsed*, which is a difficult task. Luckily, there exists a predefined class. Nevertheless, using the class requires some understanding about parsing strings.

We consider the following type definition that defines the type of functions that parse strings to values.

```
type ReadS a = String -> [(a,String)]
```

What is the intention behind the return type of ReadS? The first part of the tuple is the actual result of the parser while the second part is the remainder of the string that has not been parsed yet. For example, when we consider the string "Node Empty 42 Empty", it becomes clear that after reading the initial string Node, a tree needs to follow. In this situation, we would like to receive the remaining unparsed string to use it later.

If the string can be parsed multiple ways, the alternatives are returned as elements of a list. If the supplied string cannot be parsed, the returned value is the empty list.

Using ReadS can look like the following.

```
class Read a where
  readsPrec :: Int -> ReadS a
  readList :: ReadS [a] -- predefined
```

The two functions reads and read are defined as follows.

Thus, reads allows transforming a string into a value while checking whether the input is syntactically correct. On the other hand, read can be used when there is no doubt that the input is syntactically correct.

The evaluation of reads and read expressions looks like this.

```
reads "(3,'a')" :: [((Int,Char),String)]
= [((3,'a'),"")]

reads "(3,'a')" :: [((Int,Int),String)]
= []

read "(3,'a')" :: (Int,Char)
= (3,'a')

read "(3,'a')" :: (Int,Int)
= error: no parse

reads "3,'a')" :: [(Int,String)]
= [(3,",'a'")]
```

3.7 Lazy Evaluation

We consider the following Haskell program.

```
f x = 1

h = h
```

The expression f h can be evaluated in multiple ways. f can be evaluated first, which leads to the result 1, or an attempt is made to evaluate h – which never terminates.

From this example we see that not every computation path terminates, but this depends on the evaluation strategy. We distinguish two excellent *reduction strategies*:

- leftmost-innermost (LI): applicative order (strict functions)
- leftmost-outermost (LO): normal order (non-strict functions)

One advantage of LO reduction is that it is complete. Anything that *can* somehow be computed *will* be computed. However, LO can be inefficient because computations can be doubled.

Can LO still have advantages in practice? Indeed, LO offers the following.

- Avoidance of superfluous (possibly infinite) calculations
- Computing with infinite data structures

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For example, the following function from defines the ascending, infinite list of natural numbers, starting with the number n.

```
from :: Num a => a -> [a]
from n = n : from (n + 1)
```

As another example, we recall the function take.

take 1 (from 1) evaluates to [1] since LO works like this.

```
take 1 (from 1)
= take 1 (1:from 2)
= 1 : take 0 (from 2)
= 1 : []
```

The advantage lies in the separation of control (take 1) and data (from 1).

As a further example, we consider the prime number calculation using the sieve of Eratosthenes. The idea is as follows.

- 1. Consider the list of all numbers greater or equal to 2.
- 2. Remove all multiples of the first (prime) number.
- 3. The first element of the list is a prime number. Return to step 2. proceed the same with the remaining list.

This algorithm can be implement in Haskell as follows.

```
sieve :: [Int] -> [Int]
sieve (p:xs) = p : sieve (filter (\x -> x `mod` p > 0) xs)
primes :: [Int]
primes = sieve (from 2)
```

The argument of sieve is an input list which starts with a prime number and in which all multiples of smaller prime numbers are missing. The result is a list of all prime numbers! Now the expression take 10 primes yields the first ten prime numbers.

```
[2,3,5,7,11,13,17,19,23,29]
```

By using (!!), we can output the tenth prime number directly: primes!!9 evaluates to 29.

Infinite data structures can be used as an alternative to the accumulator technique. We consider the Fibonacci function as an example. To retrieve the n-th Fibonacci number, we create the list of all Fibonacci numbers and look up the n-th element.

```
fibgen :: Int -> Int -> [Int]
fibgen n1 n2 = n1 : fibgen n2 (n1 + n2)
fibs :: [Int]
fibs = fibgen 0 1
```

```
fib :: Int -> Int
fib n = fibs !! n
```

Now fib 10 evaluates to 55.

However, one disadvantage of the LO strategy remains: Calculations can be duplicated. We have another look at the simple function double.

```
double x = x + x
```

If we pass (double 3) as an argument to double, then the evaluation looks like this, according to the LI strategy.

```
double (double 3)
= double (3 + 3)
= double 6
= 6 + 6
= 12
```

According to the LO strategy, the result is instead as follows.

```
double (double 3)
= double 3 + double 3
= (3 + 3) + double 3
= 6 + double 3
= 6 + (3 + 3)
= 6 + 6
```

Because of the obvious inefficiency, no real programming language uses the LO strategy. One optimization of the strategy leads to *lazy evaluation*, where, instead of terms, graphs are being reduced. Variables of the program correspond to pointers to expressions and evaluating an expression updates the value of every variable that points to it. This is called *sharing*. Sharing can also be understood as normalization of the program, where for every subexpression a new variable is introduced. For the above example, this looks like the following.

```
double x = x + x

main = let y = 3

z = double y

in double z
```

The evaluation works as shown in figure 3.3. Black lines indicate a reduction step and blue lines are pointers to expressions.

This strategy was formalized by Launchbury in 1993 [?]. Lazy evaluation is optimal in respect to the length of the evaluation: There are no redundant computations as with LI and no duplicated expressions as with LO, although sometimes a lot of memory is needed.

Lazy evaluation is used in the programming language Haskell, while other languages like ML, Erlang, Scheme and Lisp use the LI strategy.

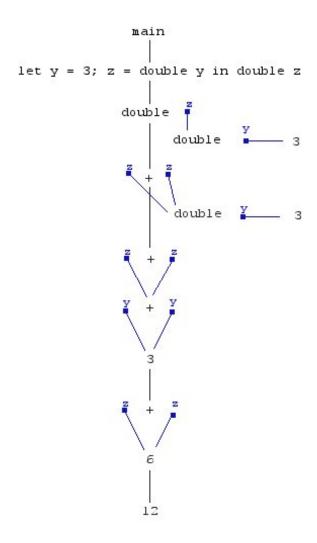


Figure 3.3: Sharing with lazy evaluation

Another advantage of lazy evaluation is that functions can be composed nicely: If we assume the existance of a producer function, for example, of the type $\operatorname{\mathsf{gen}} :: \alpha \to \beta$, and a consumer function, for example, with the type $\operatorname{\mathsf{con}} :: \beta \to \gamma$. If we compose both functions $\operatorname{\mathsf{con}}$. $\operatorname{\mathsf{gen}}$, lazy evaluation does not create large intermediate data structures, but instead, only parts that are needed at the moment are stored in memory and freed as soon as they are no longer needed.

Haskell also allows cyclic data structures like the list

```
ones = 1 : ones
```

The list can be stored in a constant amount of memory, as shown in figure 3.4.

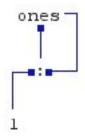


Figure 3.4: Cyclic list ones

3.8 Enumerated Lists

Because infinite structures can often be useful, Haskell has some predefined functions to create infinite lists. For example, repeat yields an infinite list of identical elements.

```
repeat :: a -> [a]
repeat x = x : repeat x
```

Thus, take 70 (repeat '-') yields a textual line.

An infinite list of repeated application of a function can be created with the function iterate.

```
iterate :: (a \rightarrow a) \rightarrow a \rightarrow [a]
iterate f x = x : iterate f (f x)
```

As an exercise, one should think about what iterate (+1) 0 evaluates to.

The example from has shown us that it is simple to create infinite arithmetic sequences in Haskell. We can also define *arithmetic sequences* and *intervals* with the step size 1 or arbitrary step size.

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Since these functions 6 are rather useful, Haskell has a special syntax [n..m] for this functionality.

The following equalities hold.

```
[1..4] == [1,2,3,4]
take 5 [2..] == [2,3,4,5,6]
take 5 [2,4..] == [2,4,6,8,10]
[1,3..10] == [1,3,5,7,9]
take 5 [3,1..] == [3,1,-1,-3,-5]
```

Not only integers but also floating point numbers and characters can be enumerated.

```
> [1,1.5 .. 10]
[1.0,1.5,2.0,2.5,3.0,3.5,4.0,4.5,5.0,5.5,6.0,6.5,7.0,7.5,8.0,8.5,9.0,9.5,10.0]
> ['a'..'z']
"abcdefghijklmnopqrstuvwxyz"
> take 20 ['A' ..]
"ABCDEFGHIJKLMNOPQRST"
```

For this reason, Haskell's Prelude contains the type class Enum.

```
class Enum a where
  succ, pred :: a -> a
  toEnum :: Int -> a
  fromEnum :: a -> Int
  enumFrom :: a -> [a] -- [n..]
  enumFromThen :: a -> a -> [a] -- [n1,n2..]
  enumFromThen :: a -> a -> [a] -- [n..m]
  enumFromThenTo :: a -> a -> a -> [a] -- [n1,n2..m]
```

Therefore, the notation can be used for sequences of all instances of Enum. There are instances, for example, for Int, Float, Char, Bool and Ordering. Therefore, the following holds.

```
> [LT ..]
[LT,EQ,GT]
```

For data types with a finite number of members, the instance only yield finite sequences,

Arithmetic sequences are have many uses. For example, the factorial function can be defined as follows.

```
fac n = foldr (*) 1 [1 .. n]
```

It is also possible to enumerate a list of objects by pairing each element with an index.

```
> zip [1..] "abcde"
[(1,'a'),(2,'b'),(3,'c'),(4,'d'),(5,'e')]
```

The combination with other familiar functions is possible too, of course.

⁶In Haskell, these functions are defined with the prefix enum.

```
> map (uncurry (++)) (zip (map show [1..]) (take 5 (repeat ". Zeile")))
["1. Zeile","2. Zeile","3. Zeile","4. Zeile","5. Zeile"]
```

3.9 List Comprehensions

As we already saw in chapter 3.8 on page 60, Haskell provides some syntactic sugar for list definitions with arithmetic sequences. Furthermore, we can even describe lists with a mathematical notation called *list comprehension*.

```
[(i,j) \mid i \leftarrow [1..3], j \leftarrow [2..4], i \neq j]
```

The above expression evaluates to [(1,2),(1,3),(1,4),(2,3),(2,4),(3,2),(3,4)]. Allowed are generatores like i <- [1..3] and boolean conditions like i /= j. let is also possible.

Thus, the following expression yields all finite initial sequences of natural numbers.

```
[[0..n] \mid n < - [0..]]
```

Finally, we consider another definition of the function concat, which turns a list of lists into a result list containing the lists' entries.

```
concat :: [[a]] -> [a]
concat xss = [y | ys <- xss, y <- ys]</pre>
```

A practical application where list comprehensions are very useful is parsing values.

As an example, we consider CSV files (comma separated values), which we can use to store a list of lists of values in one file.

```
data CSV a where CSV :: [[a]] -> CSV a
```

First we define a Show instance.

```
instance Show a => Show (CSV a) where
  show (CSV xss) = unlines (map separate xss)
  where
    separate [] = ""
  separate [x] = show x
  separate (x:xs) = show x ++ "," ++ separate xs
```

The following is the instance for Read. Here we again use the idea of parsing introduced in the chapter on classes.

```
readsPrec :: Read a => Int -> String -> [(a, String)]
```

The function uses an additional parameter that can be used to note precedents in parentheses. Here we simply ignore it or forward it unchanged to the sub-parser.

Since we do not want to consider the precedent parameter p, we simply pass it on with recursive calls using readsPrec p s. Alternatively, we could have written a shorter reads s, which would result in the initial value being assigned to the precedent parameter.

3.10 Input and Output

Haskell is a purely functional language, that is, functions have no side effects. How can input and output be integrated into such a language?

A first idea: As in other languages (for example ML, Erlang or Scheme), with true side effects.

```
main = let str = getLine
    in putStr str
```

Here getLine should read a line from the keyboard and putStr is supposed to output a string on the standard output. What could be the type of main or putStr?

In Haskell, the smallest type is () (pronounced: unit), whose only value is ()::() (corresponds to void in languages like Java). But if main had the result type (), then no input or output would be necessary to calculate the result () because of lazy evaluation...

Therefore, putStr must output the passed string as a side effect before the result () is returned.

We consider another example.

Here the semicolon ";" is supposed to cause \mathtt{Hi} to appear twice in a row as output. The problem is, however, that due to lazy evaluation \mathtt{x} is calculated only once, which means that the side effect, the output, is executed only once.

Another problem arises from the following, common scenario.

Here the lazy evaluation prevents the desired input sequence. All these problems are solved in Haskell with monads.

3.10.1 IO monad

The input and output is done in Haskell *not* as a side effect. Instead, IO functions provide an *action* for input or output as a result, that is, as a value.

```
putStr "Hi"
```

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For example, the above expression is an action that outputs Hi to the standard output if it is executed.

In principle, an action is a mapping of the following type.

```
World -> (a, World)
```

Here, World describes the present state of the whole "outer world". An action takes the current state of the world and returns a value (for example the read input) and a changed world state. Important is the fact that the world is not directly accessible. Therefore this type is abstractly called "IO a".

Output actions only change the world and return nothing. For this reason they have the type IO (). There are, for example, the following predefined output actions.

```
putStr :: String -> IO ()
putChar :: Char -> IO ()
```

IO actions, like all other functions, are "first class citizens". They can be stored in data structures, for example. An interactive program thus conceptually defines a large IO action, which is applied to the initial world at program start and finally delivers a changed world.

```
main :: IO ()
```

Combining IO actions is achieved with the sequence operator.

```
(>>) :: IO () -> IO () -> IO ()
```

This yields the following equivalences.

An infinite list of output actions is generated by repeat (putStr "Hi").

IO actions can be combined with purely functional calculations as usual. For example, we can output the results of calculations.

```
fac :: Int -> Int
fac n = if n == 0 then 1 else n * fac (n - 1)

main = putStr (show (fac 42))

Or, in a more simple way, like this.
main = print (fac 42)
```

print is a function that first converts a passed value into a string and then writes it to the standard output.

```
print :: Show a => a -> IO ()
print x = putStr (show x) >> putChar '\n'
```

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We consider another example. The following definitions of putStr are equivalent.

```
putStr :: String -> IO ()
putStr "" = return ()
putStr (c:cs) = putChar c >> putStr cs

putStr = foldr (\c -> (putChar c >>)) (return ())
```

return () is here, so to speak, the "empty IO action" which does nothing and only returns its argument.

```
return :: a -> IO a
```

There are corresponding actions for the input of data, where the type of the return value corresponds to the type of the input data.

```
getChar :: IO Char
getLine :: IO String
```

How can the result of an IO action be further used in a subsequent IO action? For we can use the "bind operator".

```
(>>=) :: IO a -> (a -> IO b) -> IO b
```

The bind operator is used as shown in the following example.

```
getChar >>= putChar
```

Here getChar has the type IO Char, putChar has the type Char -> IO (). So getChar >>= putChar has the type IO (). It reads a character and outputs it again.

Next we want to read a whole line.

On a closer look, the string

```
... >>= ...
```

resembles an assignment in an imperative language, except that the left and right sides are swapped. For this reason, a special notation has been introduced for better readability.

3.10.2 do Notation

```
With do { a_1; ...; a_n } or do a_1 \vdots
```

(note the layout rule after the do!) there is an alternative, "imperative" notation.

The expression

is equivalent to the bind expression

```
e_1 >>= \propty e_2
```

and equivalent to the instruction sequence

```
do e_1
```

for the sequence operator

```
e_1 >> e_2
```

This allows us to define the getLine operation as follows.

```
getLine = do c <- getChar
    if c == '\n'
        then return ""
    else do cs <- getLine
        return (c:cs)</pre>
```

3.10.3 Printing Intermediate Results

We want to write a function that calculates the faculty, outputting all intermediate results of the calculation. This is possible if we reformulate the function to an IO action.

Using the function looks like the following.

```
> main
n: 6
(0,1)
(1,1)
(2,2)
(3,6)
(4,24)
(5,120)
Factorial: 720
```

But such programs should be avoided! It is always better to separate input and output on one side from calculations on the other. A common, established scheme is the following.

```
main = do input <- getInput
    let res = computation input
    print res</pre>
```

Here, the line

```
let res = computation input
```

is a purely functional computation. The let does not need an in in a do block.

3.10.4 Reading and Writing Files

The Haskell libraries, based on the monadic IO concept, offer many possibilities to communicate with the environment, for example, reading and writing files, databases or socket connections for network applications. A very easy to use method for reading and writing files is predefined with the following IO actions.

```
readFile :: String \rightarrow IO String \rightarrow reads a file with a given name writeFile :: String \rightarrow String \rightarrow IO () \rightarrow writes a file with given contents
```

For example, we could copy a file like this.

```
readFile "oldfile" >>= writeFile "newfile"
```

Because these operations are performed lazily, even large files can be copied without significant memory consumption.

With the functions already known to us, we can analyze files in a simple way. The size of a file can be computed as follows.

```
> readFile "FILE" >>= print . length
```

We can calculate the number of lines in a file in this way.

```
> readFile "FILE" >>= print . length . lines
```

Here we can see how the function composition and the associated combination style can be applied well. Another example is counting the number of blank lines in a file.

```
> readFile "FILE" >>= print . length . filter (all (==' ')) . lines
```

Here, the predefined function all checks whether all elements of a list meet a given predicate (one has to understand why this definition expresses this exactly).

```
all :: (a \rightarrow Bool) \rightarrow [a] \rightarrow Bool
all p = foldr (&&) True . map p
```

Finally, we want to see how easy it is to output the lines of a file numbered by using higher order functions and infinite data structures. For this we define a function that numbers a text line by line.

If the file FILE contains the content

```
Dies ist
eine Datei
mit drei Zeilen.
```

it is numbered as follows:

```
> readFile "FILE" >>= putStrLn . enumerateLines
1: Dies ist
2: eine Datei
3: mit drei Zeilen.
```

By using some predefined functions, we can still improve the code. For example, since the combination of concat and map is often used, this combination is predefined as follows.

```
-- Maps a function from elements to lists and merges the result into one list. concatMap :: (a -> [b]) -> [a] -> [b] concatMap f = concat . map f
```

This allows us to simplify the definition of enumerateLines as follows.

If we take a closer look at the first function, we see that it merges a list of strings into one string, line by line, by adding line breaks between the individual strings. Since this is often used, this combination is predefined.

```
-- Concatenates a list of strings with terminating newlines. unlines :: [String] -> String unlines ls = concatMap (++"\n") ls
```

This allows us to simplify the definition of enumerateLines once again.

3.11 Modules

Like almost every other programming language, Haskell supports large-scale programming by dividing a program into several modules. Since modules in Haskell are organized similarly to other languages, we will only give a brief overview below.

A module defines a set of names (of functions or data constructors) that can be used when importing this module. You can limit the set of names by export declarations so that, for example, the names of only locally relevant functions are not exported, that is, are not visible on the outside.

The source code of a module therefore begins as follows.

```
module MyProgram (f, g, h) where
```

Here MyProgram is the *module name*, which must be identical to the filename (exception: hierarchical module names, which we will not discuss further here). Thus this module is stored in the file MyProgram.hs. The list of exported names follows in brackets, in this

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case the names f, g and h are exported. Normally the declarations after a where have to be indented further, but in the case of a module they can also start in the first column, so that we can write programs as usual. So our whole module could look like this.

```
module MyProgram (f, g, h) where
f = 42
g = f*f
h = f+g
```

Note that module names must always begin with a capital letter!

It is possible to deviate from this general scheme as follows.

- The export list (i.e. "(f, g, h)") can also be missing; in this case all names defined in the module will be exported.
- Type constructors can also be listed in the export list. In this case, the type constructor name is also exported, but not the corresponding data constructors. If these are also to be exported, then one must write "(..)" after type constructor names in the export list, as shown in the following example.

• If no module header is specified in the source file, the system implicitly uses the module header

```
module Main(main) where
```

We have used an interactive Haskell system like ghci to test small programs before. It is also possible to create an executable file from a Haskell program using the ghc. For this, the program (or the main part of the program) must be stored in the module Main, which in turn must contain a main function, that is, an IO action that is executed as the main program. ⁷

```
main :: IO ()
```

Then one can generate an executable machine program myexec with the following command, for example.

```
ghc -o myexec Main.hs
```

⁷This is how an IO action is associated with a world state!

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Typically, the main program uses a number of other modules that can be imported using an import declaration. For example, our main program could look like this.

```
module Main where
import Nats

main = print (add (S Z) (S Z))

If we execute the command
  ghc -o myexec Main.hs

we get an executable file myexec which we can then execute directly.
> ./myexec
  S (S Z)
```

A import declaration makes those names visible in the current module which are exported from the imported module.

There are a number of variants:

• The imported names can be limited by an enumeration. For example, the following import does not include the name mult.

```
import Nats (Nat(..),add)
```

 One can also import all names except for a few exceptions by means of a hidingconstraint.

```
import Nats hiding (mult)
```

 The standard module Prelude is always implicitly imported if it is not explicitly specified, for example in the following program the functions map and foldr are not imported.

```
import Prelude hiding (map,foldr)
```

• To avoid ambiguities, you can also access imported names within a module in a qualified manner, for example like this.

```
module Main where
import Nats
main = Prelude.print (Nats.add (S Z) (S Z))
```

• If one wants to force that imported names must always be accessed in a qualified way, this can indicated by the restriction qualified.

```
module Main where
import qualified Nats
main = print (Nats.add (Nats.S Nats.Z) (Nats.S Nats.Z))
```

• If the names are too long, the module can be renamed during import.

```
module Main where
```

```
import qualified Nats as N
main = print (N.add (N.S N.Z) (N.S N.Z))
```

In addition, there are further possibilities and rules for the use of modules, which cannot all be described here. For this you should consult Haskell's language definition.

3.12 Data Abstraction and Abstract Data Types

As we have seen from the example of integers, there are different ways to implement a data type. In fact, a central idea of programming with data is

data abstraction: It is not important how the data is represented internally, it is only important how the data is used, that is, which operations are available for this purpose.

For example, in higher programming languages it is not interesting in which bit order integer values are represented internally. What is important is that the addition and multiplication works "as usual". The same applies to other data. For example, in chapter 3.5.5 we have seen that we can implement fields differently than we know from imperative programming languages. Nevertheless, the field operations provide the correct results. Since data abstraction is an important programming technique (in all programming languages!), we will explain this in more detail below.

Abstraction from the internal structure or representation of the data allows using it without knowing the implementation. Instead, one only needs to know the interface, sometimes called API (Application Programming Interface), that is, a set of operations that can be used to work with the data. To give the interface more structure, the interface operations are classified as follows.

Constructors: Operations for constructing data

Selectors: Operations for extracting partial information from data

Operators: Operations to combine data

As a simple example, we consider rational numbers, consisting of a numerator and a denominator. A concrete data type can be immediately defined for this.

```
data Rat where
   Rat :: Int -> Int -> Rat
   deriving Eq
```

For a "natural" output, we define our own show instance.

```
instance Show Rat where
  show (Rat n d) = show n ++ "/" ++ show d
```

A constructor for rational numbers is Rat. But if we want to hide the actual implementation so that we can change it later, then it is better to use a self-defined constructor operation instead of this constructor:

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```
rat :: Int -> Int -> Rat
rat n d = Rat n d
```

In addition, we also need *selector* operations if we want to access the components of a rational number:

```
numerator :: Rat -> Int
numerator (Rat n _) = n

denominator :: Rat -> Int
denominator (Rat _ n) = n
```

A *operator* on rational numbers is, for example, an arithmetic combination like addition or multiplication. These are quite easy to define with the school knowledge on fractions.

```
addR :: Rat -> Rat -> Rat
addR (Rat n1 d1) (Rat n2 d2) = rat (n1*d2 + n2*d1) (d1*d2)
mulR :: Rat -> Rat -> Rat
mulR (Rat n1 d1) (Rat n2 d2) = rat (n1*n2) (d1*d2)
```

To prevent the user from looking into the details of the implementation and defining functions that depend on it, we hide them by exporting only the type name Rat. Our module Rat looks like this.

```
module Rat(Rat,rat,numerator,denominator,addR,mulR) where
```

```
data Rat where
   Rat :: Int -> Int -> Rat
   deriving Eq

instance Show Rat where
   show (Rat n d) = show n ++ "/" ++ show d

rat :: Int -> Int -> Rat
rat n d = Rat n d

numerator :: Rat -> Int
numerator (Rat n _) = n

denominator :: Rat -> Int
denominator (Rat _ n) = n

addR :: Rat -> Rat -> Rat
addR (Rat n1 d1) (Rat n2 d2) = rat (n1*d2 + n2*d1) (d1*d2)

mulR :: Rat -> Rat -> Rat
mulR (Rat n1 d1) (Rat n2 d2) = rat (n1*n2) (d1*d2)
```

We can now use rational numbers. For example, we can define fractions and link them.

```
oneThird :: Rat
oneThird = rat 1 3
```

```
twoThird :: Rat
twoThird = addR oneThird oneThird

If we output the value of twoThird, we get
ghci> twoThird
6/9
```

This is not wrong, but we would have expected the output 2/3. Intuitively, rational numbers should always be reduced. We achieve this by a simple change. When constructing rational numbers, numerators and denominators are always reduced with the largest common divisor. Therefore, we simply change the definition of the constructor.

```
rat :: Int -> Int -> Rat
rat n d = let g = gcd n d in Rat (div n g) (div d g)
ghci> twoThird
2/3
```

However, one disadvantage remains: negative denominators are not uniquely represented. For example, the expression

```
ghci> rat (-3) (-2) -3/-2
```

yields -3/-2 although this is the same value as 3/2. In order to make the representation unique and thus as simple as possible, we avoid the construction of rational numbers with negative denominators with the following redefinition.

Here we already see an important advantage of data abstraction. We have only improved one operation and can continue to use the remaining operations with the same interface. However, if the implementation of a data type is hidden, then how do we know how the operations will behave or what kind of results they produce? This is where another principle of data abstraction comes into play: The behavior of the operations is determined by certain laws which every implementation must meet. This means that we can use different implementations, but still rely on the fact that they work in a certain way. In principle, these laws could be any logical formula, but as a rule, we limit ourselves to equations like "addR x y == addR y x". Such a description of a data type is also called abstract data type.

Definition 3.1 (Abstract data type) An abstract data type (ADT) is a tuple (Σ, X, E) consisting of the following components.

• A program signature Σ (contains operations on the data type),

- a set X of variables that is disjoint with Σ , and
- a set E of equations of the form t = t', where $t, t' \in T_{\Sigma}(X)_s$ are terms of the same signature s.

Let us look at the rational numbers again as an example. The ADT for rational numbers is (Σ, X, E) with

- $\Sigma = (S, F)$ with $S = \{\text{Rat}, \text{Int}\}$
 - $F = \{ \mathtt{rat} :: \mathtt{Int} \ \mathtt{Int} \to \mathtt{Rat}, \mathtt{numerator} :: \mathtt{Rat} \to \mathtt{Int}, \mathtt{denominator} :: \mathtt{Rat} \to \mathtt{Int} \}$
- $X = \{n :: Int, d :: Int\}$
- $E = \left\{ \frac{\text{numerator} \quad (\text{rat } n \ d)}{\text{denominator} \ (\text{rat } n \ d)} = \frac{n}{d} \right\}$

Typically, an ADT has an explicit signature (here: Rat) that describes the elements of the data type (and other signatures used in the ADT operations). Frequently, the use of this ADT signature also reveals the category to which the ADT operations belong.

- Constructors Constructors have the ADT signature as the result signature. (here: Rat).
- Selectors Constructors have the ADT signature as the argument signature. (here: numerator and denominator).
- Operators Constructors have the ADT signature as the result and argument signature (for example "addR :: Rat, Rat → Rat", which is not specified further here.)

It is often the case that selectors and constructors are in a unique relationship, that is, the selector returns exactly the data that was constructed with a constructor. In our case, however, the equations

$$numerator (rat n d) = n$$

and

$$\texttt{denominator} \; (\texttt{rat} \; n \; d) = d$$

would not be correct since we represent fractions in a reduced form. The equation in the ADT therefore specifies that it is irrelevant how rational numbers are represented. What is important is that the ratio of numerator and denominator is always the same.

An *implementation of an ADT* is a program that implements all functions in the program signature so that the equations for all values are always satisfied instead of the variables. Many important data structures can be specified as abstract data types. For example, we have the following ADT components for list structures:

- Constructors: [] und (:)
- Selectors: head und tail

⁸In fact, this ADT also includes the ADT for integers, since we use it. But in the following we leave out ADTs which are already known and which we only use. But formally you would have to import them.

- Operators: (++)
- Equations:

```
\begin{array}{llll} \text{head } (x:xs) & = & x \\ \text{tail } (x:xs) & = & xs \end{array}
```

(and other equations for (++))

Case Study: Set Implementations

To show how data abstraction helps exchange the implementation of data types without changing anything for the user, consider sets as an example. Sets are collections of objects where there is no order and no duplicate elements. For sets we can define many operations. Here we only look at the following functions.

- empty: empty set
- insert x s: insert element x into set s
- isElem x s: is x an element of the set s?
- union s1 s2: union of the sets s1 und s2

Regardless of a concrete implementation, the following laws should always hold.

```
\begin{array}{rcl} & \text{isElem } x \text{ empty} &=& \text{False} \\ & \text{isElem } x \text{ (insert } x \text{ } s) &=& \text{True} \\ & \text{isElem } x \text{ (insert } x \text{ } s)) &=& \text{True} \\ & \text{isElem } x \text{ (union } s_1 \text{ } s_2) &=& \text{isElem } x \text{ } s_1 \text{ || isElem } x \text{ } s_2 \end{array}
```

A first simple and clear reference implementation of sets is obtained by the representation of sets as characteristic functions, as already done in the exercises.

```
module Sets(Set, empty, insert, isElem, union) where

data Set a where
    Set :: (a -> Bool) -> Set a

empty :: Set a
empty = Set (const False)

insert :: Eq a => a -> Set a -> Set a
insert x (Set s) = Set (\y -> x == y || s y)

isElem :: Eq a => a -> Set a -> Bool
isElem x (Set s) = s x

union :: Eq a => Set a -> Set a
union (Set s1) (Set s2) = Set (\x -> s1 x || s2 x)
```

We get an alternative implementation of sets if we represent sets, for example, as lists, in which no duplicates occur. With this representation, the implementation is also quite simple.

```
module SetsByLists(Set, empty, insert, isElem, union) where
import Test.QuickCheck

data Set a where
    Set :: [a] -> Set a
    deriving Show

empty :: Set a
empty = Set []

insert :: Eq a => a -> Set a -> Set a
insert x (Set s) = Set (if x `elem` s then s else x:s)

isElem :: Eq a => a -> Set a -> Bool
isElem x (Set xs) = x `elem` xs

union :: Eq a => Set a -> Set a
union (Set []) s2 = s2
union (Set (x:xs)) s2 = insert x (union (Set xs) s2)
```

Note that this implementation has exactly the same interface as the reference implementation and the ADT. This means that we can replace the set implementation in a program by simply replacing import Sets with import SetsByLists. Here we see an essential advantage of data abstraction: We can exchange the implementation without changing anything in the application programs (except for the name of the imported module).

A slightly more efficient implementation of sets can be achieved by representing sets as ordered lists by making sure that elements are inserted at the correct position in the list.

This has the advantage that we only have to compare on average half of the elements to check whether an element is present in a set.

```
isElem :: (Eq a, Ord a) => a -> Set a -> Bool
isElem x (Set xs) = oelem xs
where
  oelem [] = False
  oelem (y:ys) | x==y = True
```

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```
| x<y = False
| otherwise = oelem ys
```

However, this is not yet an improvement in complexity, that is, the runtime for finding an element is still linear in the number of elements. However, what we can significantly improve is the union operation. While the previous set union was quadratic in the number of elements of both lists (because of the repeated call of insert and thus elem), we can reduce this to a linear runtime for ordered lists by going through both lists at the same time.

Unfortunately, the (sometimes most important) operations <code>insert</code> and <code>isElem</code> are still linear in the number of set elements. We could reduce this to a logarithmic complexity by using balanced search trees. This makes the implementation more complex, but we can test it again with the existing ADT properties, because the interface of the implementation remains the same.

4.1 Motivation

Logic programming has a similar motivation as functional programming.

- Abstraction from the concrete execution of a program on the computer
- Programs as mathematical objects, but here: relations instead of functions
- Computer should find solutions (and not just calculate a value)

Compared to functional programming, logic programming is characterized by the following features.

- Less statements about the direction of calculations/data
- Specification of relationships between objects
- Flexible use of relations

Example: Relationships

As an introductory example, we would like to implement family relationships. The aim is to calculate answers to questions such as

- "Who is Mary's mother?"
- "Which grandfathers does James have?"
- "Is Mary James' aunt?"

and other similar questions.

Our concrete example looks like this:

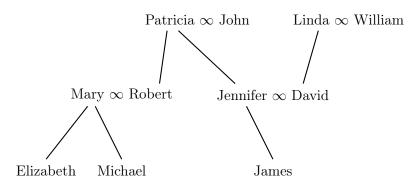
- Patricia is married to John.
- Patricia has two children: Robert and Jennifer.
- Robert is married to Mary.
- Mary has two children: Elizabeth and Michael.
- Linda is married to William.
- Linda has one child: David.
- Jennifer is married to David.
- Jennifer has one child: James.

We represent our example relationships graphically, including the following relationships.

 ∞ : Married

/ : Mother-child relationship

Then our example relationship could look like the following.



In the functional programming language Haskell, for example, these relationships can be modelled as follows: We model people as data types (one could also use String or similar instead).

We define the 'married" relationship as a function.

```
husband :: Person -> Person
husband Patricia = John
husband Linda = William
husband Mary = Robert
husband Jennifer = David
```

We define the 'mother-child' relationship as a function, too.

```
mother :: Person -> Person
mother
         Robert
                   = Patricia
mother
         Jennifer = Patricia
                   = Linda
mother
         David
mother
         Elizabeth = Mary
mother
         Michael
                   = Mary
mother
         James
                   = Jennifer
```

From these basic relationships we can derive further general relationships: The father is the husband of the mother (in a strictly Catholic relationship!).

```
father :: Person -> Person
father kind = husband (mother kind)
```

The grandchild-grandfather relationship is generally a $\it relation$.

Now we can quickly find answers to questions like "'Who is the father of Michael?" or "Is John James' grandfather?".

```
> father Michael
Robert
> grandfather James John
True
```

However, our program cannot answer the following questions directly.

- 1. What children does Robert have?
- 2. What grandfathers does James have?
- 3. Which grandchildren does John have?

This would be possible if variables in expressions were allowed.

- 1. father $k == Robert \rightarrow k = Elizabeth or <math>k = Michael$
- 2. grandfather James $g \rightarrow g = John \text{ or } g = William$
- 3. grandfather e John \rightsquigarrow e = Elizabeth or e = Michael oder e = James

This is allowed in logic languages like Prolog. They have the following characteristics.

- "Free" ("logical") variables in expressions (and rules) are allowed.
- Calculation of *solutions*, that is, values, for free variables, so that the expression is calculable ("provable").
- Problem: How to constructively find the solutions? \rightarrow later
- Calculation principle: draw conclusions from given knowledge.

A *Prolog program* is a set of facts and rules for predicates, where predicates are statements about the objects. A *statement* in turn consists of

- 1. the nature of the statement (property): 3 is prime
- 2. the objects involved: 3 is prime.

Statements are defined in Prolog in the standard prefix notation.

```
name(object_1, ..., object_n)
```

for example prime(3) or husband(mary, robert). Please note the following.

- 1. All names (also called *atoms*) are written in lower case.
- 2. The order of objects in a statement is relevant.
- 3. There must be no space before the opening parenthesis.
- 4. There are also other spellings (operators, later).

As facts we call statements that are assumed to be correct. Syntactically, facts are terminated by a period and a whitespace (space, line feed) at the end.

```
husband(patricia, john).
husband(linda, william).
husband(mary, robert).
husband(jennifer, david).
mother(robert,patricia).
```

```
mother(jennifer,patricia).
mother(david,linda).
mother(elizabeth,mary).
mother(michael,mary).
mother(james,jennifer).
```

Facts alone are not sufficient because they correspond to a notebook or a relational database. For more complex problems we need *rules* or *conclusions* to derive new correct statements.

If statement1 and statement2 are correct, then statement3 is correct.

We write this in Prolog as follows.

```
Statement3: - Statement1, Statement2.
```

Here the comma "," stands for the logical and (\land) , ":-" stands for a conclusion arrow (\Leftarrow) .

For example, the rule "The father of Elizabeth is Robert if Robert is the husband of Mary and Mary is the mother of Elizabeth" can be defined as follows in Prolog.

```
father(elizabeth,robert) :-
    husband(mary,robert),
    mother(elizabeth,mary).
```

This rule is of course very special, but Prolog allows a generalization of this rule. Instead of fixed names, we can also specify unknown objects that are also referred to as *variables*. Please note the following.

- Variable names begin with an uppercase letter
- Variables stand for any other object
- Rules or facts with variables represent infinitely many rules

For example, we can use variables to formulate the following general rules for father and grandfather relationships.

Variables in rules have the following meaning: The rules are correct conclusions for all values which can be used instead of the variables (similar to equation definitions in Haskell).

Query

Facts and rules together correspond to knowledge about a problem. After we have entered them, we can query the Prolog system: Queries are statements whose truth content is to be tested.

```
?- father(michael,robert).
yes
?- father(james,robert).
no
```

By means of variables in queries we can now use our knowledge flexibly. Variables in queries have the meaning: For which values instead of variables is the statement correct? The query "Who is the husband of Mary?" can be formulated as follows.

```
?- husband(mary,Man).
Man = robert
```

And the request "Which grandchildren does John have?" like this.

```
?- grandfather(Grandchild, john).
Grandchild = elizabeth
```

Entering a semicolon ";" prompts the Prolog system to look for further solutions.

```
?- grandfather(Grandchild, john).
Grandchild = elizabeth ;
Grandchild = michael ;
Grandchild = james ;
no
```

Here the Prolog system indicates with a "no" that no further solutions were found.

Summary: Concepts of Logic Programming:

A logic program consists essentially of the following components.

- atoms (elementary objects)
- facts (valid statements)
- rules (if-then-statements)
- queries (Is a statement valid?)
- variables (For which values is a statement valid?)

A property or relationship can be defined with both facts and rules. Therefore, facts and rules are also called *clauses* for this property or relationship. Generally, the latter is also called *predicate* (applied to objects: either true or false) or *relation*.

So far Prolog is what can be described with first order predicate logic. There are constants, predicates, variables (functions) and quantification via individual variables (to be precise: logic programming is based on a subset of first order predicate logic). Because first order predicate logic is not decidable, there can be a set of clauses for every powerful proof system where a certain request cannot be answered with yes or no. The same is true for Prolog, that is, Prolog programs may not terminate. This depends on the evaluation strategy, which will be presented later.

4.2 Syntax of Prolog

As in any programming language, objects in Prolog are either elementary (that is *numbers* or *atoms*) or structured. For the exact definition of the syntax we distinguish four categories of character sets.

Uppercase letters: A B ... Z

Lowercase letters: a b ... z

Numerals: 0 1 ... 9

Special characters: + - * / < = > ` \ : . ? @ # \$ & ^ ~

Prolog objects, also called *terms*, are structured as follows.

Numbers are sequences of numerals (or floating point numbers with usual syntax) **Atoms** form indivisible Prolog objects.

- sequence of lowercase letters, uppercase letters, numerals, "_", beginning with a lowercase letter; or
- sequence of special characters; or
- any special character, enclosed in ', for example 'an atom!'; or
- "special atoms" (not arbitrarily usable): , ; ! []

Constants are numerals or atoms.

Structures are created by combining several objects into one. A structure consists of:

- functor (corresponds to constructor)
- components (arbitrary Prolog objects)

An example of a structure: date(1,6,27). Here date denotes the functor, 1, 6 and 27 are the components. Between functor and components must be *no spaces*.

Structures can also be nested.

```
person(james, smith, date(1,6,27))
```

The functor of a structure is relevant here: date(1,6,27) is not the same as time(1,6,27).

Lists

As in Haskell, lists are the most important structuring option for any number of objects. Lists are defined in Prolog inductively as follows.

- Empty list: []
- ullet Structure of the form .(E,L) where E is the first element of the list and L the remainder of the list.

A list with the elements a, b and c could look like this.

```
.(a, .(b, .(c, [])))
```

There is also shorter syntax for lists in Prolog: $[E_1, E_2, ..., E_n]$ stands for a list with the elements $E_1, E_2, ..., E_n$, [E|L] stands for . (E,L).

Therefore, the following lists are equivalent.

```
.(a,.(b,.(c,[])))
[a,b,c]
[a|[b,c]]
[a,b|[c]]
```

Text is represented in Prolog by lists of ASCII values:

The text "Prolog" corresponds to the list [80,114,111,108,111,103].

Operators

Prolog also has *operators*. Thus "The sum of 1 and the product of 3 and 4" can be described by the structure +(1,*(3,4)). But there is also the natural spelling 1+3*4 which describes the same expression.

The operator notation of structures looks like this in Prolog.

- 1. Structures with one component
 - a) Prefix operator: -2 means -(2)
 - b) Postfix operator: 2 fac fac means fac(fac(2))
- 2. Structures with two components
 - a) Infix operators: 2+3 means +(2,3)

With infix operators, the problem of ambiguity arises immediately.

- \bullet 1-2-3 can be interpreted as
 - "-(-(1,2),3)": then "-" is left-associative- "-(1,-(2,3))": then "-" is right-associative
- 12/6+1
 - -+(/(12,6),1): "/" binds stronger than "+" -/(12,+(6,1)): "+" binds stronger then "/"

The Prolog programmer can define operators themselves. To do this, they must specify associativity and binding strength. The *directive* ":- op(...)" is used for this (you can read more about it in the Prolog manual). The usual mathematical operators like + - * are predefined. Of course it is always possible to set brackets:12/(6+1).

Variables

Variables in Prolog are described by a sequence of letters, numbers and _, beginning with an uppercase letter or _.

- date(1,4,Year) means all first days of April
- [a|L] corresponds to the list with a as the first element
- [A,B|L] means all lists with at least two elements

Variables can occur more than once in a query or clause: Thus [E,E|L] corresponds to all lists with at least two elements, whereby the first two elements are identical.

Prolog offers anonymous variables. _ represents an object whose value is of no interest. Every occurrence of _ stands for another value. As an example, we use our relationship example.

```
isHusband(Person) :- husband(_, Person).
```

The query ?- mother(_,M). queries the Prolog system for all mothers.

Any constant, variable or structure in Prolog is a *term*. A *basic term* is a term without variables.

Programming with List Structures

We now look at an example for programming with list structures. Our goal is to define a predicate member(E,L) that is true if E appears in the list L. We must express our knowledge about the properties of member as facts and rules. An intuitive solution would be to specify many rules for this.

- If E is the first element of L then member (E,L) is true.
- If E is the second element of L then member (E,L) is true.
- if E is the third element of L then member(E,L) is true.
- ...

Since this would lead to infinitely many rules, we think about the following.

member(E,L) is true if E is the first element of L or if it occurs in the tail of L.

We can express this in Prolog as follows.

```
member(E, [E|_]).
member(E, [_|R]) :- member(E, R).
```

Now we can query the Prolog system.

```
?- member(X,[1,2,3]).
X=1 ;
X=2 ;
X=3 ;
no
```

Term Equality

Comparison operators in languages like Java or Haskell, for example ==, always refer to the equality after evaluation the expressions on both sides. In Prolog, however, = denotes the structural equality of terms: Nothing is evaluated!

```
?- 5 = 2+3.
no
?- date(1,4,Year) = date(Day,4,2009).
```

```
Year = 2009
Day = 1
```

4.3 Elementary Programming Techniques

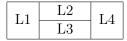
In this chapter we show some basic logical programming techniques.

4.3.1 Enumeration of Search Space

We consider coloring a map with, for example, four countries with the four colors red, yellow, green and blue. We are looking for an arrangement in which adjoining countries have different colors. The four countries are arranged as follows.

- L1 borders L2 and L3.
- L2 borders L1, L3 and L4.
- L3 borders L1, L2 and L4.
- L4 borders an L2 and L3.

Visualization:



What do we know about the problem?

1. There are four colors.

```
color(red).
color(yellow).
color(green).
color(blue).
```

2. Every country has one of these colors.

```
coloring(L1,L2,L3,L4) :- color(L1), color(L2), color(L3), color(L4).
```

3. When are two colors different?

```
different(red,yellow).
different(red,green).
different(red,blue).
...
different(green,blue).
```

4. Correct solution: Bordering countries have different colors.

```
correctColoring(L1,L2,L3,L4) :-
    different(L1,L2),
    different(L1,L3),
    different(L2,L3),
    different(L2,L4),
    different(L3,L4).
```

5. Complete solution of the problem:

```
?- coloring(L1,L2,L3,L4), correctColoring(L1,L2,L3,L4).
L1 = red
L2 = yellow
L3 = green
L4 = red
(more solutions by pressing ";")
```

This example is typical for a situation where one doesn't know how to get a solution in a systematic way. To get a solution with the help of logic programming, we need the following things.

- Generating potential solutions (coloring): We describe the structure of possible solutions.
- Characterization of the correct solutions
- Overall approach:

```
solution(L) :- possibleSolution(L), correctSolution(L).
```

This technique is called *generate-and-test*.

The complexity depends on the set of possible solutions (also called *search space*). In this example, there are $4^4 = 256$ possible solutions. This is acceptable in this case, but sometimes this can also be considerably worse.

Sorting Numbers

To see that the complexity can also become very large, let us consider the next example of sorting numbers, sort(UL,SL). Here UL is a list of numbers and SL a sorted variant of UL.

1. When is a list sorted? That is, what are correct solutions?

If each element is less than or equal to the next.

Expressed in standard Prolog operations on lists (here the predicate x = < y is met if the values of x and y are numbers that are in a less-equal relationship).

```
sorted([]).
sorted([_]).
sorted([E1,E2|L]) :- E1 =< E2, sorted([E2|L]).</pre>
```

2. What are possible solutions?

The sorted list SL is a permutation of UL. A permutation contains the same elements but possibly in a different order. We define permutation by deleting elements:

```
perm([],[]).
perm(L1,[E|R2]) :- remove(E,L1,R1), perm(R1,R2).
```

The definition of remove decides, whether the element that should be removed is the head of the list.

```
\label{eq:remove} $$\operatorname{E}(E, E, R).$$ $\operatorname{remove}(E, A, R).$$ $\operatorname{remove}(E, R, R).$$ :- $\operatorname{remove}(E, R, R).$$
```

3. We get the complete solution according to the technique.

4. This logical specification can be executed.

```
?- sort([3,1,4,2,5],SL).
SL = [1,2,3,4,5]
```

The complexity of this example for a n-element list is in the order of O(n!), because a n-element list has n! possible permutations. This means that for a list with ten elements there are already 3,628,800 possible solutions, so this approach is useless in practice.

In such cases only a more detailed problem analysis (development of better methods, for example, sorting algorithms) can help.

4.3.2 Pattern-oriented Representation of Knowledge

A typical example of a pattern-oriented knowledge representation is list processing. In the case of lists, we differentiate between the cases of empty and non-empty lists. This results in a small, possibly even single-element search space.

As an example we consider the predicate append(L1,L2,L3), which should concatenate two lists L1 and L2 to one list L3.

In more detail, the following should hold.

append(L1,L2,L3)
$$\iff$$
 L1 = $[a_1,...,a_m] \land$ L2 = $[b_1,...,b_n] \land$ L3 = $[a_1,...,a_m,b_1,...,b_n]$

We can define this by case distinction on the structure of the first list L1.

- 1. If L1 is empty, then L3 equals L2.
- 2. If L1 is not empty, then the first element of L3 is equal to the first element of L1 and the remaining list of L3 is the concatenation of the elements of the residual list of L1 and L2.

We can implement this in Prolog as follows.

```
append([],L,L).
append([E|R],L,[E|RL]) :- append(R,L,RL).
```

This predicate is defined *pattern-oriented*. The first clause is only applicable for empty lists L1, the second only for non-empty lists. For a given list L1 only one clause fits.

Examples:

$$\vdash$$
 (1. clause) ?- .

Note:

- The search space has one element.
- The calculation is completely deterministic.
- The processing time is linearly dependent on the input.

4.3.3 Using Relations

Often the problems are of a functional nature: n input values should be assigned to an output value. Mathematical translation:

$$f: M_1 \times \dots \times M_n \to M$$
$$(x_1, \dots, x_n) \mapsto y$$

The implementation of functions is possible in Prolog in the form of relations: the relation $f(X_1, X_2, ..., X_n, Y)$ is fulfilled if Y is the result value of $X_1, ..., X_n$ of f.

However, this relation is defined by clauses, not by a functional expression! This has important consequences, because in Prolog one can use this relation in different ways.

Use as function: x_i are fixed values, and for the result Y we use a variable.

```
?- f(x_1,...,x_n,Y).
Y = y
```

The definition can also be used as an inverse function or relation by specifying the "result value", that is, now y is a fixed value and X_i are variables.

```
?- f(X_1,...,X_n,y).

X_1 = x_1

...

X_n = x_n
```

Consequence: If one programs a function in Prolog, there is also the inverse function or relation available.

The application of the above append as a function looks like this.

```
?- append([1,2],[3,4],L).
L = [1,2,3,4]
```

And as inverse function append can be used as follows.

```
?- append(X,[3,4],[1,2,3,4]).

X = [1,2]

?- append([a], Y, [a,b,c]).

y = [b,c]
```

We can even use it as a reverse relation, for example, to decompose a list.

```
?- append(X,Y,[1,2]).

X = []

Y = [1,2] ;

X = [1]

Y = [2] ;

X = [1,2]

Y = [] ;
```

This flexibility can be used to define new functions and relations.

Adding an element to a list:

We remember the following for logic programming.

- Thinking in relations instead of in functions!
- All parameters are equal (no input/output parameters)!
- Existing predicates are useful. We should pay attention with new predicates in regard to generality or other applications.

4.3.4 Peano Numbers

Another example of logical programming is the *peano* representation of natural numbers, which we have already seen in chapter ??. In the Peano representation, a natural number is defined as follows.

- 0 is a natural number which we represent in Prolog with the functor o.
- If n is a natural number, then s(n) (the successor) is also a natural number. We represent this in Prolog with s(n).

We can define a unary predicate isPeano as follows.

```
isPeano(o).
isPeano(s(N)) :- isPeano(N).
```

On one hand, this can be used to check whether a certain value is a Peano number. In logic programming, however, we can also use it to list all possible Peano numbers.

```
?- isPeano(N).

X = o;

X = s(o);

X = s(s(o));

X = s(s(s(o)))
```

Next, we define arithmetic operations for Peano numbers. First we start with the successor and the partial predecessor function.

```
succ(X, s(X)).
pred(s(X),X).
```

Adding two Peano numbers is also simple and works like concatenating lists.

```
add(o ,Y,Y ) . add(s(X),Y,s(Z)) :- add(X,Y,Z).
```

The addition can also be used vice versa, for example, to determine which numbers can be added together to obtain a given value.

```
?- add(X,Y,s(s(o))).

X = o, Y = s(s(o));

X = s(o), Y = s(o);

X = s(s(o)), Y = o;
```

Now we can define subtraction using addition.

```
sub(X,Y,Z) := add(Y,Z,X).
```

In the next step we define multiplication and the natural order ("smaller or equal").

A few example queries:

```
?- mult(s(s(o)),s(s(o)),V).
V = s(s(s(s(o))));
false
?- leq(s(s(o)),s(o)).
false
?- leq(N,s(s(o))).
N = o;
N = s(o);
N = s(s(o));
false
?- mult(X,Y,s(s(o))).
X = s(o), Y = s(s(o));
X = s(s(o)), Y = s(o);
```

After that the system unfortunately does not terminate because Prolog cannot recognize from the definition of multiplication when it is no longer useful to try further values for X and Y. In the next chapter we look at exactly how this happens and which statements Prolog can solve.

4.4 Computations in Logic Programming

Computations in Prolog essentially correspond to proving statements. But how does Prolog prove statements? To understand this, we first consider a simplified technique: the simple resolution principle.

We know the following elements of logic programming.

- Facts are provable statements. A statement is the application of a predicate to objects, sometimes referred to as a literal.
- If $L:-L_1,...,L_n$. is a rule and the literals $L_1,...,L_n$ are provable, then L is also provable. This rule is called *modus ponens* or *separation rule*.
- Queries are statements to be checked with the following semantics: If ?- $L_1, ..., L_n$. is a query, then it is checked whether $L_1, ..., L_n$ is provable with the given facts and rules.

This leads to the following idea: To check the statement of a query, look for a matching rule and reverse the modus ponens.

Simple Resolution Principle: Reduce the literal proof L to the literal proof $L_1, ..., L_n$, if $L:-L_1, ..., L_n$ is a rule. Here facts are interpreted as rules with an empty body.

We consider the following example.

```
husband(mary, john).
mother(kate, mary).
father(kate, john) :-
    husband(mary, john),
    mother(kate, mary).
```

By means of the simple resolution principle we can carry out the following derivation.

If a query can be reduced in several steps to the empty query by means of the resolution principle, then the request is provable.

Unification

The problem is often that the rules often do not fit "directly": For example, the rule father(K,V):- husband(M,V), mother(K,M).

does not match the query

```
?- father(kate, john).
```

But K and V are variables, so we can use arbitrary objects. If we replace K with kate and V with john, we can apply the rule and thus the resolution principle.

Definition 4.1 (Substitution) A substitution σ is a mapping $\sigma: X \to T_{\Sigma}(X)$ of variables $x:: s \in X$ to terms $t \in T_{\Sigma}(X)_s$.

Applying a substitution σ to a term $t \in T_{\Sigma}(X)$, written as $t\sigma$, is defined inductively as follows.

- If $t :: s \in X$ then $t\sigma = \sigma(t)$.
- If $t :: s \in \Sigma$ (that is, if t is a constant) then $t\sigma = t$.
- If $t = (f \ t_1 \dots t_n)$ then $t\sigma = (f \ t_1 \sigma \dots t_n \sigma)$.

So the substitution only replaces variables – everything else remains unchanged. Usually one demands that a substitution σ changes only finitely many variables, that is, the set

$$\{x \mid x :: s \in X \text{ with } \sigma(x) \neq x\}$$

is finite. Then σ can also be represented as a finite set of pairs of the form

$$\{x \mapsto \sigma(x) \mid x :: s \in X \text{ with } \sigma(x) \neq x\}$$

Substitutions are also used in logic programming, where we note terms in Prolog notation and $\sigma(t)$ for the application of a substitution σ to a term t.

In our example, the substitution σ is defined as follows.

$$\sigma = \{ \mathtt{K} \mapsto \mathtt{kate}, \mathtt{V} \mapsto \mathtt{john} \}$$

Applying this substitution then looks like this.

$$\sigma(\text{father}(K,V)) = \text{father}(\sigma(K),\sigma(V))) = \text{father}(\text{kate,john})$$

In the case of logic programming we have to replace variables not only in rules, but also in queries like

```
?- husband(mary, M).
```

so that we can compute the result of a query.

This is done using *unification*, which describes the process of replacing variables in terms so that the terms become syntactically identical. For the terms date(Day,Month,83)

and date(3,M,J) there are several possible substitutions that make them equal.

$$\sigma_1 = \{ \texttt{Tag} \mapsto 3, \texttt{Monat} \mapsto 4, \texttt{M} \mapsto 4, \texttt{J} \mapsto 83 \}$$

$$\sigma_2 = \{ \texttt{Tag} \mapsto 3, \texttt{Monat} \mapsto \texttt{M}, \texttt{J} \mapsto 83 \}$$

Both σ_1 and σ_2 make the two terms equal, but σ_1 is more special.

Definition 4.2 (Unifier) A substitution σ is called unifier for t_1 and t_2 if $\sigma(t_1) = \sigma(t_2)$. In this case, t_1 and t_2 are called unifiable.

 σ is a most general unifier, MGU, if for all unifiers σ' a substitution ϕ exists with $\sigma' = \phi \circ \sigma$, where the composition $\phi \circ \sigma$ is defined by $\phi \circ \sigma(t) = \phi(\sigma(t))$.

It is important to use MGUs instead of more specific unifiers because it gives us less proof options and therefore less to look for. So the question arises: Are there always MGUs and how can we calculate them?

The answer was found by *Robinson* in 1965 [?]: There are always MGUs for unifiable terms. For the calculation we define the term disagreement set of terms.

Definition 4.3 If t, t' are terms, then the disagreement set) ds(t, t') is defined by.

- 1. If t = t': $ds(t, t') = \emptyset$
- 2. If t or t' are variables and $t \neq t'$: $ds(t, t') = \{t, t'\}$
- 3. If $t = f(t_1, ..., t_n)$ and $t' = g(s_1, ..., s_m)$ $(n, m \ge 0)$:
 - If $f \neq g$ or $m \neq n$: $ds(t, t') = \{t, t'\}$
 - If f = g or m = n and $t_i = s_i$ for all i < k and $t_k \neq s_k$: $ds(t, t') = ds(t_k, s_k)$

Intuitively, this definition means: ds(t,t') contains the partial terms of t and t' at the left innermost position, where t and t' are different.

This directly results in the following unification algorithm.

Unification Algorithm:

Input: terms (literals) t_0, t_1

Output: MGU σ for t_0, t_1 if unifiable, "fail" otherwise

- 1. $k := 0; \sigma_0 := \{\}$
- 2. If $\sigma_k(t_0) = \sigma_k(t_1)$, then σ_k is MGU
- 3. If $ds(\sigma_k(t_0), \sigma_k(t_1)) = \{x, t\}$ with x variable and x does not occur in t, then: σ_{k+1} := $\{x \mapsto t\} \circ \sigma_k$; k := k + 1; go to 2; otherwise: "fail"

We want to try the algorithm with some examples.

- 1. $t_0 = \text{husband(mary,M)}, t_1 = \text{husband(F,john)}$
 - $ds(t_0, t_1) = \{F, mary\}$
 - $\sigma_1 = \{ \mathtt{F} \mapsto \mathtt{mary} \}$
 - $ds(\sigma_1(t_0), \sigma_1(t_1)) = \{M, john\}$
 - $\sigma_2 = \{ \mathtt{M} \mapsto \mathtt{john}, \mathtt{F} \mapsto \mathtt{mary} \}$
 - $ds(\sigma_2(t_0), \sigma_2(t_1)) = \emptyset$
 - $\Longrightarrow \sigma_2 \text{ is MGU}$
- 2. The next example shows how unification can also fail.

$$t_0 = \operatorname{\mathsf{equ}}(\mathtt{f(1),g(X)}), \, t_1 = \operatorname{\mathsf{equ}}(\mathtt{Y,Y})$$

- $ds(t_0, t_1) = \{Y, f(1)\}$
- $\sigma_1 = \{ Y \mapsto f(1) \}$
- $ds(\sigma_1(t_0), \sigma_1(t_1)) = \{g(X), f(1)\}$
- \Longrightarrow not unifiable
- 3. A last example shows the reasoning for checking whether x does not occur in t in step 3.

$$t_0 = X, t_1 = f(X)$$

- $ds(t_0, t_1) = \{X, f(X)\}$
- \implies not unifiable, because X occurs in f(X)!

The check in step 3 of the algorithm is also called *occur check*) and is relevant to its correctness. Many Prolog systems do not use this test for reasons of efficiency, because it is rarely successful, that is, it does not happen with most practical programs that a unification fails due to the occurrence test. Theoretically this can lead to an incorrect unification and to the generation of cyclic terms.

The following theorem applies to the unification algorithm:

Theorem 4.1 (Robinson's Unification Theorem [?]) Let t_0, t_1 be terms. If these are unifiable, then the above algorithm outputs an MGU for t_0, t_1 . If they are not, then it outputs "fail".

Proof: Termination:

- 1. A loop pass only occurs if $ds(\sigma_k(t_0), \sigma_k(t_1))$ contains at least one variable.
- 2. One variable is eliminated in each loop pass (that is, in $\sigma_{k+1}(t_i)$, i = 0.1, x no longer occurs).
- 3. Since t_0 and t_1 only contain a finite number of variables, there are only a finite number of loop passes due to 1. and 2.
- ⇒ The algorithm always terminates.

Correctness:

1. We assume that t_0 and t_1 are not unifiable. If the algorithm stops in step 2, then

 t_0 and t_1 are unifiable. Since the algorithm stops in any case and t_0 and t_1 are not unifiable, the algorithm must stop in step 3, that is, "fail" is the output.

2. Let t_0 and t_1 be unifiable and θ any unifier for t_0 and t_1 . We show:

For all $k \geq 0$ there is a substitution γ_k with $\theta = \gamma_k \circ \sigma_k$ and in the k-th run there is no "fail" output.

Due to termination, this means that the output is actually an MGU.

Proof by induction over k:

$$k = 0$$
: Let $\gamma_0 := \theta$. Then $\gamma_0 \circ \sigma_0 = \theta \circ \{\} = \theta$

 $k \Rightarrow k+1$: Induction hypothesis: $\theta = \gamma_k \circ \sigma_k$, that is, γ_k is unifier for $\sigma_k(t_0)$ and $\sigma_k(t_1)$.

Either: $\sigma_k(t_0) = \sigma_k(t_1)$: Then there is no k+1-th loop pass.

Or: $\sigma_k(t_0) \neq \sigma_k(t_1)$: Also

$$\emptyset \neq ds(\sigma_k(t_0), \sigma_k(t_1)) = \{x, t\}$$

and x does not occur in t (in all other cases $\sigma_k(t_0)$ and $\sigma_k(t_1)$ would not be unifiable!). It follows:

- in the (k+1)-th pass the output it not "fail"
- $\sigma_{k+1} = \{x \mapsto t\} \circ \sigma_k$

Now let $\gamma_{k+1} := \gamma_k \setminus \{x \mapsto \gamma_k(x)\}$ (that is, remove the substitution for x from γ_k). We conclude:

```
\begin{array}{l} \gamma_{k+1} \circ \sigma_{k+1} \\ = \gamma_{k+1} \circ \{x \mapsto t\} \circ \sigma_k \\ = \{x \mapsto \gamma_{k+1}(t)\} \circ \gamma_{k+1} \circ \sigma_k & \text{(since } x \text{ in } \gamma_{k+1} \text{ is not replaced)} \\ = \{x \mapsto \gamma_k(t)\} \circ \gamma_{k+1} \circ \sigma_k & \text{(since } x \text{ does not occur in } t) \\ = \gamma_k \circ \sigma_k & (\gamma_k(x) = \gamma_k(t) \text{ and definition of } \gamma_{k+1}) \\ = \theta & \text{(induction hypothesis)} \end{array}
```

This proves the induction statement and thus also the proposition.

We conclude from this: Unifiable terms always have a most general unifier. We briefly consider the **complexity of the unification algorithm**: In the worst case, the algorithm has an exponential runtime with respect to the size of the input terms. This is due to exponentially growing terms.

Let $t_0 = p(x_1, \dots, x_n)$ and $t_1 = p(f(x_0, x_0), f(x_1, x_1), \dots, f(x_{n-1}, x_{n-1}))$ then:

- $\sigma_1 = \{x_1 \mapsto f(x_0, x_0)\}$
- $\sigma_2 = \{x_2 \mapsto f(f(x_0, x_0), f(x_0, x_0))\} \circ \sigma_1$
- . .

 σ_k replaces x_k with a term with $2^k - 1$ f symbols. Thus the occurrence test necessary for σ_n has an exponential runtime.

The following remarks apply to the runtime of the algorithm.

- 1. Even without an occurrence test, the algorithm has an exponential runtime, since the exponentially growing terms have to be built up.
- 2. A better runtime is achieved if the explicit representation of the terms is omitted and, for example, graphs are used. Thus runtime improvements up to linear algorithms are possible (see, for example, [?, ?, ?]).
- 3. An exponential growth of the terms is extremely rare in practice. Therefore the classical algorithm is often sufficient in connection with the "sharing" of variables, that is, variables are not replaced directly by terms but by references to terms (cf. Sharing in Haskell).

General Resolution Principle

The general resolution principle combines resolution and unification and is also called SLD resolution (Linear Resolution with Selection Function for Definite Clauses). A selection function determines which literal is selected from a query in the next proof step. Possible selection rules are, for example, FIRST (always select the first literal) or LAST (always select the last literal).

Definition 4.4 (SLD resolution) Given is a selection rule and the query

?-
$$A_1, \ldots, A_{i-1}, A_i, A_{i+1}, \ldots, A_m$$
.

where the selection rule selects the literal A_i from the query.

If

$$L :- L_1, \ldots, L_n$$
.

is a rule (with new variables, where n = 0 is allowed) and σ is a most common unifier for A_i and L, then the query

?-
$$\sigma(A_1,\ldots,A_{i-1},L_1,\ldots,L_n,A_{i+1},\ldots,A_m)$$
.

can be derived in an SLD resolution step from the query and the rule regarding the selection rule. If G denotes the original query and G' the derived query, then we also note this resolution step in the form $G \vdash_{\sigma} G'$.

As an example we consider once again the equality in Prolog. In Prolog systems the clause

$$=(X,X)$$
.

is predefined, with "=" declared as an infix operator.

Conclusion: The query

?-
$$t_0 = t_1$$
.

is provable if t_0 and t_1 are unifiable.

The requirement that a *rule with new variables* (this is also called *variant* of a rule) must be taken in a resolution step is justified because the variables in a rule stand for arbitrary values and can therefore be chosen arbitrarily without changing the meaning of a rule.

For example, the clause

```
=(X,X).
```

is equal to the clause

$$=(Y,Y)$$
.

Choosing a rule with new variables is sometimes necessary to avoid failures caused by naming conflicts. If we have clause

```
p(X).
```

(which states that the predicate p is provable for every argument) and we try to prove the query

```
?-p(f(X)).
```

then no resolution step would be possible without new rule variables, since the unification of p(X) and p(f(X)) fails due to the occurrence test.

However, if we take the resolution step with the rule variant

```
p(Y).
```

the unification of p(Y) and p(f(X)) succeeds.

Note that renaming rule variables is also necessary to avoid conflicts between query variables and local variables in rules.

For example, if we look at the rule

```
p :- X=a.
```

and the query

$$?- p, X=b.$$

If the local rule variable X is renamed to Y in the first resolution step, then this request is provable.

```
?- p, X=b.

- Y=a, X=b.

- \{Y\mapsto a\}

?- X=b.

- \{X\mapsto b\}
```

If, on the other hand, this renaming does not take place, then the request would not be provable.

```
?- p, X=b.

- X=a, X=b.

- {X→a}
```

```
?- a=b. 
Fail!
```

As a further example of the general resolution principle, we consider the following program.

Evaluation Strategy and SLD tree

So far, we have only shown what individual SLD steps are and how they can be combined to form a successful derivation. However, there may be many different derivations for a query, some successful and some unsuccessful. An exact evaluation strategy should therefore determine how these different derivatives are constructed or searched.

To get an overview of the different SLD steps and SLD derivations for a query, we summarize them in a tree structure, which is also called the *SLD tree*.

Definition 4.5 (SLD tree) Given is a program P and a request G. A SLD tree for G is a tree whose nodes are marked with requests (even empty requests without literals are allowed).

- 1. The root is marked with G.
- 2. If N is a node that is marked with G_0 and G_1, \ldots, G_n are all requests that can be derived from G_0 using a clause from P (and the given selection rule), then N has exactly the children N_1, \ldots, N_n that are marked with G_1, \ldots, G_n , respectively.
- 3. An empty query without literals does not have a child node.

As an example, let us consider the following program (where we number the rules).

- (1) p(X,Z) := q(X,Y), p(Y,Z).
- (2) p(X,X).
- (3) q(a,b).

The SLD tree for this program and the query "?- p(S,b)." is shown in Figure 4.1. From this SLD tree it can be seen that there are three different SLD derivations for the original query.

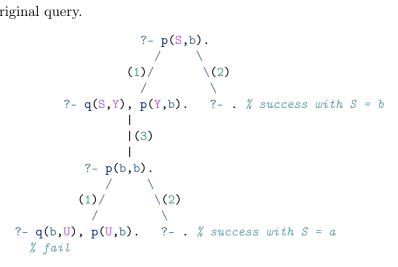


Figure 4.1: SLD tree

An evaluation strategy can thus be interpreted as a rule for searching an SLD tree. A safe strategy, that is, a strategy that always finds a solution if it exists, would be a breadth-first run through the SLD tree. A disadvantage of the breadth-first run is the high memory effort, because we always have to remember the nodes of the respective level. For this reason Prolog does without a secure strategy and uses a more efficient strategy, which is *incomplete*, that is, in some cases it does not find a successful SLD derivation.

Evaluation Strategy of Prolog

The evaluation strategy of Prolog uses the selection strategy FIRST, that is, Prolog always selects the leftmost literal to prove, and passes through the SLD tree with a depth-first pass. Of course, the SLD tree is not actually built, but Prolog uses an evaluation method that ultimately corresponds to a depth-first run (from left to right) in the SLD tree. This is realized by the following backtracking method.

- 1. Clauses are ordered, the order in which they are defined in the program.
- 2. In a resolution step, the *first* appropriate clause for the left literal is picked. At dead ends the last steps are undone and the next alternative is tried (*backtrack*).
- 3. When applying a rule, the variables are replaced by terms by the unification. Then

a variable is bound to a term variable binding also instantiated.

We illustrate the evaluation strategy of Prolog using the following example.

However, the following program shows the problems of the backtracking method, that is, the incompleteness of the evaluation strategy of Prolog. :

```
p :- p.
p.
?- p.
⊢ ?- p.
⊢ ?- p.
```

The system ends up in an infinite loop instead of returning yes. Prolog is therefore incomplete as a "theorem prover".

We consider another example that shows the relevance of the clause order.

```
\begin{split} & \mathsf{last}( [\mathsf{K} | \mathsf{R}] , \ \mathsf{E}) \ :- \ \mathsf{last}( \mathsf{R}, \mathsf{E}) \, . \\ & \\ & ?- \ \mathsf{last}( \mathsf{L}, \ \mathsf{3}) \, . \\ & \qquad \qquad \vdash \ \{ \mathsf{L} \ \mapsto \ [\mathsf{K1} | \mathsf{R1}] \} \\ ?- \ & \mathsf{last}( \mathsf{R1}, \ \mathsf{3}) \, . \\ & \qquad \qquad \vdash \ \{ \mathsf{R1} \ \mapsto \ [\mathsf{K2} | \mathsf{R2}] \} \\ ?- \ & \mathsf{last}( \mathsf{R2}, \ \mathsf{3}) \, . \\ & \dots \end{split}
```

Thus the derivation does not end.

One can therefore remember the following recommendation: clauses for special cases should always be *before* more general clauses! Because if we swap the clauses in the last example, the request will be terminated immediately.

```
?- last(L, 3). \vdash \{L \mapsto [3]\}
```

?- .

Now that we have understood how Prolog tries to prove statements, we can look at more advanced features of Prolog that are very useful for programming in Prolog.

4.5 Negation

The following example shows why negation is often useful.

```
siblings(S, P) :- mother(S, M), mother(P, M).
```

Here, however, a condition like "not S = P" is still missing, otherwise everyone who has a mother would be sibling of themselves.

In Prolog the *negation as failure (NAF)* is implemented: "\+ p" is provable if all proofs for p fail.

```
?- \+ mary = elizabeth.
yes
```

One should note, however, that the negation as a failure does not correspond to the negation from frist-order logic. If we consider the example

```
p :- \ \ \ p.
```

logical negation implies the following.

$$\neg p \Rightarrow p \equiv \neg(\neg p) \lor p \equiv p \lor p \equiv p$$

This means that p is true but a Prolog system gets into an endless loop when trying to prove p.

Clark [?] has therefore given the negation in Prolog a somewhat different meaning (the details of which we skip here) and introduced the NAF rule (negation as finite failure) as the operational principle of negation.

If all proofs for p are finite and fail, then \+ p is provable.

The NAF rule can be effectively implemented by checking the entire SLD tree of the negated predicate and then negating the result (true or false).

However, there is another problem: The negation in Prolog is incorrect if the negated literal contains variables.

```
p(a,a).
p(a,b).
?- \+ p(b,b).
yes
?- \+ p(X,b).
no
```

Logically the answer $\{X \mapsto b\}$ should have been calculated for the last query, but because of the NAF rule variables are never bound in a negation!

The consequence of this is: When proving "+ p", p must not contain any variables! Now let us have another look at our relationship example.

```
siblings(S,P) :-
    mother(S, M),
    mother(P, M),
    \+ S = P.  % okay here because S and P are always bound
```

It is therefore advisable to move negated literals as far to the right of the rule as possible to ensure that the negative literal is free of variables if it is to be proven.

A way out offered in some Prolog systems is delayed negation.

Idea: Delay proof of negative literals until they contain no variables to obtain a logically certain negation.

If negation is realized in this way, we could also write our relationship example in this way.

```
siblings(S,P) :- + S = P, mother(S,M), mother(P,M).
```

A proof could then look as follows.

The delay of the evaluation of literals is theoretically justified, because the selection of literals in logic programming can be arbitrary.

Delay of the evaluation can be implemented quite easily if the Prolog system also contains couroutining, which is the case with many of today's Prolog systems. So you can, for example, in SICStus-Prolog or SWI-Prolog, instead of "\+ p", write the following.

```
when(ground(p),\+ p)
```

This delays the evaluation of "\+ p" until p is variable-free.

4.6 The Cut Operator

With "Cut" (!) we can partially suppress backtracking, that is, conceptually we can "cut off" parts of the SLD tree. This can be useful for different reasons, as listed in the following.

- 1. efficiency (memory and runtime)
- 2. marking functions

- 3. preventing runtime errors (for example with is, later)
- 4. avoiding non-terminating search paths

In Prolog! can be used instead of literals in a rule's body.

```
p :- q, !, r.
```

Operational this means that if the rule above is used to prove p, then the following applies.

- 1. If q is not provable: choose next rule for p.
- 2. If q is provable: p is only provable if r is provable. In other words, no alternative proof for q and no other rule for p is tried.

We consider the following example.

```
yes :- ab(X), !, X = b.
yes.
ab(a).
ab(b).
?- yes.
```

Logically, the query can be proved in two ways. Operational, however, the first rule is applied, X bound to a, followed by a cut, then a failure because X = b cannot be proved, and finally no alternative is tried out. Thus the request is not provable in Prolog. So the cut should be used very carefully!

A cut is often used to distinguish between cases.

```
p :- q, !, r. p :- s.
```

This corresponds (informally) to the following rule.

```
p :- if q then r else s.
```

In fact, there is a special syntax in Prolog for this.

```
p :- q -> r; s.
```

As an example we want to implement the maximum relation.

```
max(X,Y,Z) :- X \ge Y, !, Z = X.

max(X,Y,Z) :- Z = Y. % logically, this is nonsense!
```

Alternatively, we could write it like this.

```
max(X,Y,Z) :- X >= Y -> Z = X ; Z = Y.
```

Furthermore, with the help of the cut we can also define the negation from the last chapter

```
p :- \ q. as the following. p :- q, !, fail.
```

Here fail is a predicate that cannot be proven.

4.7 Arithmetic in Prolog

Since Prolog is a universal programming language, you can also use arithmetic expressions in Prolog. A *arithmetic expression* is a structure with numbers and functors such as +, -, *, / or mod.

For example, the predicate is(X,Y) is predefined, where is is an infix operator. "X is Y" is valid or provable if

- 1. Y is a variable-free arithmetic expression at the time of proof, and
- 2. X = Z holds, if Z is the evaluated value of Y.

Examples:

```
?- 16 is 5 * 3 + 1.

yes

?- X is 5 * 3 + 1.

X = 16

?- 2 + 1 is 2 + 1.
```

4.7.1 Arithmetic Comparison Predicates

With arithmetic comparisons in Prolog both arguments are evaluated before the comparison with is. Prolog offers the following operators for comparisons.

Predicate	Meaning
X =:= Y	value equality
X =\= Y	value inequality
X < Y	less than
X > Y	greater than
X >= Y	greater than or equal
X =< Y	smaller than or equal

A definition of the faculty function in Prolog looks like this.

```
\begin{array}{l} \texttt{fac(0,1).} \\ \texttt{fac(N,F)} := N > 0, \\ & \texttt{N1 is N-1,} \\ & \texttt{fac(N1, F1), \% the order is important!} \\ & \texttt{F is F1 * N.} \end{array}
```

When using arithmetic in Prolog, note that the is predicate is partial: If in "X is Y", Y is not a variable-free arithmetic expression, the calculation is aborted with an error message. Therefore, the order in which is is used is important.

```
?- X = 2, Y is 3 + X. % left-to-right evaluation Y = 5 X = 2 ?- Y is 3 + X, X = 2. ERROR: is/2: Arguments are not sufficiently instantiated
```

The arithmetic in Prolog is therefore logically incomplete.

```
?- 5 is 3 + 2. 
yes 
?- X is 3 + 2. 
X = 5 
?- 5 is 3 + X. 
ERROR: is/2: Arguments are not sufficiently instantiated
```

A global consequence of using this arithmetic is that Prolog programs with arithmetic are often only available in certain modes.¹

4.8 Metaprogramming

Metaprogramming means to change or mix the levels of data and programs during programming. For example, it is possible to interpret data terms as queries or rules or one can summarize results about the structure of calculations as data. Prolog offers many possibilities, some of which are discussed in this chapter.

4.8.1 Higher-Order Predicates

In functional programming we got to know the concept of higher-order functions, that is, functions that process other functions as arguments or deliver results. This made it possible to make programs more compact and reusable. Although Prolog is based on first order logic, Prolog offers similar possibilities as functional programming.

To pass a predicate or predicate call as argument to other predicates and then interpret it as query, Prolog offers a family of call predicates. In the simplest case one can call the predicate call with an argument, whereby the argument term is interpreted and processed as a query.

```
?- call(append([1,2],[3,4],X)).
X = [1, 2, 3, 4]
?- call(is(X,3+4)).
X = 7
```

We can also add more arguments to call, which then has the effect that these arguments are added to the first argument before it is called.

```
?- call(is(X),3+4). X = 7
```

¹The mode of a predicate specifies for which formal parameter which type of current parameter (variable, basic term, . . .) is allowed.

```
?- call(is,X,3+4). X = 7
```

This enables us to define universal predicates, as in functional programming, with which we can make our code more compact. For example, we can define the function map from functional programming as a predicate as follows.

```
map_list(_,[],[]).
map_list(P,[X|Xs],[Y|Ys]) :- call(P,X,Y), map_list(P,Xs,Ys).
```

An example of using this predicate is the following.

```
?- map_list(is, L, [3+4,5+6,10+5]).
L = [7, 11, 15]
```

Since this predicate is quite useful, as we know from functional programming, it is predefined in many Prolog systems as a family of predicates maplist that apply a predicate to one or more lists.²

```
?- maplist(is,[X,Y],[3+4,7+X]).
X = 7
Y = 14
?- maplist(<(0),[0,1,2,3]).
no
?- maplist(<(0),[1,2,3]).
yes</pre>
```

4.8.2 Encapsulation of Non-determinism

In many situations one would not only like to get *one* non-deterministic solution, but identify *all* solutions and continue the computation with these solutions. As an example, we look again at our modeling of family relationships. Here we can, for example, determine all mothers or grandfathers and then use them in the following definitions. For this we can use the meta-predicate findall as follows.

```
mothers(Ms) :- findall(M,mother(_,M),Ms).
grandfathers(Gs) :- findall(G,grandfather(_,G),Gs).
```

codefindall is used as follows: The second argument is a literal for which all solutions should be calculated. In the first example a term, in which mother as a functor has nothing to do with the predicate mother, is evaluated as a predicate when determining all solutions. This is why we speak of metaprogramming, because terms are now interpreted as predicates. The list of calculated solutions is unified with the third argument. Thereby the form of the individual list elements is specified by the first argument of findall, in our example simply the possible assignments of the variables M and G respectively.

An example query yields the following.

```
?- mothers(L).
L = [patricia, patricia, mary, jennifer, mary, linda].
```

²In SWI-Prolog these are directly available, while in SICStus-Prolog you have to import the library lists with ":- use_module(library(lists)).".

```
?- grossvaeter(L).
L = [john, john, william, john].
```

Thus all possible solutions are enumerated and for each solution the assignment of the variables (M or G) is entered into the result list. This becomes even clearer if we also add the children or grandchildren to our list in the program.

```
mothers(Ms) :- findall((E,M),mother(E,M),Ms).
grandfathers(Gs) :- findall((E,G),grandfathers(E,G),Gs).

?- mothers(L).
L = [ (robert, patricia), (jennifer, patricia), (david, linda), (elizabeth, mary), (michael, mary), (james, jnnifer)].

?- grossvaeter(L).
L = [(elizabeth, john), (michael, john), (james, john), (james, william)].
```

So we can only use the predicate findall if the SLD search tree for the predicate in the second argument is finite and all solutions can be effectively determined by Prolog. Similar to the negation, findall would not terminate otherwise.

One often calls findall a capsule, in which the search or the non-determinism is encapsulated and all solutions are calculated. Here one sometimes receives, as seen above, certain solutions several times, since all solutions are written into a list. In the example patricia is a mother of two children and is therefore also listed twice in the list.

However, it is often useful to encapsulate solutions only with regard to certain variables and to keep the non-determinism over the other variables, for which one can use the meta-predicate bagof.

```
mothers(Ms) :- bagof(M,mother(_,M),Ms).
grandfathers(Gs) :- bagof(G,grandfather(_,G),Gs).

?- muetter(L).
L = [jennifer];
L = [patricia];
L = [patricia];
L = [linda];
L = [mary];
L = [mary].

?- grandfathers(L).
L = [john, william];
L = [john];
L = [john].
```

setof works similar to bagof. The only difference, but often important in practice, is that the solution list does not contain duplicate elements and is additionally sorted.

However, one should generally take care to be careful when collecting solutions. This

often means that the declarative character of a Prolog program is lost and one programs rather (non-deterministically) procedurally.

4.9 Difference Lists

Difference lists are not a construct of Prolog, but they were developed within the framework of Prolog and are used, for example, to process natural language sentences efficiently. The motivation to develop difference lists stems from the desire to concatenate lists more

```
append([],L,L).
append([E|R],L,[E|RL]) :- append(R,L,RL).
```

quickly. Let us look at our well-known predicate append.

Since this relation is defined by the structure of the first argument, the runtime is linear to the length of the first list. We could improve the runtime if we had direct access to the end of the first list. For this purpose we represent a list as a difference list, that is, as "difference" of two lists.

As an example, consider the following terms that represent the list [a,b,c].

```
[a,b,c,d] - [d]
[a,b,c] - []
[a,b,c,d,e,f] - [d,e,f]
[a,b,c|L] - L
```

Generally, when using difference lists, we represent a list $[e_1, \ldots, e_n]$ through the pair $[e_1, \ldots, e_n | L]$ -L. The empty list then corresponds to the pair L-L

The advantage is obvious: If M-L is a difference list, then L is the "end" of the list, that is, we have access to the end of the list in constant time!

Thus we can define the concatenation of difference lists by a fact, which is executed in constant time.

```
append_dl(L-M, M-N, L-N).
```

Example: The concatenation of [1,2] and [3,4] is done by a single resolution step.

```
?- append_dl([1,2|L1]-L1, [3,4|L2]-L2, L3). \rightarrow L3 = [1,2,3,4|L2]-L2
```

Runtime improvement with difference lists

Consider reversing the order of all elements in a list as an example. The direct and simple solution is the following.

```
rev([],[]).
rev([E|R],L) :- rev(R,UR), append(UR,[E],L).
```

The runtime is quadratic depending on the length of the first list because for each list element E a concatenation of UR with [E] is necessary.

Since the result list is concatenated by rev, we use difference lists for the second argument of rev.

```
rev_dl([],L-L).
rev_dl([E|R],L-M) :- rev_dl(R,UR-T), append_dl(UR-T,[E|N]-N,L-M).
```

Since append_dl is defined by a simple fact, we calculate it directly and represent the difference list by two arguments.

```
rev_dl([],L,L).
rev_dl([E|R],L,M) :- rev_dl(R,L,[E|M]).
rev(L,M) :- rev_dl(L,M,[]).
```

This gives us a linear runtime for this improved definition!

Difference lists can also be used for many other problems. However, you have to keep in mind that difference lists are not a universal replacement for normal lists, because they can only be concatenated once!

The following example illustrates this.

```
?- DL = [a|L]-L, append_dl(DL,[b|M]-M,X), append_dl(DL,[c|N]-N,Y). \rightarrow no!
```

As a programmer one has to ensure this property.

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