LEDA User Manual

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Introduction

One of the major differences between combinatorial computing and other areas of computing such as statistics, numerical analysis and linear programming is the use of complex data types. Whilst the built-in types, such as integers, reals, vectors, and matrices, usually suffice in the other areas, combinatorial computing relies heavily on types like stacks, queues, dictionaries, sequences, sorted sequences, priority queues, graphs, points, segments, ... In the fall of 1988, we started a project (called **LEDA** for Library of Efficient Data types and Algorithms) to build a small, but growing library of data types and algorithms in a form which allows them to be used by non-experts. We hope that the system will narrow the gap between algorithms research, teaching, and implementation. The main features of LEDA are:

- 1) A clear separation between (abstract) data types and the data structures used to implement them. This distinction is frequently not made in the combinatorial algorithms literature, but is crucial for a library.
- 2) Generic data types: Most of the data types in LEDA have type parameters. For example, a dictionary has a key type K and an information type I and a specific dictionary type is obtained by setting, say, K to int and I to real.
- 3) LEDA is extendible: Users can include own data types either by implementing data structures from scratch in C++ or by combining already existing LEDA data types.
- 4) Ease of use: All data types and algorithms are precompiled C++ modules which can be linked with application programs.

This manual contains the specifications of all data types and algorithms currently available in LEDA. Users should be familiar with the C++ programming language (see [S86] or [L89]). The main concepts and some implementation details of LEDA are described in [MN89]. The manual is structured as follows: In chapter one, which is a prerequisite for all other chapters, we discuss the basic concepts and notations used in LEDA. The other chapters define the data types and algorithms available in LEDA and give examples of their use. These chapters can be consulted independently from one another.

 ϵ

1. Basics

We start with an example. The following program counts the number of occurrences of each string in a sequence of strings

```
\begin{tabular}{ll} \#include < LEDA/d\_array.h> \\ & declare2(d\_array,string,int) \\ & main() \\ & \{ & d\_array(string,int) \ N(0); \\ & string \ s; \\ & while \ (cin >> s \&\& \ s \ != "stop") \ N[s] + +; \\ & for all\_defined(s,N) \ cout << s << " " << N[s] << "\n"; \\ & \} \\ \end{tabular}
```

The program can be compiled using the LEDA library (cf. section 1.9). When executed it reads a sequence of strings from the standard input until the string "stop" is encountered and then prints the number of occurrences of each string on the standard output. More examples of LEDA programs can be found throughout this manual.

The program above uses the data type dictionary array $(d_{-}array)$ from the library. This is expressed by the include statement (cf. section 1.8 for more details). The specification of the data type $d_{-}array$ can be found in section 4.4. We use it also as a running example to discuss the principles underlying LEDA in sections 1.1 to 1.6.

1.1 Specifications

In general the specification of a LEDA data type consists of five parts: a definition of the set of objects comprising the (parameterized) abstract data type, a description of how to derive a concrete data type from a parameterized data type, a description of how to create an object of the data type, the definition of the operations available on the objects of the data type, and finally, informations about the implementation. The five parts appear under the headers definition, type declaration, creation, operations, and implementation respectively.

Definition

This part of the specification defines the objects (also called instances or elements) comprising the data type using standard mathematical concepts and notation.

Example, the generic data type dictionary array:

An object a of type $d_{array}(I, E)$ is an injective function from the data type I to the set of variables of data type E. The types I and E are called the index and the element type respectively, a is called a dictionary array from I to E.

Note that the types I and E are parameters in the definition above. A concrete dictionary array type is declared by a type declaration which we discuss next.

• Type Declaration

This part gives the syntax for deriving specific data types from parameterized or generic data types, i.e., it shows how to set the formal type parameters of a generic data type to concrete data types. For a generic data type XYZ with k type parameters the statement

$$declarek(XYZ,t_1,\ldots,t_k)$$

introduces a new data type with name "XYZ $(t_1, t_2, ..., t_k)$ ", where $t_1, ..., t_k$ are the names of the actual type parameters. (Due to a limitation of the implementation language C++, the number k of type parameters also appears in the name of the declare-macro.) For example,

$declare2(d_array, string, int)$

introduces a new data type with name " $d_array(string, int)$ ". An object of data type $d_array(string, int)$ is a injective mapping from the set of all strings to the set of variables of type int.

Only simple data types are allowed as actual type parameters in declarations of parameterized data types. Simple data types are the C++ built in types char and int, all C++ pointer types, the LEDA types bool, real, string, vector, and matrix (cf. section 2), all basic two-dimensional objects from section 6.1 (point, segment, line, polygon, circle), and all item types (cf. section 1.5). In order to realize generic data types with more complicated subtypes (such as dictionaries, lists, graphs, ...) pointers to these types must be used.

• Creation

A variable of a (previously declared) data type is introduced by a C++ variable declaration. For all LEDA data types variables are initialized at the time of declaration. In many

cases the user has to provide arguments used for the initialization of the variable. In general a declaration

$$XYZ(t_1,\ldots,t_k) \quad y(x_1,\ldots,x_\ell);$$

introduces a variable y of the data type with name "XYZ (t_1, \ldots, t_k) " and uses the arguments x_1, \ldots, x_ℓ to initialize it. For example,

$$d_{-}array(string, int) A(0)$$

introduces A as a dictionary array from strings to integers, and initializes A as follows: an injective function a from string to the set of unused variables of type int is constructed, and is assigned to A. Moreover, all variables in the range of a are initialized to a. The reader may wonder how LEDA handles an array of infinite size. The solution is , of course, that only that part of a is explicitly stored which has been accessed already.

For most data types, in particular for the simple types, the assignment operator is available for variables of that type. However, assignment is in general not a constant time operation, e.g., if s_1 and s_2 are variables of type string then the assignment $s_1 = s_2$ takes time proportional to the length of the value of s_2 .

Remark: For most of the complex data types of LEDA, e.g., dictionaries, lists, and priority queues, it is convenient to interpret a variable name as the name for an object of the data type which evolves over time by means of the operations applied to it. This is appropriate, whenever the operations on a data type only "modify" the values of variables, e.g., it is more natural to say an operation on a dictionary D modifies D than to say that it takes the old value of D, constructs a new dictionary out of it, and assigns the new value to D. Of course, both interpretations are equivalent. From this more object-oriented point of view, a variable declaration, e.g., dictionary(string, int) D, is creating a new dictionary object with name D rather than introducing a new variable of type dictionary(string, int); hence the name "creation" for this part of a specification.

Operations

In this section the operations of the data types are described. For each operation the description consists of two parts

a) The interface of the operation is defined using the C++ function declaration syntax. In this syntax the result type of the operation (void if there is no result) is followed by the operation name and an argument list specifying the type of each argument. For example,

list_item L.insert (E x, list_item it, rel_pos p = after) defines the interface of the insert operation on a list L of elements of type E. (cf. section 3.7). Insert takes as arguments an element x of type E, a list_item it and an optional relative position argument p. It returns a list_item as result.

E& A[I x]

defines the interface of the access operation on a dictionary array A. It takes an element of I as an argument and returns a variable of type E.

b) The effect of the operation is defined. Often the arguments have to fulfill certain preconditions. If such a condition is violated the effect of the operation is not defined. Some, but not all, of these cases result in error messages and abnormal termination of the program (see also section 7.5).

For the insert operation on lists this definition reads:

A new item with contents x is inserted after (if p = after) or before (if p = before) item it into L. The new item is returned. (precondition: item it must be in L)

For the access operation on dictionary arrays the definition reads: returns the variable A(x).

• Implementation

The implementation section lists the data structures used to implement the data type and gives the time bounds for the operations and the space requirement. For example,

Dictionary arrays are implemented by red black trees. Access operations A[x] take time $O(\log dom(A))$. The space requirement is O(dom(A)).

1.2 Arguments

Optional Arguments

The trailing arguments in the argument list of an operation may be optional. If these trailing arguments are missing in a call of an operation the default argument values given in the specification are used. For example, if the relative position argument in the list insert operation is missing it is assumed to have the value after, i.e., L.insert(it, y) will insert the item < y > after item it into L.

Argument Passing

There are two kinds of argument passing in C++, by value and by reference. An argument x of type type specified by "type x" in the argument list of an operation or user defined function will be passed by value, i.e., the operation or function is provided with a copy of x. The syntax for specifying an argument passed by reference is "type& x". In this case the operation or function works directly on x (the variable x is passed not its value).

Passing by reference must always be used if the operation is to change the value of the argument. It should always be used for passing large objects such as lists, arrays, graphs and other LEDA data types to functions. Otherwise a complete copy of the actual argument is made, which takes time proportional to its size, whereas passing by reference always takes constant time.

• Functions as Arguments

Some operations take functions as arguments. For instance the bucket sort operation on lists requires a function which maps the elements of the list into an interval of integers. We use the C++ syntax to define the type of a function argument f:

$$T \quad (*f)(T_1,T_2,\ldots,T_k)$$

declares f to be a function taking k arguments of the data types T_1, \ldots, T_k , respectively, and returning a result of type T, i.e, $f: T_1 \times \ldots \times T_k \longrightarrow T$.

1.3 Overloading

Operation and function names may be overloaded, i.e., there can be different interfaces for the same operation. An example is the translate operations for points (cf. section 6.1).

```
point p.translate(vector v)
point p.translate(real α, real dist)
```

It can either be called with a vector as argument or with two arguments of type real specifying the direction and the distance of the translation.

An important overloaded function is discussed in the next section: Function compare, used to define linear orders for simple data types.

1.4 Linear Orders

Many data types, such as dictionaries, priority queues, and sorted sequences require linearly ordered subtypes. Whenever a type T is used in such a situation, e.g. in declare2(dictionary, T, ...) the function

$$int compare(T\&, T\&)$$

must be declared and must define a linear order on the data type T.

A binary relation rel on a set T is called a linear order on T if for all $x, y, z \in T$:

```
    x rel y
    x rel y and y rel z implies x rel z
    x rel y or y rel x
    x rel y and y rel x implies x = y
```

A function int compare(T&,T&) is said to define the linear order rel on T if

$$compare(x,y)$$
 $\begin{cases} <0, & \text{if } x \ rel \ y \ \text{and } x \neq y \\ =0, & \text{if } x=y \\ >0, & \text{if } y \ rel \ x \ \text{and } x \neq y \end{cases}$

For each of the simple data types char, int, real, string, and point a function compare is predefined and defines the so-called default ordering on that type. The default ordering is the usual \leq - order for char, int, and real, the lexicographic ordering for string, and for point the lexicographic ordering of the cartesian coordinates. For all other simple types T there is no default ordering, and the user has to provide a compare function whenever a linear order on T is required.

Example: Suppose pairs of real numbers shall be used as keys in a dictionary with the lexicographic order of their components. First we declare type pair as the type of pointers to pairs of real numbers, and then we define the lexicographic order on pair by declaring an appropriate compare function.

```
struct Pair {
  real x;
  real y;
};
```

typedef Pair* pair;

```
int compare(pair\&\ p,\ pair\&\ q) { if (p 	o x < q 	o x) return -1; if (p 	o x > q 	o x) return 1; if (p 	o y < q 	o y) return -1; if (p 	o y > q 	o y) return 1; return 0; }
```

Sometimes, a user may need additional linear orders on a simple data type T which are different from the order defined by compare, e.g., he might want to order points in the plane by the lexicographic ordering of their cartesian coordinates and by their polar coordinates. In this example, the former ordering is the default ordering for points. The user can introduce an alternative ordering on the data type point (cf. section 6.1) by defining an appropriate comparing function int cmp(point&, point&) and then declaring the type POINT(cmp) with "declare(POINT, cmp)". POINT is a parameterized data type (cf. section 1.1) with one parameter which must be the name of a comparing function. All data types POINT(cmp) derived from POINT are equivalent to the data type point, with the only exception that if POINT(cmp) is used as an actual parameter in a type declaration, e.g. in "declare2(dictionary, POINT(cmp), ...)", the resulting type (dictionary(POINT(cmp),...)) is based on the linear order defined by cmp. For every simple data type t (except of pointer types) there exists such an equivalent parameterized type T which can be used to define additional linear orders on t by declaring types T(cmp) as described for points. The name of T is always the name of t written with capital letters.

```
In the example, we first declare a function pol\_cmp and the type POINT(pol\_cmp).

int\ pol\_cmp(point\&\ x,\ point\&\ y)
{ //lexicographic ordering on polar coordinates}
}

declare(POINT,\ pol\_cmp)

Now, dictionaries based on either ordering can be defined.

declare2(dictionary,\ point,\ \dots)

declare2(dictionary,\ POINT(pol\_cmp),\ \dots)
```

Remark: We have chosen to associate a fixed linear order with most of the simple types (by predefining the function *compare*). This order is used whenever operations require a linear order on the type, e.g., the operations on a dictionary. Alternatively, we could have required the user to specify a linear order each time he uses a simple type in a situation where an ordering is needed, e.g., a user could define

```
declare3(dictionary,point,lexicographic_ordering,...)
and
declare3(dictionary,point,polar_ordering,...)
```

This alternative would handle the cases where two or more different orderings are needed more elegantly. However, we have chosen the first alternative because of the smaller implementation effort.

1.5 Items

Many of the advanced data types in LEDA (e.g. dictionaries), are defined in terms of so-called items. An item is a container which can hold an object relevant for the data type. For example, in the case of dictionaries a dic_item contains a pair consisting of a key and an information. A general definition of items will be given at the end of this section.

We now discuss the role of items for the dictionary example in some detail. A popular specification of dictionaries defines a dictionary as a partial function from some type K to some other type I, or alternatively, as a set of pairs from $K \times I$, i.e., as the graph of the function. In an implementation each pair (k,i) in the dictionary is stored in some location of the memory. Efficiency dictates that the pair (k,i) cannot only be accessed through the key k but sometimes also through the location where it is stored, e.g., we might want to lookup the information i associated with key k (this involves a search in the data structure), then compute with the value i a new value i', and finally associate the new value with k. This either involves another search in the data structure or, if the lookup returned the location where the pair (k,i) is stored, can be done by direct access. Of course, the second solution is more efficient and we therefore wanted to provide it in LEDA.

In LEDA items play the role of positions or locations in data structures. Thus an

object of type dictionary(K, I), where K and I are types, is defined as a collection of items (type dic_item) where each item contains a pair in $K \times I$. We use $k \in K$ there is at most one $k \in K$ that $k \in K$ there is at most one $k \in K$ that $k \in K$ there is definition may be rephrased as follows: A dictionary $k \in K$ there is at most one $k \in K$ such that the pair $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ that the pair $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ there is at most one $k \in K$ there is at most one $k \in K$ that the pair $k \in K$ there is at most one $k \in K$.

The functionality of the operations

```
dic_item D.lookup(K k)

I D.inf(dic_item it)

void D.change_inf(dic_item it, I i')
```

is now as follows: D.lookup(k) returns an item it with contents (k,i), D.inf(it) extracts i from it, and a new value i' can be associated with k by $D.change_inf(it,i')$.

Let us have a look at the insert operation for dictionaries next:

```
dic_item D.insert(K k, I i)
```

There are two cases to consider. If D contains an item it with contents (k, i') then i' is replaced by i and it is returned. If D contains no such item, then a new item, i.e., an item which is not contained in any dictionary, is added to D, this item is made to contain (k, i) and is returned. In this manual (cf. section 4.3) all of this is abbreviated to

```
dic_item D.insert(K k, I i) associates the information i with the key k.

If there is an item k, i in k then i is replaced by i, else a new item k, i is added to k. In both cases the item is returned.
```

We now turn to a general discussion. With some LEDA types XYZ there is an associated type XYZ_item of items. Nothing is known about the objects of type XYZ_item except that there are infinitely many of them. The only operations available on XYZ_items besides the one defined in the specification of type XYZ is the equality predicate "==" and the assignment operator "=". The objects of type XYZ are defined as sets or sequences of XYZ_items containing objects of some other type Z. In this situation an XYZ_item containing an object $z \in Z$ is denoted by $z \in Z$. A new or unused $z \in Z$ is any $z \in Z$ is not part of any object of type $z \in Z$.

Remark: For some readers it may be useful to interpret a dic_item as a pointer to a variable of type $K \times I$. The differences are that the assignment to the variable contained in a dic_item is restricted, e.g., the K-component cannot be changed, and that in return for this restriction the access to dic_items is more flexible than for ordinary variables, e.g., access through the value of the K-component is possible.

1.6 Input and output

Some parameterized data types (e.g. list) provide the operations print and read for printing their contents to the standard output and for initializing an instance of this type by inserting elements read from the standard input. There are two overloaded functions which can be used for defining input and output functions for user-defined pointer types which are used by read and print operations (and sometimes for error messages).

```
void\ Read(T\&)\ \{\ldots\} defines an input function for objects of type T.
void\ Print(T\&)\ \{\ldots\} defines an output function for objects of type T.
```

Example: We declare the data type list(pair) (see section 3.7) and want to read and print lists of pairs. Note that the Read and Print functions have to be declared before the declaration of the list type.

1.7 Iteration

For many data types LEDA provides iteration macros. These macros can be used to iterate over the elements of lists, sets and dictionaries or the nodes and edges of a graph. Iteration macros can be used similarly to the C++ for statement. Examples are

for all nodes(v, G) { the nodes of G are successively assigned to v}

for all_adj_nodes (w, v) { the neighbor nodes of v are successively assigned to w}

Note: Update operations on an object x are not allowed inside the body of an iteration statement for x.

1.8 Header Files

LEDA data types and algorithms can be used in any C++ program as described in this manual. The specifications (class declarations) are contained in header files. To use a specific data type its header file has to be included into the program. In general the header file for data type XYZ is <LEDA/XYZ.h>. Exceptions are

```
<LEDA/basic.h>
```

This header file contains the declarations for the simple data types bool, real, string (section 2), and the macros and functions described in section 7.

```
<LEDA/graph.h>
```

contains the declarations for graphs and related data types and the declarations of all graph algorithms (section 5).

```
<LEDA/plane.h>
```

contains the two-dimensional objects point, segment, line, polygon, and circle and some basic two-dimensional algorithms (section 6.1).

```
<LEDA/sunview.h>
```

contains a version of the graphic window data type gwindow providing an interface to the SunView window system.

1.9 Libraries

The implementions of all LEDA data types and algorithms are precompiled and contained in three libraries which can be linked with C++ application programs.

a) libL.a

This is the main LEDA library, it contains the implementations of all simple data types (section 2), basic data types (section 3), dictionaries and priority queues (section 4). To compile a program prog.c using any of these data types the libL.a library has to be used like this:

CC prog.c libL.a

b) libG.a

This is the LEDA graph library. It contains the implementations of all graph data types and algorithms (section 5). To compile a program using any graph u data types or algorithms both the libG.a and libL.a library have to be used:

CC prog.c libG.a libL.a

c) libP.a

This is the LEDA library for geometry in the plane. It contains the implementations of all data types and algorithms for two-dimensional geometry (section 6). To compile a program using two-dimensional data types or algorithms all libraries have to be used:

CC prog.c libP.a libG.a libL.a -lm (-lsuntool -lsunwindow -lpixrect)

Note that the libraries must be given in this order, the *suntool*, *sunwindow*, and *pixrect* libraries must be added if a SunView graphic window (cf. section 6.7) is used.

2. Simple Data Types

Simple data types are the C++ built in types char, int, the LEDA data types bool, real, string, vector, matrix, all C++ pointer types and all item types. Simple data types may be used as actual type parameters for generic data types, e.g. dictionary(real, string).

2.1 Boolean Values (bool)

An instance of the data type bool has either the value true or false. The usual C++ logical operators && (and), || (or), ! (negation) are defined for bool.

2.2 Real Numbers (real)

Data type real is the LEDA equivalent of the C++ built in type double. Variables of the data type real behave exactly like variables of type double (arithmetic, compare and input/output operators are the same). The only difference between real and double lies in the fact that real is allowed as subtype (type parameter) for generic data types. There is automatic type conversion from real to double. Thus, all functions taking double arguments accept also arguments of type real and vice versa. In particular the mathematical functions declared in <math.h> can be used with real arguments. The ~operator is defined to explicitly convert an instance of the data type real to a C++ double. This allows the use of the C++ functions printf and form for formatted ouput of reals.

Example:

```
#include <math.h>

real r = 3.1415;

real s = \sin(r);

cout << form("sine of %f = %f\n",~r,~s);
```

2.3 Strings (string)

Data type string is the LEDA equivalent of char* in C++. The differences to the char*-type are that assignment, compare and concatenation operators are defined and that argument passing by value works properly, i.e., there is passed a copy of the string and not only a copy of a pointer. Furthermore a few useful operations for string manipulations are available. The operator converts a string instance to a char*.

1. Creation of a string

- a) string s;
- b) $string \ s(char * c);$

introduces a variable s of type string initialized with the empty string. Variant b) takes as argument a string constant c (char*) and initializes s with c.

2. Operations on a string s

int	$s.\mathrm{length}()$	returns the length of string s
char	$s \ [int \ i]$	returns the character at position i Precondition: $0 \le i \le s.\text{length}()-1$
string	s (int i , int j)	returns the substring of s starting at position i and ending at position j Precondition: $0 \le i \le j \le s.\text{length()-1}$
string	s.tail(int i)	$egin{aligned} ext{returns } s(i, s. ext{length()-1)} \ Precondition: \ 0 \leq i \leq s. ext{length()} \end{aligned}$
string	$s.\mathrm{head}(int\ i)$	$egin{aligned} & ext{returns } s(0, \ i) \ & Precondition: \ 0 \leq i \leq s. ext{length()-1} \end{aligned}$
int	s.pos(string s1)	returns the first position of $s1$ in s if $s1$ is a substring of s , -1 otherwise
string	s.insert(string s1, int i)	$egin{aligned} ext{returns } s. ext{head}(i) + s1 + s. ext{tail}(i+1) \ Precondition: } 0 \leq i \leq s. ext{length}()-1 \end{aligned}$
string	s.replace(string s1, strin	g $s2)$
		returns $s.\text{head}(s.\text{pos}(s1)-1) + s2$ + $s.\text{tail}(s.\text{pos}(s1)+s1.\text{length}())$ Precondition: $s1$ is a substing of s .
char*	~ 8	converts s into a C++ string (char*)
string &	s = s1	assigns the value of s1 to s and returns it

string	s + s1	returns the concatenation of s and $s1$
string&	s + = s1	appends $s1$ to s and returns s
bool	s == s1	returns true iff s and s1 are equal
bool	s! = s1	returns true iff s and s1 are not equal
bool	s < s1	returns true iff s is lexicographically smaller than s1
bool	s > s1	returns true iff s is lexicographically larger than $s1$
bool	$s \ll s1$	returns $(s < s1) \mid\mid (s == s1)$
bool	s >= s1	returns $(s > s1) \mid\mid (s == s1)$
ostream &	$O \ll s$	writes string s to the output stream O
istream &	I>>s	reads string s from the input stream I

3. Implementation

Strings are implemented by C++ character vectors. All operations on a string s take time O(s.length()), except of s[] and s.length() which take constant time.

2.4 Real-Valued Vectors (vector)

An instance of the data type vector is a vector of real variables.

1. Creation of a vector

- a) vector v(int d);
- b) vector v(real a, real b);
- c) vector v(real a, real b, real c);

creates an instance v of type vector; v is initialized to the zero vector of dimension d (variant a), the two-dimensional vector (a, b) (variant b) or the three-dimensional vector (a, b, c) (variant c).

2. Operations on a vector v

int	$v.\dim()$	returns the dimension of v .
real	$v.\mathtt{length}()$	returns the Euclidean length of v
real	$v.{ m angle}(vector \ w)$	returns the angle between v and w .
real &	v $[int i]$	returns i-th component of v . Precondition: $0 \le i \le v.\dim() - 1$.
vector	$v + v_1$	$egin{aligned} ext{Addition} \ ext{\it Precondition:} & v. ext{dim()} = v_1. ext{dim()}. \end{aligned}$
vector	$v - v_1$	$egin{aligned} ext{Subtraction} \ ext{$Precondition:} \ v. ext{dim()} = v_1. ext{dim()}. \end{aligned}$
real	$v * v_1$	$egin{aligned} & ext{Scalar multiplication} \ & ext{$Precondition:} & v. ext{dim()} = v_1. ext{dim()}. \end{aligned}$
vector	v * r	Componentwise multiplication with real r
vector&	$v = v_1$	Componentwise assignment; returns v $Precondition: v.dim() = v_1.dim().$
bool	$v == v_1$	Test for equality
bool	$v != v_1$	Test for inequality
ostream &	O << v	writes v componentwise to the output stream O
istream &	I>>v	reads v componentwise from the input stream I

3. Implementation

Vectors are implemented by arrays of real numbers. All operations on a vector v take time O(v.dim()), except of dim and [] which take constant time. The space requirement is O(v.dim()).

2.5 Real-Valued Matrices (matrix)

An instance of the data type matrix is a matrix of real variables.

1. Creation of a matrix

matrix M(int n, int m);

creates an instance M of type matrix, M is initialized to the $n \times m$ - zero matrix.

2. Operations on a matrix M

int	$M.\mathrm{dim}1()$	returns n , the number of rows of M .
int	$M.\mathrm{dim}2()$	returns m , the number of cols of M .
vector	M.row(int i)	returns the <i>i</i> -th row of M (an m -vector). Precondition: $0 \le i \le n-1$.
vector	M.col(int i)	returns the <i>i</i> -th column of M (an n -vector). Precondition: $0 \le i \le m-1$.
matrix	$M.\mathrm{trans}()$	returns M^T ($m \times n$ - matrix).
real	$M.\mathrm{det}()$	returns the determinant of M . Precondition: M is quadratic.
matrix	M.inv()	returns the inverse matrix of M . Precondition: $M.\det() \neq 0$.
vector	$M.\operatorname{solve}(vector\ b)$	returns vector x with $M \cdot x = b$. Precondition: $M.\dim 1() = M.\dim 2() = b.\dim()$ and $M.\det() \neq 0$.
real &	M (int i, int j)	returns $M_{i,j}$. Precondition: $0 \le i \le n-1$ and $0 \le j \le m-1$.
matrix &	$M = M_1$	Componentwise assignment; returns M . Precondition: $M.\dim 1() = M_1.\dim 1()$ and $M.\dim 2() = M_1.\dim 2()$.

matrix	$M + M_1$	Addition
		Precondition: $M.\dim 1() = M_1.\dim 1()$ and
		$M.\mathrm{dim}2() = M_1.\mathrm{dim}2().$
matrix	$M - M_1$	Subtraktion
		Precondition: $M.\dim 1() = M_1.\dim 1()$ and
		$M.\dim 2() = M_1.\dim 2().$
matrix	$M * M_1$	Multiplication
		Precondition: $M.\dim 2() = M_1.\dim 1()$.
matrix	M * r	Multiplication with real
vector	M * v	Multiplication with vector
		Precondition: M.dim2() = v.dim().
ostream &	O << M	writes matrix M to the output stream O
istream &	I>>M	reads matrix M from the input stream I

3. Implementation

Data type matrix is implemented by two-dimensional arrays of real numbers. All operations take time O(nm), except of det, inv, and solve which take time O(n!), and dim1, dim2, row, and col, which take constant time. The space requirement is O(nm).

3. Basic Data Types

3.1 One Dimensional Arrays (array)

An instance A of the data type array is a mapping from an interval I = [a..b] of integers, called the index set of A, to a set of variables of a data type E, called the element type of A. A(i) is called the element at position i.

1. Declaration of an array type

declare(array, E)

introduces a new data type with name array(E) consisting of all arrays with element type E.

2. Creation of an array

array(E) A(int a, int b);

creates an instance A of type array(E) with index set [a..b].

3. Operations on an array A

A [int i]returns A(i). Precondition: $a \le i \le b$ E&A.low()returns the minimal index a intA.high()returns the maximal index b intvoidA.sort(int (*cmp)(E&, E&)) sorts the elements of A, using function cmpto compare two elements, i.e., if (in_a, \ldots, in_b) and (out_a, \ldots, out_b) denote the values of the variables $(A(a), \ldots, A(b))$ before and after the call of sort, then $cmp(out_i, out_j) \leq 0$ for $i \leq j$ and there is a permutation π of [a..b] such that $out_i = in_{\pi(i)}$ for $a \leq i \leq b$.

intA.binary_search(E x, int (*cmp)(E&, E&))

> performs a binary search for x. Returns iwith A[i] = x if x in A, A.low() -1otherwise. Function cmp is used to compare two elements. Precondition: A must be sorted according to cmp.

4. Implementation

Arrays are implemented by C++ vectors. The access operation takes time O(1), the sorting is realized by quicksort (time $O(n \log n)$) and the binary_search operation takes time $O(\log n)$, where n = b - a + 1. The space requirement is O(|I|).

3.2 Two Dimensional Arrays (array2)

An instance A of the data type array2 is a mapping from a set of pairs $I = [a..b] \times [c..d]$, called the index set of A, to a set of variables of a data type E, called the element type of A, for two fixed intervals of integers [a..b] and [b..c]. A(i,j) is called the element at position (i,j).

1. Declaration of a two dimensional array type

declare(array2, E)

introduces a new data type with name array2(E) consisting of all two-dimensional arrays with element type E.

2. Creation of a two-dimensional array

$$array2(E)$$
 $A(a, b, c, d);$

creates an instance A of type array2(E) with index set $[a..b] \times [c..d]$.

3. Operations on a two-dimensional array A

E&	$A \ (int \ i, \ int \ j)$	returns A(i,j).	
		Precondition: $a \leq i \leq b$ and $c \leq j \leq d$.	
int	$A.\mathrm{low1}()$	$\mathbf{returns} \ a$	
int	$A.{ m high1}()$	$\mathbf{returns} \ b$	
int	$A.\mathrm{low2}()$	returns c	
int	A.high2()	$\mathbf{returns} \ d$	

4. Implementation

Two dimensional arrays are implemented by C++ vectors. All operations take time O(1), the space requirement is O(|I|).

3.3 Stacks (stack)

An instance S of the data type stack is a sequence of elements of a data type E, called the element type of S. Insertions or deletions of elements take place only at one end of the sequence, called the top of S. The size of S is the length of the sequence, a stack of size zero is called the empty stack.

1. Declaration of a stack type

declare(stack, E)

introduces a new data type with name stack(E) consisting all all stacks with element type E.

2. Creation of a stack

stack(E) S;

creates an instance S of type stack(E). S is initialized with the empty stack.

3. Operations on a stack S

$oldsymbol{E}$	S.top()	returns the top element of S Precondition: S is not empty.
$oldsymbol{E}$	$S.\mathtt{pop}()$	deletes and returns the top element of S Precondition: S is not empty.
\boldsymbol{E}	$S.\mathtt{push}(E x)$	adds x as new top element to S .
void	$S.\mathtt{clear}()$	makes S the empty stack.
int	$S.\mathtt{size}()$	returns the size of S .
bool	$S.\mathtt{empty}()$	returns true if S is empty, false otherwise.

4. Implementation

Stacks are implemented by singly linked linear lists. All operations take time O(1), except clear which takes time O(n), where n is the size of the stack.

3.4 Queues (queue)

An instance Q of the data type queue is a sequence of elements of a data type E, called the element type of Q. Elements are inserted at one end (the rear) and deleted at the other end (the front) of Q. The size of Q is the length of the sequence, a queue of size zero is called the empty queue.

1. Declaration of a queue type

declare(queue, E)

introduces a new data type with name queue(E) consisting all all queues with element type E.

2. Creation of a queue

queue(E) Q;

creates an instance Q of type queue(E). Q is initialized with the empty queue.

3. Operations on a queue Q

E	Q.top()	returns the front element of Q Precondition: Q is not empty.
E	Q.pop()	deletes and returns the front element of Q <i>Precondition</i> : Q is not empty.
E	$Q.\mathtt{append}(E x)$	appends x to the rear end of Q .
void	$Q.\mathtt{clear}()$	makes Q the empty queue.
int	$Q.\mathtt{size}()$	returns the size of Q .
bool	$Q.\mathtt{empty}()$	returns true if Q is empty, false otherwise.

4. Implementation

Queues are implemented by singly linked linear lists. All operations take time O(1), except clear which takes time O(n), where n is the size of the queue.

3.5 Bounded Stacks (b_stack)

An instance S of the data type b_stack is a stack (see section 2.3) of bounded size.

1. Declaration of a bounded stack type

 $declare(b_stack, E)$

introduces a new data type with name $b_stack(E)$ consisting all bounded stacks with element type E.

2. Creation of a bounded stack

 $b_{-}stack(E)$ S(n);

creates an instance S of type $b_stack(E)$ that can hold up to n elements. S is initialized with the empty stack.

3. Operations on a b_stack S

\boldsymbol{E}	$S. ext{top}()$	returns the top element of S Precondition: S is not empty.
\boldsymbol{E}	$S.\mathtt{pop}()$	deletes and returns the top element of S <i>Precondition</i> : S is not empty.
\boldsymbol{E}	$S.\mathtt{push}(E x)$	adds x as new top element to S Precondition: S .size() < n .
void	$S.\mathtt{clear}()$	makes S the empty stack.
int	$S.{ m size}()$	returns the size of S .
bool	S.empty()	returns true if S is empty, false otherwise.

4. Implementation

Bounded Stacks are implemented by C++ vectors. All operations take time O(1). The space requirement is O(n).

3.6 Bounded Queues (b_queue)

An instance Q of the data type $b_{-}queue$ is a queue (see section 2.4) of bounded size.

1. Declaration of a bounded queue type

 $declare(b_queue, E)$

introduces a new data type with name $b_queue(E)$ consisting all bounded queue with element type E.

2. Creation of a bounded queue

 $b_{-}queue(E) \quad Q(n);$

creates an instance Q of type $b_{-}queue(E)$ that can hold up to n elements. Q is initialized with the empty queue.

3. Operations on a b_q ueue Q

E	Q.top()	returns the front element of Q Precondition: Q is not empty.
E	$Q.\mathtt{pop}ig(ig)$	deletes and returns the front element of Q <i>Precondition</i> : Q is not empty.
E	$Q.\mathtt{append}(E x)$	appends x to the rear end of Q Precondition: Q .size() < n .
void	$Q.\mathtt{clear}()$	makes Q the empty queue.
int	$Q.\mathrm{size}()$	returns the size of Q .
bool	$Q.\mathtt{empty}()$	returns true if Q is empty, false otherwise.

4. Implementation

Bounded Queues are implemented by circular arrays. All operations take time O(1). The space requirement is O(n).

3.7 Linear Lists (list)

An instance L of the data type list is a sequence of items ($list_item$). Each item in L contains an element of a data type E, called the element type of L. The number of items in L is called the length of L. If L has length zero it is called the empty list. In the sequel < x > is used to denote a list item containing the element x and L[i] is used to denote the contents of list item i in L.

1. Declaration of a list type

declare(list, E)

introduces a new data type with name list(E) consisting of all lists with element type E.

2. Creation of list

list(E) L;

creates an instance L of type list(E) and initializes it to the empty list.

3. Operations on a list L

a) Access Operations

int	$L.{ m length}()$	returns the length of L .
int	$L.{f size}()$	returns L .length().
bool	$L.\mathtt{empty}()$	returns true if L is empty, false otherwise.
$list_item$	$L.\mathrm{first}()$	returns the first item of L .
$list_item$	$L.\mathrm{last}()$	returns the last item of L .
$list_item$	$L.\mathtt{succ}(list_item\ it)$	returns the successor item of item it , nil if $it = L.last()$. Precondition: it is an item in L .
$list_item$	$L.\mathtt{pred}(list_item\ it)$	returns the predecessor item of item it , nil if $it = L.$ first(). Precondition: it is an item in L .
$list_item$	L.cyclic_succ(list_item it)	returns the cyclic successor of item it , i.e., $L.first()$ if $it = L.last()$, $L.succ(it)$ otherwise.
$list_item$	$L. \operatorname{cyclic_pred}(list_item\ it)$	returns the cyclic predecessor of item it, i.e,

		$L.{ m last}() \ { m if} \ it = L.{ m first}(), \ L.{ m pred}(it) \ { m otherwise}.$
$list_item$	$L.\mathrm{search}(E x)$	returns the first item of L that contains x , nil if x is not an element of L
$oldsymbol{E}$	$L. contents(list_item\ it)$	returns the contents $L[it]$ of item it
		Precondition: it is an item in L .
\boldsymbol{E}	$L.inf(list_item\ it)$	${\tt returns} \ L. {\tt contents}(it).$
$oldsymbol{E}$	$L.\mathtt{head}()$	returns the first element of L , i.e. the contents
		of $L.$ first().
		Precondition: L is not empty.
E	L.tail()	returns the last element of L , i.e. the contents
		of $L.last()$.
		Precondition: L is not empty.
int	$L.\mathrm{rank}(E x)$	returns the rank of x in L , i.e. its first
		position in L as an integer from $[1\dots L]$
		(0 if x is not in L).

b) Update Operations

s) opua	o operations		
$list_item$	$ist_item \ L.insert(E \ x, list_item \ it, \ direction \ dir = after)$		
		inserts a new item $\langle x \rangle$ after (if $dir = after$) or before (if $dir = before$) item it into L and returns it. <i>Precondition</i> : it is an item in L .	
$list_item$	$L.\mathrm{push}(E x)$	adds a new item $< x >$ at the front of L and returns it ($L.insert(x, L.first(), before)$)	
$list_item$	$L.\mathtt{append}(E x)$	appends a new item $< x >$ to L and returns it $(L.insert(x, L.last(), after))$	
E	$L. { m del_item}(list_item \ it)$	deletes the item it from L and returns its contents $L[it]$. Precondition: it is an item in L .	
E	$L.\mathtt{pop}()$	deletes the first item from L and returns its contents. Precondition: L is not empty.	
E	L.Pop()	deletes the last item from L and returns its contents. Precondition: L is not empty.	
void	L.assign(list_item it, E x)	makes x the contents of item it . Precondition: it is an item in L .	
void	L.conc(list&~L1)	appends list $L1$ to list L and makes $L1$ the empty list	

void $L.split(list_item\ it, list\&\ L1,\ L2)$

> splits L at item it into lists L1 and L2and makes L the empty list. More precisely, if $L = x_1, ..., x_{k-1}, it, x_{k+1}, ..., x_n$ then

 $L1 = x_1, \ldots, x_{k-1}$ and $L2 = it, x_{k+1}, \ldots, x_n$

Precondition: it is an item in L.

L.apply(void (*f)(E&))for all items $\langle x \rangle$ in L function f is void

called with argument x (passed by reference).

L.sort(int (*cmp)(E&, E&)) sorts the items of L using the ordering defined void

by the compare function $cmp: E \times E \longrightarrow int$,

< 0, if a < b

with cmp(a,b) = 0, if a = b

<0, if a>b

More precisely, if $L = (x_1, ..., x_n)$ before the sort then $L = (x_{\pi(1)}, \ldots, x_{\pi(n)})$ for some permutation π of [1..n] and $cmp(L[x_j], L[x_{j+1}]) \leq 0$ for $1 \le j < n$ after the sort.

L.bucket_sort(int i, int j, int (*f)(E&)) void

> sorts the items of L using bucket sort, where $f: E \longrightarrow int$ with $f(x) \in [i..j]$ for all elements x of L. The sort is stable, i.e., if f(x) = f(y) and $\langle x \rangle$ is before $\langle y \rangle$ in L then $\langle x \rangle$ is before $\langle y \rangle$ after the sort.

the items of L are randomly permuted.

L.permute()void

L.clear()void

makes L the empty list

c) Input and Output Operations

L.read(string s, char delim) void

> Prints string s on the standard output and then reads a sequence of objects of type E terminated by the delimiter delim from the standard input using the overloaded Read function (section 1.5) L is made a list of appropriate length and the sequence is stored in L.

L.read(string s)void

calls $L.read(s, '\n')$.

void

 $L.read(char\ delim)$

calls L.read("", delim).

voidL.read() calls $L.read("", '\n')$.

voidL.print(string s, char space)

Prints the contents of list L to the standard

output using the overload *Print* function to print each element. The elements are separated by the space character *space*. String s is used as a header.

 $void \qquad L.print(string \ s) \qquad calls \ L.print(s, ``).$ $void \qquad L.print(char \ space) \qquad calls \ L.print("", \ space).$ $void \qquad L.print() \qquad calls \ L.print("", ``).$

d) Iterators

Each list L has a special item called the iterator of L. There are operations to read the current value or the contents of this iterator, to move it (setting it to its successor or predecessor) and to test whether the end (head or tail) of the list is reached. If the iterator contains a $list_item \neq nil$ we call this item the position of the iterator. Iterators are used to implement iteration statements on lists.

void L.set_iterator(list_item it) assigns item it to the iterator Precondition: it is in L or it = nil. void L.init_iterator() assigns nil to the iterator

list_item L.get_iterator() returns the current value of the iterator

 $list_item$ L.move_iterator($direction \ dir = forward$)

moves the iterator to its successor (predecessor) if dir = forward (backward) and to the first (last) item if it is undefined (= nil), returns the iterator.

bool L.current_element($E\&\ x$) if the iterator is defined (\neq nil) its contents is assigned to x and true is returned else false is returned

 $L.move_iterator(forward) + return L.current_element(x)$

 $L.prev_element(E\&x)$ $L.move_iterator(backward) +$

 $return L.current_element(x)$

e) Operators

 $egin{array}{lll} & E & L[list_item \ it] & {
m returns} \ L.{
m contents}(it) \ & {
m list}(E)\& & L1 = L2 & {
m assignment:} \end{array}$

The assignment operator makes list L1 a copy of list L2. More precisely if L2 is the sequence of items $x_1, x_2, \ldots x_n$ then L1 is made a sequence of item $y_1, y_2, \ldots y_n$ with $L1[y_i] = L2[x_i]$ for $1 \le i \le n$.

5. Iteration

```
for all list items (it, L) { "the items of L are successively assigned to it" }

for all (x, L) { "the elements of L are successively assigned to x" }
```

6. Implementation

The data type list is realized by doubly linked linear lists. All operations take constant time except for the following operations. Search and rank take linear time O(n), bucket_sort takes time O(n+j-i) and sort takes time $O(n \cdot c \cdot \log n)$ where c is the time complexity of the compare function. n is always the current length of the list.

3.8 Sets (set)

An instance S of the data type set is collection of elements of a linearly ordered type U, called the element type of S. The size of S is the number of elements in S, a set of size zero is called the empty set.

1. Declaration of a set type

declare(set, U)

introduces a new data type with name set(U) consisting of all sets with element type U. Precondition: U is linearly ordered.

2. Creation of a set

set(U) S;

creates an instance S of type set(U) and initializes it to the empty set.

3. Operations on a set S

void	$S.\mathrm{insert}(U oldsymbol{x})$	adds x to S
void	$S.\operatorname{del}(U \boldsymbol{x})$	deletes x from S
bool	$S.\operatorname{member}(U \boldsymbol{x})$	returns true if x in S , false otherwise
U	$S.{ m choose}()$	returns an element of S (error if S is empty) Precondition: S is not empty.
bool	$S.\mathtt{empty}()$	returns true if S is empty, false otherwise
int	$S.\mathrm{size}()$	returns the size of S
void	$S.\mathtt{clear}()$	makes S the empty set

4. Iteration

forall(x, S) { "the elements of S are successively assigned to x" }

5. Implementation

Sets are implemented by red black trees. Operations insert, del, member take time $O(\log n)$, empty, size take time O(1), and clear takes time O(n), where n is the current size of the set.

3.9 Integer Sets (int_set)

An instance S of the data type int_set is a subset of a fixed interval [a..b] of the integers.

1. Creation of an int_set

$$int_set S(a,b);$$

creates an instance S of type int_set for elements from [a..b] and initializes it to the empty set.

2. Operations on a int_set S

void	S.insert(int x)	adds x to S Precondition: $a \le x \le b$.
void	$S.\operatorname{del}(int \ x)$	deletes x from S Precondition: $a \le x \le b$.
bool	S.member(int x)	returns true if x in S , false otherwise Precondition: $a \le x \le b$.
void	$S.\mathtt{clear}()$	makes S the empty set
int_set	S1 = S2	assignment
int_set	$S1 \mid S2$	returns the union of $S1$ and $S2$
int_set	S1 & S2	returns the intersection of $S1$ and $S2$
int_set	~ _S	returns the complement of S

3. Implementation

Integer sets are implemented by bit vectors. Operations insert, delete, member, empty, and size take constant time. Clear, intersection, union and complement take time O(b-a+1).

3.10 Partitions (partition)

An instance of the data type partition consists of a finite set of items (predefined type partition_item) and a partition of this set into blocks.

1. Creation of a partition

partition P;

Creates an instance P of type partition and initializes it to the empty partition.

2. Operations on a partition P

partition_item P.make_block() returns a new partition_item it and adds

the block $\{it\}$ to partition P.

partition_item P.find(partition_item p)

returns a canonical item of the block that

contains item p, i.e., if $P.same_block(p, q)$

then $P.\operatorname{find}(p) = P.\operatorname{find}(q)$.

Precondition: p is an item in P.

bool P.same_block(partition_item p, partition_item q)

returns true if p and q belong to the same

block of partition P.

Precondition: p and q are items in P.

void P.union_blocks(partition_item p, partition_item q)

unites the blocks of partition P containing

items p and q.

Precondition: p and q are items in P.

3. Implementation

Partitions are implemented by the union find algorithm with weighted union and path compression (cf. [T83]). Any sequence of n make_block and $m \ge n$ other operations takes time $O(m\alpha(m,n))$, where α is a functional inverse of Ackerman's function.

4. Example

Spanning Tree Algorithms (cf. graph)

3.11 Dynamic collections of trees (tree_collection)

An instance D of the data type $tree_collection$ is a collection of vertex disjoint rooted trees, each of whose vertices has a real-valued cost and contains an information of type I, called the information type of D.

1. Declaration of a dynamic tree collection type

 $declare(tree_collection, I)$

introduces a new data type with name $tree_collection(I)$ consisting of all dynamic tree collections with information type I.

2. Creation of a tree_collection

 $tree_collection(I)$ D;

creates an instance D of type $tree_collection(I)$, initialized with the empty collection.

3. Operations on a tree_collection D

 d_vertex D.maketree(I x) Adds a new tree to D containing a single

vertex v with cost zero and information x,

and returns v.

I $D.\inf(d_vertex\ v)$ Returns the information of vertex v.

 $d_{-}vertex$ D.findroot $(d_{-}vertex \ v)$ Returns the root of the tree containing v.

 d_vertex D.findcost($d_vertex v, real \& x$)

Sets x to the minimum cost of a vertex on the tree path from v to findroot(v) and returns the last vertex (closest to the root) on this

path of cost x.

void D.addcost($d_vertex\ v, real\ x$)

Adds real number x to the cost of every vertex on the tree path from v to findroot(v).

void $D.link(d_vertex \ v, \ d_vertex \ w)$

Combines the trees containing vertices v and w by adding the edge (v, w). (We regard tree edges as directed from child to parent.)

Precondition: v and w are in different trees

and v is a root.

void $D.cut(d_vertex v)$

Divides the tree containing vertex v into two trees by deleting the edge out of v.

Precondition: v is not a tree root.

4. Implementation

Dynamic collections of trees are implemented by partitioning the trees into vertex disjoint paths and representing each path by a self-adjusting binary tree (see [T83]). All operations take amortized time $O(\log n)$ where n is the number of maketree operations.

4. Priority Queues and Dictionaries

4.1 Priority Queues (priority_queue)

An instance Q of the data type $priority_queue$ is a collection of items (type pq_item). Every item contains a key from a type K and an information from a linearly ordered type I. K is called the key type of Q and I is called the information type of Q. The number of items in Q is called the size of Q. If Q has size zero it is called the empty priority queue. We use < k, i > to denote a pq_item with key k and information i. on I.

1. Declaration of a priority queue type

 $declare2(priority_queue, K, I)$

introduces a new data type with name $priority_queue(K, I)$ consisting of all priority queues with key type K and information type I. Precondition: I is linearly ordered.

2. Creation of a priority queue

 $priority_queue(K, I)$ Q;

creates an instance Q of type $priority_queue(K, I)$ and initializes it with the empty priority queue.

3. Operations on a priority_queue Q

K	$Q.\mathrm{key}(pq_item\ it)$	returns the key of item it . Precondition: it is an item in Q .
I	$Q.inf(pq_item\ it)$	returns the information of item it . Precondition: it is an item in Q .
pq_item	Q.insert(K k, I i)	adds a new item $< k, i >$ to Q and returns it .
pq_item	$Q.{ m find_min}()$	returns an item with minimal information (nil if Q is empty)
void	$Q.\mathrm{del_item}(pq_item~it)$	removes the item it from Q . Precondition: it is an item in Q .
K	$Q.\mathtt{del_min}()$	removes the item with minimal information from Q and returns its key. Precondition: Q is not empty.

pq_item Q.decrease_inf(pq_item it, I i) makes i the new information of item it

Precondition: it is an item in Q and i

is not larger then inf(it).

void Q.change_key(pq_item it, K k) makes k the new key of item it

Precondition: it is an item in Q.

void Q.clear() makes Q the empty priority queue

bool Q.empty() returns true, if Q is empty, false otherwise

int Q.size() returns the size of Q.

4. Implementation

Priority queues are implemented by Fibonacci heaps ([FT84]. Operations insert, del_item, del_min take time $O(\log n)$, find_min, decrease_inf, key, inf, empty take time O(1) and clear takes time O(n), where n is the size of Q. The space requirement is O(n).

5. Example

Dijkstra's Algorithm (cf. section 8.1)

4.2 Bounded Priority Queues (b_priority_queue)

An instance Q of the data type $b_priority_queue$ is a priority_queue (cf. section 4.1) whose items contain informations from a fixed interval [a..b] of integers.

1. Declaration of a bounded priority queue type

 $declare(b_priority_queue, K)$

introduces a new data type with name $b_priority_queue(K)$ consisting of all bounded priority queues with key type K.

2. Creation of a bounded priority queue

 $b_priority_queue(K)$ Q(a,b);

creates an instance Q of type $b_priority_queue(K)$ with information type [a..b] and initializes it with the empty priority queue.

3. Operations on a $b_priority_queue Q$

The operations are the same as for the data type $priority_queue$ with the additional precondition that any information argument must be in the range [a..b].

4. Implementation

Bounded priority queues are implemented by arrays of linear lists. Operations insert, find_min, del_item, decrease_inf, key, inf, and empty take time O(1), del_min (= del_item for the minimal element) takes time O(d), where d is the distance of the minimal element to the next bigger element in the queue (= O(b-a) in the worst case). clear takes time O(b-a+n) and the space requirement is O(b-a+n), where n is the current size of the queue.

4.3 Dictionaries (dictionary)

An instance D of the data type dictionary is a collection of items (dic_item) . Every item in D contains a key from a linearly ordered data type K, called the key type of D, and an information from a data type I, called the information type of D. The number of items in D is called the size of D. A dictionary of size zero is called the empty dictionary. We use k, k to denote an item with key k and information k is said to be the information associated with key k. For each $k \in K$ there is at most one item k, k to k.

1. Declaration of a dictionary type

declare 2 (dictionary, K, I)

introduces a new data type with name dictionary(K, I) consisting of all dictionaries with key type K and information type I. Precondition: K is linearly ordered.

2. Creation of a dictionary

dictionary(K,I) D;

creates an instance D of type dictionary(K, I) and initializes D to the empty dictionary.

3. Operations on a dictionary D

	K	$D.\text{key}(dic_item \ it)$	returns the key of item it. Precondition: it is an item in D.
	I	$D.\inf(dic_item\ it)$	returns the information of item it .
		,	Precondition: it is an item in D.
	dic_item	D.insert(K k, I i)	associates the information i with the key k . If there is an item $< k, j >$ in D then j is replaced by i, else a new item $< k, i >$ is added to D . In both cases the item is returned.
	dic_item	D.lookup(K k)	returns the item with key k (nil if no such item exists in D).
	I	D.access(K k)	returns the information associated with key k Precondition: there is an item with key k in D .
	void	$D.\operatorname{del}(K k)$	deletes the item with key k from D (null operation, if no such item exists).
1	void	$D.\operatorname{del_item}(dic_item\ it)$	removes item it from D .

```
Precondition: it is an item in D.

void D.change_inf(dic_item it, I i) makes i the information of item it.

Precondition: it is an item in D.

void D.clear() makes D the empty dictionary.

bool D.empty() returns true if D is empty, false otherwise.

int D.size() returns the size of D.
```

4. Iteration

for all_dic_items (i, D) { "the items of D are successively assigned to i" }

5. Implementation

Dictionaries are implemented by leaf oriented red black trees. Operations insert, lookup, delitem, del take time $O(\log n)$, key, inf, empty, size, change inf take time O(1), and clear takes time O(n). Here n is the current size of the dictionary. The space requirement is O(n).

6. Example

Using a dictionary to count the number of occurrences of the elements in a sequence of strings, terminated by string "stop".

```
#include <LEDA/dictionary.h>

\begin{aligned}
\text{declare2}(\text{dictionary, string, int}) \\
\text{main()} \\
\{ \\
\text{dictionary(string, int) } D; \\
\text{string } s; \\
\text{dic.item } it; \\
\text{while (} (\text{cin} >> s) && (s != \text{"stop"}) ) \\
& \{ it = D.\text{lookup}(s); \\
& \text{if } (it == nil) \ D.\text{insert}(s, 1); \\
& \text{else } D.\text{change.inf}(it, D.\text{info}(it) + 1); \\
& \} \\
\text{forall_dic_items}(it, D) \text{ cout } << D.\text{info}(it) << \text{" " } << D.\text{key}(it) << \text{"} \n"; \\
}
```

4.4 Dictionary Arrays (d_array)

An instance A of the data type d_{-array} (dictionary array) is an injective mapping from a linearly ordered data type I, called the index type of A, to a set of variables of a data type E, called the element type of A.

1. Declaration of a d_array type

 $declare2(d_array, I, E)$

introduces a new data type with name $d_{-}array(I, E)$ consisting of all d_arrays with index type I and element type E. Precondition: I is linearly ordered.

2. Creation of a d_array

$$d_{-}array(I, E) \quad A(x);$$

creates an injective function a from I to the set of unused variables of type E, assigns x to all variables in the range of a and initializes A with a.

3. Operations on a $d_{array} A$

E& $A \ [I \ x]$ returns the variable A(x)bool A.defined($I \ x$) returns true if $x \in dom(A)$, false otherwise; here dom(A) is the set of all $x \in I$ for which A[x] has already been executed.

4. Iteration

 $forall_defined(x, A)$ { "the elements from dom(A) are successively assigned to x" }

5. Implementation

Dictionary arrays are implemented by red black trees. Access operations A[x] take time $O(\log dom(A))$. The space requirement is O(dom(A)).

6. Example

Program 1: Using a dictionary array to count the number of occurences of the elements in a sequence of strings.

```
\label{eq:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:lemma:
```

Program 2: Using a d_array to realize an english/german dictionary.

```
#include <LEDA/d_array.h>

declare2(d_array,string,string)
main()
{
    d_array(string,string) trans;
    trans["hello"] = "hallo";
    trans["world"] = "Welt";
    trans["book"] = "Buch";
    trans["key"] = "Schluessel";
    string s;
    forall_defined(s, trans) cout << s << " " << trans[s] << "\n";
}</pre>
```

4.5 Hashing arrays (h_array)

An instance A of the data type h_{-array} (hashing array) is an injective mapping from a data type I, called the index type of A, to a set of variables of a data type E, called the element type of A. I must be char, int, a pointer type, or an item type.

1. Declaration of an hashing array type

 $declare2(h_array, I, E)$

introduces a new data type with name $h_{array}(I, E)$ consisting of all h_arrays with index type I and element type E.

2. Creation of a h_array

$$h_{-}array(I, E) \quad A(x);$$

creates an injective function a from I to the set of unused variables of type E, assigns x to all variables in the range of a and initializes A with a.

3. Operations on a h-array A

E& $A \ [I \ x]$ returns the variable A(x)bool A.defined(I x) returns true if $x \in dom(A)$, false otherwise; here dom(A) is the set of all $x \in I$ for which A[x] has already been executed.

4. Iteration

for all_defined (x, A) { "the elements from dom(A) are successively assigned to x" }

5. Implementation

Hashing arrays are implemented by dynamic perfect hashing ([DKMMRT88]). Access operations A[x] take time O(1). Hashing arrays are more efficient than dictionary arrays.

4.6 Sorted Sequences (sortseq)

An instance S of the data type sortseq is a sequence of items (seq_item). Every item contains a key from a linearly ordered data type K, called the key type of S, and an information from a data type I, called the information type of S. The number of items in S is called the size of S. A sorted sequence of size zero is called empty. We use < k, i > to denote a seq_item with key k and information i (called the information associated with key k). For each $k \in K$ there is at most one item $< k, i > \in S$.

The linear order on K may be time-dependent, e.g., in an algorithm that sweeps an arrangement of lines by a vertical sweep line we may want to order the lines by the y-coordinates of their intersections with the sweep line. However, whenever an operation (except of reverse_items) is applied to a sorted sequence S, the keys of S must form an increasing sequence according to the currently valid linear order on K. For operation reverse_items this must hold after the execution of the operation.

1. Declaration of a sorted sequence type

declare2(sortseq, K, I)

introduces a new data type with name sortseq(K, I) consisting of all sorted sequences with key type K and information type I.

2. Creation of a sorted sequence

sortseq(K, I) S;

creates an instance S of type sortseq(K,I) and initializes it to the empty sorted sequence.

3. Operations on a sortseq S

K	S.key(seq_item it)	returns the key of item it Precondition: it is an item in S.
I	$S.inf(seq_item\ it)$	returns the information of item it Precondition: it is an item in S.
seq_item	S.lookup(K k)	returns the item with key k (nil if no such item exists in S)
seq_item	S.insert(K k, I i)	associates information i with key k : If there is an item $< k, j >$ in S then j is

replaced by i, else a new item < k, i > is added to S. In both cases the item is returned.

 $seg_item S.insert_at_item(seg_item it, K k, I i)$

Like insert(k,i), the item it gives the position of the item $\langle k, i \rangle$ in the sequence Precondition: it is an item in S with either key(it) is maximal with key(it) < k or

key(it) is minimal with key(it) > k

 $seq_item S.locate(K k)$ returns the item < k', i >in S such that

k' is minimal with k' >= k (nil if no

such item exists).

seq_item S.succ(seq_item it) returns the successor item of it, i.e., the

> item $\langle k, i \rangle$ in S such that k is minimal with k > key(it) (nil if no such item exists).

Precondition: it is an item in S.

seq_item S.pred(seq_item it) returns the predecessor item of it, i.e., the

> item $\langle k, i \rangle$ in S such that k is maximal with k < key(it) (nil if no such item exists).

Precondition: it is an item in S.

returns the item with maximal key $seq_item S.max()$

(nil if S is empty).

returns the item with minimal key $seq_item S.min()$

(nil if S is empty).

removes the item it from S. void $S. del_item(seg_item it)$

Precondition: it is an item in S.

S.del(K k)removes the item with key k from Svoid

(null operation if no such item exists).

voidS.change_inf($seq_item\ it,\ I\ i$) makes i the information of item it.

Precondition: it is an item in S.

void $S.reverse_items(seq_item a, seq_item b)$

> the subsequence of S from a to b is reversed. Precondition: Item a appears before item b

in S.

voidS.clear()makes S the empty sorted sequence.

S.size()returns the size of S. int

returns true if S is empty, false otherwise. S.empty()bool

4. Iteration

for all_seq_items(i, S) { "the items of S are successively assigned to i" }

5. Implementation

Sorted sequences are implemented by (2,4)-trees. Operations lookup, locate, insert, del take time $O(\log n)$, operations succ, pred, max, min, key, inf, insert_at_item and del_item take time O(1). Clear takes time O(n) and reverse_items $O(\ell)$, where ℓ is the length of the reversed subsequence. The space requirement is O(n). Here n is the current size of the sequence.

6. Example

Using a sorted sequence to list all elements in a sequence of strings lying lexicographically between two given search strings.

5. Graphs and Related Data Types

5.1 Directed graphs (graph)

An instance G of the data type graph consists of a set of nodes V and a set of edges E (node and edge are predefined data types). Every edge $e \in E$ is a pair of nodes $(v,w) \in V \times V$, v is called the source of e and w is called the target of e. With every node v the list of its adjacent edges $adj_list(v) = \{ e \in E \mid source(e) = v \}$, called the adjacency list of v, is associated.

1. Creation of a graph

graph G;

creates an instance G of type graph and initializes it to the empty graph.

2. Operations on a graph G

a) Access operations

int	$G.\mathrm{indeg}(node\ v)$	returns the indegree of node v
int	$G. { m outdeg}(node \ v)$	returns the outdegree of node v
node	G.source(edge e)	returns the source node of edge e
node	G.target(edge e)	returns the target node of edge e
int	$G.number_of_nodes()$	returns the number of nodes in G
int	$G.\mathtt{number_of_edges}()$	returns the number of edges in G
list(node)	$G.\mathtt{all_nodes}()$	returns the list of all nodes of G
list(edge)	$G.\mathtt{all_edges}()$	returns the list of all edges of G
list(edge)	$G.\mathtt{adj_edges}(node\ v)$	returns the list of all edges adjacent to v
list(node)	$G.\mathtt{adj_nodes}(node\ v)$	returns the list of all nodes adjacent to v
edge	$G. { m first_adj_edge}(node\ v)$	returns the first edge in the adjacency list of v
edge	$G.\mathtt{last_adj_edge}(node\ v)$	returns the last edge in the adjacency list of v
edge	$G.adj_succ(edge\ e)$	returns the successor of edge e in the
		adjacency list of source(e)
		(nil if it does not exist)
edge	$G.\mathtt{adj_pred}(edge\ e)$	returns the predecessor of edge e in the
		adjacency list of source(e)
		(nil if it does not exist)

edge	G.cyclic_adj_succ(edge e)	returns the cyclic successor of edge e in the adjacency list of $source(e)$
edge	$G. \operatorname{cyclic_adj_pred}(edge\ e)$	returns the cyclic predecessor of edge e in the adjacency list of $source(e)$
node	$G.{ m choose_node}()$	returns a node of G (nil if G is empty)
edge	$G.{ m choose_edge}()$	returns an edge of G (nil if G is empty)
b) Updat	te operations	
node	$G.\mathtt{new_node}()$	adds a new node to G and returns it
void	$G. ext{del_node}(node \ v)$	deletes v and all edges adjacent to v from G . Precondition: $indeg(v) = 0$.
edge	$G.\mathtt{new_edge}(node\ v,\ w)$	adds a new edge (v, w) to G by appending it to the adjacency list of v and returns it.
edge	G.new_edge(edge e, node w,	rel_pos $dir = after$) adds a new edge $e' = (source(e), w)$ to G by inserting it after $(dir = after)$ or before $(dir = before)$ edge e into the adjacency list of $source(e)$, returns e' .
void	$G. del_edge(edge e)$	deletes the edge e from G
void	$G. del_all_nodes()$	deletes all nodes from G
void	$G. \mathbf{del_all_edges}()$	deletes all edges from G
edge	$G.\mathtt{rev_edge}(edge\ e)$	reverses the edge $e = (v, w)$ by removing it from G and inserting the edge $e' = (w, v)$ into G by appending it to the adjacency list of w , returns e'
void	G.rev()	all edges in G are reversed
void	$G.sort_nodes(int(*cmp)(node))$	&, node&))
		the nodes of G are sorted according to the ordering defined by the comparing function cmp. Subsequent executions of forall_nodes step through the nodes in this order. (cf. TOPSORT1 in section 8.1)
void	$G. {f sort_edges}(int(*cmp)(edged))$	the edges of G are sorted according to the ordering defined by the comparing function cmp. Subsequent executions of forall_edges step through the edges in this order. (cf. TOPSORT1 in section 8.1)

list(edge) G.insert_reverse_edges()

for every edge (v, w) in G the reverse edge (w, v) is inserted into G. The list of all inserted edges is returned.

void G.clear()

makes G the empty graph

c) Iterators

With the adjacency list of every node v is associated a list iterator called the adjacency iterator of v (cf. list). There are operations to initialize the adjacency iterator, to move it to the successor or predecessor list item, to access its contents (an edge) and to test if it is defined, (\neq nil). Adjacency iterators are used to implement iteration statements (forall_adj_edges, forall_adj_nodes).

void G.init_adj_iterator($node\ v$) assigns nil to the adjacency iterator of node v

bool G.current_adj_edge(edge& e, node v)

if the adjacency iterator of v is defined (\neq nil) its contents is assigned to e and true is returned

else false is returned.

bool G.next_adj_edge(edge& e, node v)

moves the adjacency iterator of v forward (to the first item of $adj_list(v)$ if it is nil) and returns

 $G. \operatorname{current_adj_edge}(e, v)$

bool G.current_adj_node(node& w, node v)

if G.current_adj_edge(e, v) = true then assign target(e) to w and return true, else return

false

bool G.next_adj_node(node& w, node v)

if G.next_adj_edge(e, v) = true then assign target(e) to w and return true, else return

false

void G.reset() assign nil to all adjacency iterators in G

d) Miscellaneous operations

void G.write(string s) writes a compressed representation of G to

the file with name s

void G. read (string s) read a compressed representation of G from

the file with name s

void	$G.\mathtt{print_node}(node \ v)$	writes a readable representation of node v to the standard output
void	$G.\mathtt{print_edge}(edge\ e)$	writes a readable representation of edge e to the standard output
void	G.print()	writes a readable representation of G to the standard output

e) Operators

graph& G = G1

makes a copy of G1 and assigns it to G.

3. Iteration

```
forall_nodes(v,G) { "the nodes of G are successively assigned to v" }

forall_edges(e,G) { "the edges of G are successively assigned to e" }

forall_adj_edges(e,w)

{ "the edges adjacent to node w are successively assigned to e" }

forall_adj_nodes(v,w)

{ "the nodes adjacent to node w are successively assigned to v" }
```

4. Implementation

Graphs are implemented by adjacency lists. Most operations take constant time, except of all_nodes, all_edges, del_all_nodes, del_all_edges, clear, write, and read which take time O(n+m), where n is the current number of nodes and m is the current number of edges. The space requirement is O(n+m).

5. Examples

See section 8.1.

5.2 Undirected graphs (ugraph)

An instance G of the data type ugraph consists of a set of nodes V and a set of undirected edges E. Every edge $e \in E$ is a set of two nodes $\{v, w\}$, v and w are called the endpoints of e. With every node v is associated the list of its adjacent edges $adj_list(v) = \{e \in E \mid v \in e\}$.

1. Creation of an undirected graph

ugraph G;

creates an instance G of type ugraph and initializes it to the empty undirected graph.

2. Operations on a ugraph G

Most operations are the same as for directed graphs. The following operations are either additional or have different effects.

node	$G.opposite(node\ v,\ edge\ e)$	returns w if $e = \{v, w\}$, nil otherwise
int	$G. {\tt degree}(node \ v)$	returns the degree of node v .
edge	$G.\mathtt{new_edge}(node\ v,\ node\ w)$	inserts the undirected edge $\{v, w\}$ into G by appending it to the adjacency lists of both v and w and returns it
edge	$G.\mathtt{new_edge}(node\ v,\ node\ w,$	edge e1, edge e2, $dir1 = after$, $dir2 = after$) inserts the undirected edge $\{v, w\}$ after (if $dir1 = after$) or before (if $dir1 = before$) the edge e1 into the adjacency list of v and after (if $dir2 = after$) or before (if $dir2 = before$) the edge e2 into the adjacency list of w and returns it
edge	$G.adj_succ(edge\ e,\ node\ v)$	returns the successor of edge e in the adjacency list of v .
edge	$G.adj_pred(edge\ e,\ node\ v)$	returns the predecessor of edge e in the adjacency list of v .
edge	G.cyclic_adj_succ(edge e, node	e v)
		returns the cyclic successor of edge e in the adjacency list of v .
edge	G.cyclic_adj_pred(edge e, nod	e v)
		returns the cyclic predecessor of edge e in the

adjacency list of v.

3. Implementation

Undirected graphs are implemented like directed graphs by adjacency lists. The adjacency list of a node v contains all edges $\{v, w\}$ of the graph. Most operations take constant time, except of all_nodes, all_edges, del_all_nodes, del_all_edges, clear, write, and read which take time O(n+m), where n is the current number of nodes and m is the current number of edges. The space requirement is O(n+m).

5.3 Planar Maps (planar_map)

An instance M of the data type $planar_{-}map$ is the combinatorial embedding of a planar graph.

1. Creation of a planar_map

```
planar_map \ M(graph \ G);
```

creates an instance M of type $planar_map$ and initializes it to the planar map represented by the directed graph G. Precondition: G represents an undirected planar map, i.e. for every edge (v, w) in G the reverse edge (w, v) is also in G and there is a planar embedding of G such that for every node v the ordering of the edges in the adjacency list of v corresponds to the counter-clockwise ordering of these edges around v in the embedding.

2. Operations on a planar_map M

Most operations are the same as for directed graphs. The following operations are either additional or have different effects.

face	$M.{ t adj_face}(edge\ e)$	returns the face of M to the right of e .
list(face)	$M.\mathtt{all_faces}()$	returns the list of all faces of M .
list(face)	$M.{ m adj_faces}(node\ v)$	returns the list of all faces of M adjacent to node v in counter-clockwise order.
list(edge)	$M.\mathrm{adj_edges}(face\ f)$	returns the list of all edges of M bounding face f in clockwise order.
list(node)	$M.{ m adj_nodes}(face\ f)$	returns the list of all nodes of M adjacent to face f in clockwise order.
edge	M.reverse(edge e)	returns the reversal of edge e in M .

edge	$M.{ m first_face_edge}()$	returns the first edge of face f in M .
edge	$M.\mathtt{succ_face_edge}(edge\ e)$	returns the successor edge of e in face f i.e., the next edge in clockwise order.
edge	$M.\mathtt{pred_face_edge}(edge\ e)$	returns the predecessor edge of e in face f , i.e., the next edge in counter-clockwise order.
edge	M.new_edge(edge e1, edge e	(e_2)
		inserts the edge $e = (source(e_1), source(e_2))$ and its reversal edge into M . Precondition: e_1 and e_2 are bounding the same face F . The operation splits F into two new faces.
edge	$M.\mathrm{del_edge}(edge\ e)$	deletes the edge e from M . The two faces adjacent to e are united to one face.
list(edge)	$M. { m triangulate}()$	triangulates all faces of M by inserting new edges. The list of inserted edges is is returned.
void	$M. { m straight_line_embedding} (r$	$node_array(int)$ $xcoord$, $node_array(int)$ $ycoord$) computes a straight line embedding for M with integer coordinates $xcoord[v]$, $ycoord[v]$) in the range $02(n-1)$ for every node v of M .

3. Iteration

Additional iteration macros are

```
forall_faces(f, M) { "the faces of M are successively assigned to f" }

forall_adj_edges(e, f)

{ "the edges adjacent to face f are successively assigned to e" }
```

4. Implementation

Planar maps are implemented by parameterized directed graphs. All operations take constant time, except of, new_edge and del_edge which take time O(f) where f is the number of edges in the created faces, and triangulate and straight_line_embedding take time O(n) where n is the current size (number of edges) of the planar map.

5.4 Parameterized Graphs (GRAPH)

A parameterized graph G is a graph whose nodes and edges contain additional (user defined) informations. Every node contains an element of a data type vtype, called the node type of G and every edge contains an element of a data type etype called the edge type of G. We use $\langle v, w, y \rangle$ to denote an edge (v, w) with information y and $\langle x \rangle$ to denote a node with information x.

All operations defined on instances of the data type graph are also defined on instances of any parameterized graph type GRAPH(vtype, etype). For parameterized graphs there are additional operations to access or update the informations associated with its nodes and edges. Instances of a parameterized graph type can be used wherever an instance of the data type graph can be used, e.g., in assignments and as arguments to functions with formal parameters of type graph or graph. If a function f(graph G) is called with an argument G of type GRAPH(vtype, etype) then inside G only the basic graph structure of G (the adjacency lists) can be accessed. The node and edge informations are hidden. This allows the design of generic graph algorithms, i.e., algorithms accepting instances of any parametrized graph type as argument.

1. Declaration of a parameterized graph type

declare2(GRAPH, vtype, etype)

introduces a new data type with name GRAPH(vtype, etype) consisting of all parameterized graphs with node type vtype and edge type etype.

2. Creation of a parameterized graph

GRAPH(vtype, etype) G;

creates an instance G of type GRAPH(vtype, etype) and initializes it to the empty graph.

3. Operations on a GRAPH G

In addition to the operations of the data type graph (see section 2):

vtype G.inf(node v) returns the information of node v

etype G.inf(edge e) returns the information of edge e

void G.assign(node v, vtype x) makes x the information of node v

void G.assign(edge e, etype y) makes y the information of edge e

node G.new_node(vtype x) adds a new node < x > to G and returns it

edge G.new_edge(node v, w, etype x)

adds a new edge $e = \langle v, w, x \rangle$ to G by appending it to the adjacency list of v and returns e.

edge G.new_edge($edge\ e,\ node\ w,\ etype\ x,\ dir = after$)

adds a new edge $e' = \langle source(e), w, x \rangle$ to G by inserting it after (dir = after) or before (dir = before) edge e into the adjacency list of source(e) and returns e'.

4. Operators

vtype& G [node v] returns G.inf(v).

etype& G [edge e] returns G.inf(e).

5. Implementation

Parameterized graphs are derived from directed graphs. All additional operations for manipulating the node and edge informations take constant time.

5.5 Parameterized undirected graphs (UGRAPH)

A parameterized undirected graph G is an undirected graph whose nodes and edges contain additional (user defined) informations. Every node contains an element of a data type vtype, called the node type of G and every edge contains an element of a data type etype called the edge type of G. We use $\langle \{v,w\},y\rangle$ to denote the undirected edge $\{v,w\}$ with information y and $\langle x\rangle$ to denote a node with information x.

1. Declaration of a parameterized undirected graph type

declare2(UGRAPH, vtype, etype)

introduces a new data type with name UGRAPH(vtype, etype) consisting of all undirected parameterized graphs with node type vtype and edge type etype.

2. Creation of a parameterized undirected graph

UGRAPH(vtype, etype) G;

creates an instance G of type UGRAPH(vtype, etype) and initializes it to the empty graph.

3. Operations on a UGRAPH G

In addition to the operations of the data type ugraph (see section 5.3):

 $vtype/etypG.inf(node/edge\ a)$ returns the information of node/edge a

void G.assign($node/edge \ a, \ vtype/etype \ x$)

makes x the information of node/edge a

node G.new_node(vtype x) adds a new node < x > to G and returns it

edge G.new_edge(node v, node w, etype x)

inserts the undirected edge $< \{v, w\}, x >$ into G by appending it to the adjacency lists of both v and w and returns it

edge G.new_edge(node v, node w, reflected, etc. v) rel_posdir1 =) inserts the undirected edge v, v, v after (if dir1 = after) or before (if dir1 = before) the edge v into the adjacency list of v and after (if dir2 = after) or before (if dir2 = before) the edge v into the adjacency list of v and v and v returns it.

4. Implementation

Parameterized undirected graphs are derived from undirected graphs. All additional operations for manipulating the node and edge informations take constant time.

5.6 Parameterized planar maps (PLANAR_MAP)

A parameterized planar map M is a planar map whose nodes and faces contain additional (user defined) informations. Every node contains an element of a data type vtype, called the node type of M and every face contains an element of a data type ftype called the face type of M. All operations of the data type $planar_map$ are also defined for instances of any parameterized planar_map type. For parameterized planar maps there are additional informations to access or update the node and face informations.

1. Declaration of a parameterized planar_map type

 $declare2(PLANAR_MAP, vtype, ftype)$

introduces a new data type with name $PLANAR_MAP(vtype, ftype)$ consisting of all parameterized planar maps with node type vtype and face type ftype. Precondition: The data type GRAPH(vtype, ftype), i.e., the parameterized directed graph type with node entries of type vtype and edge entries of type ftype, has been declared before.

2. Creation of a parameterized planar map

 $PLANAR_MAP(vtype, ftype) M(GRAPH(vtype, ftype) G);$

creates an instance M of type $PLANAR_MAP(vtype, ftype)$ and initializes it to the planar map represented by the parameterized directed graph G. The node entries of G are copied into the corresponding nodes of M and every face f of M is assigned the information of one of its bounding edges in G. Precondition: G represents a planar map.

3. Operations on a PLANAR_MAP M

In addition to the operations of the data type planar_map:

vtype $M.inf(node\ v)$ returns the information of node vftype $M.inf(face\ f)$ returns the information of face f

```
woid M.assign(node\ v,\ vtype\ x) makes x the information of node v woid M.assign(face\ f,\ ftype\ y) makes y the information of face f M.new\_edge(edge\ e_1,\ edge\ e_2,\ ftype\ y) inserts the edge e = (source(e_1), source(e_2)) and its reversal edge e' into M. Precondition: e_1 and e_2 are bounding the same face F. The operation splits F into two new faces f, adjacent to edge e and f', adjacent to edge e' with \inf(f) = \inf(F) and \inf(f') = y.
```

4. Implementation

Parameterized planar maps are derived from planar maps. All additional operations for manipulating the node and edge informations take constant time.

5.7 Node and edge arrays (node_array, edge_array)

An instance A of the data type $node_array$ ($edge_array$) is a partial mapping from the node set V (edge set E) of a (u)graph G to a set of variables of a data type E, called the element type of the array. The domain I of A is called the index set of A and A(x) is called the element at position x. A is said to be valid for all nodes (edges) in I.

1. Declaration of node and edge array types

 $declare(node/edge_array, E)$

introduces a new data type with name $node_array(E)$ ($edge_array(E)$) consisting of all node (edge) arrays with element type E.

2. Creation of a node array (edge array)

- a) node/edge_array A;
- b) $node/edge_array \ A(graph \ G);$
- c) node/edge_array A(graph G, E x);

creates an instance A of type $node_array(E)$ or $edge_array(E)$. Variant a) initializes the index set of A to the empty set, Variants b) and c) initialize the index set of A

to be the entire node (edge) set of graph G, i.e., A is made valid for all nodes (edges) currently contained in G. Variant c) in addition initializes A(i) with x for all nodes (edges) i of G.

3. Operations

void	$A.\mathrm{init}(graph\ G)$	sets the index set I of A to the node (edge) set of G , i.e., makes A valid for all nodes (edges) of G .
void	$A.init(graph \ G, \ E \ x)$	makes A valid for all nodes (edges) of G and sets $A(i) = x$ for all nodes (edges) of G
E&	$A \ [node/edge \ i]$	access the variable $A(i)$. Precondition: A must be valid for node (edge)

4. Implementation

Node (edge) arrays for a graph G are implemented by C++ vectors and an internal numbering of the nodes and edges of G. The access operation takes constant time, init takes time O(n), where n is the number of nodes (edges) currently in G. The space requirement is O(n).

Important: A node (edge) array is only valid for the nodes (edges) contained in G at the moment of the array declaration or initialization (init). Access operations for later added nodes (edges) are not allowed. Node and edge arrays for dynamic graphs can be realized using hashing arrays (cf. section 4.5), e.g. by declare2($h_array, edge, int$).

5.8 Two dimensional node arrays (node_matrix)

An instance M of the data type $node_matrix$ is a partial mapping from the set of node pairs $V \times V$ of a graph to a set of variables of a data type E, called the element type of M. The domain I of M is called the index set of M. M is said to be valid for all node pairs in I. A node matrix can also be viewed as a node array with element type $node_array(E)$ ($node_array(node_array(E))$).

1. Declaration of a node matrix type

 $declare(node_matrix, E)$

introduces a new data type with name $node_matrix(E)$ consisting of all node matrices with element type E.

2. Creation of a node_matrix

- a) $node_matrix(E)$ M;
- b) $node_matrix(E)$ M(G);
- c) $node_matrix(E)$ M(G, x);

creates an instance M of type $node_matrix(E)$. Variant a) initializes the index set of M to the empty set, Variants b) and c) initialize the index set of A to be the set of all node pairs of graph G, i.e., M is made valid for all pairs in $V \times V$ where V is the set of nodes currently contained in G. Variant c) in addition initializes M(v, w) with x for all nodes $v, w \in V$.

3. Operations on a node_matrix M

void	$M.\mathrm{init}(graph\ G)$	sets the index set of M to $V \times V$, where V is the set of all nodes of G
void	$M.init(graph \ G, \ E \ x)$	sets the index set of M to $V \times V$, where V is the set of all nodes of G and initializes $M(v,w)$ to x for all $v,w \in V$.
E&	$M \ (node \ v, \ node \ w)$	returns the variable $M(v, w)$. Precondition: M must be valid for v and w .
$node_array(E)$ & $M[v]$		returns the node_array $M(v)$.

4. Implementation

Node matrices for a graph G are implemented by vectors of node arrays and an internal numbering of the nodes of G. The access operation takes constant time, the init operation takes time $O(n^2)$, where n is the number of nodes currently contained in G. The space requirement is $O(n^2)$. Note that a node matrix is only valid for the nodes contained in G at the moment of the matrix declaration or initialization (init). Access operations for later added nodes are not allowed.

5.9 Sets of nodes and edges (node_set, edge_set)

An instance S of the data type $node_set$ ($edge_set$) is a subset of the nodes (edges) of a graph G. S is said to be valid for the nodes (edges) of G.

1. Creation of a node or edge set

```
node\_set S(G);
edge\_set S(G);
```

creates an instance S of type $node_set$ ($edge_set$) valid for all nodes (edges) currently contained in graph G and initializes it to the empty set.

2. Operations on a node/edge set S

void	$S.\mathrm{insert}(x)$	adds node (edge) x to S
void	$S.\operatorname{del}(x)$	removes node (edge) x from S
bool	$S.\mathrm{member}(x)$	returns true if x in S , false otherwise
$node/edge\ S. { m choose}()$		return a node (edge) of S
int	$S.{ m size}()$	returns the size of S
bool	$S.\mathtt{empty}()$	returns true iff S is the empty set
void	$S.\mathtt{clear}()$	makes S the empty set

3. Implementation

A node (edge) set S for a graph G is implemented by a combination of a list L of nodes (edges) and a node (edge) array of list_items associating with each node (edge) its position in L. All operations take constant time, except of clear which takes time O(|S|). The space requirement is O(n), where n is the number of nodes (edges) of G.

5.10 Node partitions (node_partition)

An instance of the data type $node_partition$ is a partition of the nodes of some graph G.

1. Creation of a node partition

 $node_partition P(G);$

creates an instance P of type $node_partition$ containing for every node v in G a block $\{v\}$.

2. Operations on a node_partition P

bool	$P.\mathtt{same_block}(node\ v,\ node\ w)$	returns true if v and w belong to the same block of P .
void	P.union_blocks(node v, node w)	unites the blocks of P containing nodes v and w .
node	$P.\operatorname{find}(node\ v)$	returns a canonical node of the block that contains node v .

3. Implementation

A node partition for a graph G is implemented by a combination of a partition P and a node array of $partition_item$ associating with each node in G a partition item in P. Initialization takes linear time, union_blocks takes time O(1) (worst-case), and same_block and find take time $O(\alpha(n))$ (amortized). The space requirement is O(n), where n is the number of nodes of G.

5.11 Node priority queues (node_pq)

An instance Q of the data type $node_{-}pq$ is a partial function from the nodes of a graph G to some linearly ordered type I.

1. Declaration of a node priority queue type

$declare(node_pq, I)$

introduces a new data type with name $node_{-}pq(I)$ consisting of all node priority queues with information type I.

2. Creating a node priority queue

$$node_pq(I)$$
 $Q(G)$;

creates an instance Q ot type $node_{-}pq(I)$ for the nodes of graph G with $dom(Q) = \emptyset$.

3. Operations on a node_pq Q

void	$Q.insert(node \ v, \ I \ i)$	adds the node v with information i to Q . Precondition: $v \notin dom(Q)$.
void	$Q. ext{decrease_inf}(node\ v,\ I\ i)$	makes i the new information of node v (precondition: $i \leq Q(v)$)
node	$Q.\mathrm{find_min}()$	returns a node with the minimal information (nil if Q is empty)
void	$Q.\mathrm{del}(node\ v)$	removes the node v from Q
node	$Q.\mathrm{del_min}()$	removes a node with the minimal information from Q and returns it (nil if Q is empty)
void	$Q.\mathtt{clear}()$	makes Q the empty node priority queue.
bool	$Q.\mathrm{empty}()$	returns true if Q is the empty node priority queue, false otherwise.

4. Implementation

Node priority queues are implemented by fibonacci heaps and node arrays. Operations insert, del_node, del_min take time $O(\log n)$, find_min, decrease_inf, empty take time O(1) and clear takes time O(m), where m is the size of NQ. The space requirement is O(n), where n is the number of nodes of G.

5.12 Graph Algorithms

This sections gives a summary of the graph algorithms contained in LEDA. All algorithms are generic, i.e., they accept instances of any user defined parameterized graph type GRAPH(vtype, etype) as arguments.

5.12.1 Basic Algorithms

• Topological Sorting

bool TOPSORT(graph& G, node_array(int)& ord)

TOPSORT takes as argument a directed graph G(V, E). It sorts G topologically (if G is acyclic) by computing for every node $v \in V$ an integer ord[v] such that $1 \leq ord[v] \leq |V|$ and ord[v] < ord[w] for all edges $(v, w) \in E$. TOPSORT returns true if G is acyclic and false otherwise.

Running Time: O(|V| + |E|)

• Depth First Search

 $list(node) DFS(graph\& G, node s, node_array(bool)\& reached)$

DFS takes as argument a directed graph G(V, E), a node s of G and a node_array reached of boolean values. It performs a depth first search starting at s visiting all reachable nodes v with reached[v] = false. For every visited node v reached[v] is changed to true. DFS returns the list of all reached nodes.

Running Time: O(|V| + |E|)

list(edge) DFS_NUM(graph& G, node_array(int)& df snum, node_array(int)& compnum)

DFS_NUM takes as argument a directed graph G(V, E). It performs a depth first search of G numbering the nodes of G in two different ways. dfsnum is a numbering with respect to the calling time and compnum a numbering with respect to the completion time of the recursive calls. DFS_NUM returns a depth first search forest of G (list of tree edges).

Running Time: O(|V| + |E|)

• Breadth First Search

list(node) BFS(graph& G, node s, node_array(int)& dist)

BFS takes as argument a directed graph G(V, E) and a node s of G. It performs a breadth first search starting at s computing for every visited node v the distance (length of a shortest path) dist[v] from s to v. BFS returns the list of all reached nodes.

Running Time: O(|V| + |E|)

• Connected Components

int COMPONENTS(ugraph& G, node_array(int)& compnum)

COMPONENTS takes an undirected graph G(V, E) as argument and computes for every node $v \in V$ an integer compnum[v] from [0...c-1] where c is the number of connected components of G and v belongs to the i-th connected component iff compnum[v] = i. COMPONENTS returns c.

Running Time: O(|V| + |E|)

• Strong Connected Components

int STRONG_COMPONENTS(graph& G, node_array(int)& compnum)

STRONG_COMPONENTS takes a directed graph G(V, E) as argument and computes for every node $v \in V$ an integer compnum[v] from [0...c-1] where c is the number of strongly connected components of G and v belongs to the i-th strongly connected component iff compnum[v] = i. STRONG_COMPONENTS returns c.

Running Time: O(|V| + |E|)

• Transitive Closure

 $graph TRANSITIVE_CLOSURE(graph\& G)$

TRANSITIVE_CLOSURE takes a directed graph G(V, E) as argument and computes the transitive closure of G(V, E). It returns a directed graph G'(V', E') with V' = V and $(v, w) \in E' \Leftrightarrow$ there is a path form v to w in G.

Running Time: $O(|V| \cdot |E|)$

5.12.2 Network Algorithms

Most of the following network algorithms are overloaded. They work for both integer and real valued edge costs.

• Single Source Shortest Paths

DIJKSTRA takes as arguments a directed graph G(V,E), a source node s and an edge_array cost giving for each edge in G a non-negative cost. It computes for each node v in G the distance dist[v] from s (cost of the least cost path from s to v) and the predecessor edge pred[v] in the shortest path tree.

```
Running Time: O(|E| + |V| \log |V|)
```

BELLMAN_FORD takes as arguments a graph G(V,E), a source node s and an edge_array cost giving for each edge in G a real (integer) cost. It computes for each node v in G the distance dist[v] from s (cost of the least cost path from s to v) and the predecessor edge pred[v] in the shortest path tree. BELLMAN_FORD returns false if there is a negative cycle in G and true otherwise

Running Time: $O(|V| \cdot |E|)$

• All Pairs Shortest Paths

```
void \ ALL\_PAIRS\_SHORTEST\_PATHS(graph\&\ G,\ edge\_array(int)\&\ cost, \\ node\_matrix(int)\&\ dist) void \ ALL\_PAIRS\_SHORTEST\_PATHS(graph\&\ G,\ edge\_array(real)\&\ cost)
```

 $node_matrix(real) \& dist)$

ALL_PAIRS_SHORTES_PATHS takes as arguments a graph G(V, E) and an edge_array cost giving for each edge in G a real (integer) valued cost. It computes for each node pair (v, w) of G the distance dist(v, w) from v to w (cost of the least cost path from v to w).

Running Time: $O(|V| \cdot |E| + |V|^2 \log |V|)$

• Maximum Flow

int MAX_FLOW(graph& G, node s, node t, edge_array(int)& cap, edge_array(int)& flow)

int MAX_FLOW(graph& G, node s, node t, edge_array(real)& cap, edge_array(real)& flow)

MAX_FLOW takes as arguments a directed graph G(V, E), a source node s, a sink node t and an edge_array cap giving for each edge in G a capacity. It computes for every edge e in G a flow flow[e] such that the total flow from s to t is maximal and $flow[e] \leq cap[e]$ for all edges e. MAXFLOW returns the total flow from s to t.

Running Time: $O(|V|^3)$

• Maximum Cardinality Bipartite Matching

list(edge) MAX_CARD_BIPARTITE_MATCHING(graph& G, list(node)& A, list(node)& B)

MAX_CARD_BIPARTITE_MATCHING takes as arguments a directed graph G(V, E) and two lists A and B of nodes. All edges in G must be directed from nodes in A to nodes in B. It computes a maximum cardinality bipartite matching of G, i.e., a maximal set of edges M such that no two edges in M share an end point (target or source). MAX_CARD_BIPARTITE_MATCHING returns M as a list of edges.

Running Time: $O(|E|\sqrt{|V|})$

• Maximum Weight Bipartite Matching

list(edge) MAX_WEIGHT_BIPARTITE_MATCHING(graph& G,

list(node)& A, list(node)& B, edge_array(int)& weight) list(edge) MAX_WEIGHT_BIPARTITE_MATCHING(graph&~G, list(node)&~A, list(node)&~B, $edge_array(real)\&~weight)$

MAX_WEIGHT_BIPARTITE_MATCHING takes as arguments a directed graph G, two lists A and B of nodes and an edge_array giving for each edge an integer (real) weight. All edges in G must be directed from nodes in A to nodes in B. It computes a maximum weight bipartite matching of G, i.e., a set of edges M such that the sum of weights of all edges in M is maximal and no two edges in M share an end point. MAX_WEIGHT_BIPARTITE_MATCHING returns M as a list of edges.

Running Time: $O(|V| \cdot |E|)$

Spanning Tree

list(edge) SPANNING_TREE(ugraph& G)

SPANNING_TREE takes as argument an undirected graph G(V, E). It computes a spanning tree T of G, SPANNING_TREE returns the list of edges of T.

Running Time: O(|V| + |E|)

Minimum Spanning Tree

 $list(edge) \ \text{MIN_SPANNING_TREE}(ugraph\&G, \ edge_array(int)\& \ cost)$

list(edge) MIN_SPANNING_TREE(ugraph&G, edge_array(real)& cost)

MIN_SPANNING_TREE takes as argument an undirected graph G(V, E) and an edge_array cost giving for each edge an integer cost. It computes a minimum spanning tree T of G, i.e., a spanning tree such that the sum of all edge costs is minimal. MIN_SPANNING_TREE returns the list of edges of T.

Running Time: $O(|E| \log |V|)$

5.12.3 Algorithms for Planar Graphs

• Planarity Test

bool PLANAR(graph&G)

PLANAR takes as input a directed graph G(V, E) and performs a planarity test for G. If G is a planar graph it is transformed into a planar map (a combinatorial embedding such that the edges in all adjacency lists are in clockwise ordering). PLANAR returns true if G is planar and false otherwise.

Running Time: O(|V| + |E|)

• Triangulation

list(edge) TRIANGULATE_PLANAR_MAP(graph& G)

TRIANGULATE_PLANAR_MAP takes a directed graph G representing a planar map. It triangulates the faces of G by inserting additional edges. The list of inserted edges is returned.

Running Time: O(|V| + |E|)

• Straight Line Embedding

int STRAIGHT_LINE_EMBEDDING(graph& G, node_array(int)& xcoord, node_array(int)& ycoord)

STRAIGHT_LINE_EMBEDDING takes as argument a directed graph G representing a planar map. It computes a straight line embedding of G by assigning nonnegative integer coordinates (xcoord and ycoord) in the range 0..2(n-1) to the nodes. STRAIGHT_LINE_EMBEDDING returns the maximal coordinate.

Running Time: $O(|V|^2)$

5.13 Miscellaneous

5.13.1 Some useful functions

void complete_graph(graph& G, int n)

creates a complete graph G with n nodes.

void random_graph(graph& G, int n, int m)

creates a random graph G with n nodes and m edges.

void test_graph(graph& G)

creates interactively a user defined graph G.

void test_bigraph(graph& G, nodelist& A, nodelist& B)

creates interactively a user defined bipartite graph G with sides A and B. All edges are directed from A to B.

bool compute_correspondence(graph& G, edge_array(edge)& reversal)

computes for every edge e = (v, w) in G its reversal reversal[e] = (w, v) in G (if present). Returns true if every edge has a reversal and false otherwise.

void eliminate_parallel_edges(graph& G)

removes all parallel edges from G.

5.13.2 Predefined parameterized types

list(node)	list(edge)
$node_array(int) \ node_array(bool) \ node_array(real) \ node_array(node)$	$edge_array(int) \ edge_array(bool) \ edge_array(real) \ edge_array(node)$
$node_array(edge)$	$edge_array(edge)$
$node_matrix(int)$	
$node_matrix(bool)$	
$node_matrix(real)$	

6. Data Types For Two-Dimensional Geometry

6.1 Basic two-dimensional objects

LEDA provides a collection of simple data types for two-dimensional geometry, such as points, segments, lines, circles, and polygons. All these types can be used as type parameters in parameterized data types. Their declarations are contained in the header file < LEDA/plane.h>. The corresponding list types list(point), list(segment), list(line), list(circle), and list(polygon) are also declared in this file. Furthermore, some basic algorithms (section 6.1.6) are included.

Note: This section is preliminary and will probably change in future versions of the library. Above all, there is missing a hierarchy of the data types and a general concept for infinite objects.

6.1.1 Points (point)

An instance of the data type point is a point in the two-dimensional plane \mathbb{R}^2 . We use (a, b) to denote a point with first (or x-) coordinate a and second (or y-) coordinate b.

1. Creation of a point

- a) point p(real x, real y);
- b) point p;

introduces a variable p of type point initialized to the point (x, y). Variant b) initializes p to the point (0, 0).

2. Operations on a point p

real	p.xcoord()	returns the first coordinate of point p
real	p.ycoord()	returns the second coordinate of point p
real	$p.{ m distance}(point \ q)$	returns the euclidean distance between p and q .
real	$p.{ m distance}()$	returns the euclidean distance between p and $(0,0)$.
point	p.translate(vector v)	returns $p + v$, i.e., p translated by vector v . Precondition: $v.\dim() = 2$.

point $p.translate(real \ \alpha, real \ d)$

returns the point created by translating p in direction α by distance d. The direction is given by its angle with a right principled having the principled by

right oriented horizontal ray.

point p.rotate(point q, real α) returns the point created by a rotation of p

about point q by angle α .

3. Operators

point & point = point assignment

bool point == point test for equality

bool point != point test for inequality

point point + vector translation by vector

Input and output operators:

ostream & ostream << point writes a point to an output stream

istream & istream >> point reads the coordinates of a point (two reals)

from an input stream

6.1.2 Segments (segment)

An instance s of the data type segment is a directed straight line segment in the two-dimensional plane, i.e., a straight line segment [p,q] connecting two points $p,q \in \mathbb{R}^2$. p is called the start point and q is called the end point of s. The length of s is the euclidean distance between p and q. The angle between a right oriented horizontal ray and s is called the direction of s. The segment [(0,0),(0,0)] is said to be empty.

1. Creation of a segment

- a) $segment \ s(point \ p, \ point \ q);$
- b) segment $s(point p, real \alpha, real d)$;
- c) segment s;

introduces a variable s of type segment. s is initialized to the segment from p to q (variant a), to the segment with start point p, direction α , and length d (variant b) or to the empty segment (variant c).

2. Operations on a segment s

point	s.start()	returns the start point of segment s.
point	$s.\mathrm{end}()$	returns the end point of segment s.
real	s.xcoord1()	returns the x-coordinate of s.start().
real	s.ycoord1()	returns the y-coordinate of s.start().
real	s.xcoord2()	returns the x-coordinate of s.end().
real	s.ycoord2()	returns the y-coordinate of s.end().
real	$s.\mathrm{length}()$	returns the length of s.
real	s.direction()	returns the direction of s as an angle in the intervall $(-\pi, \pi]$.
real	s.angle(segment t)	returns the angle between s and t, i.e., t.direction() - s.direction().
real	s.angle()	returns s.direction().
bool	s.horizontal()	returns true iff s is horizontal.
bool	s.vertical()	returns true iff s is vertical.
real	s.slope()	returns the slope of s .
		Precondition: s is not vertical.
bool	s.intersection(segment t,	point& p)
		if s and t are not collinear and intersect the intersection point is assigned to p and true is
		returned, otherwise false is returned.
segment	s.rotate(point q, real α)	returns the segment created by a rotation of s
		about point q by angle α .
segment	$s.rotate(real \ lpha)$	returns $s.rotate(s.start(),\alpha)$.
segment	s. translate(vector v)	
		returns $s + v$, i.e., the segment created by translating s by vector v. <i>Precondition</i> : v has dimension 2.
segment	s.translate(real alpha, re	$al \ d)$
		returns the segment created by a translation of s in direction α by distance d .

3. Operators

segment & segment = segment assignment bool segment == segment test for equality

bool	segment != segment	test for inequality
segment	segment + vector	translation by vector

Input and output operators:

```
ostream&ostream << segment</th>writes a segment to an output stream.istream&istream >> segmentreads the coordinates of a segment (four reals)from an input stream.
```

6.1.3 Straight Lines (line)

An instance l of the data type line is a directed straight line in the two-dimensional plane. The angle between a right oriented horizontal line and l is called the direction of l.

1. Creation of a line

```
a) line l(point p, point q);
```

- b) line l(segment s);
- c) line $l(point p, real \alpha);$
- d) line l;

introduces a variable l of type line. l is initialized to the line passing through points p and q directed form p to q (variant a), to the line supporting segment s (variant b), to the line passing through point p with direction α (variant c), or a line through (0,0) with direction 0 (variant d).

2. Operations on a line l

real	$l. { m direction}()$	returns the direction of l .
real	$l.{ m angle}(lineg)$	returns the angle between l and g , i.e., g .direction() - l .direction().
real	$l.{ m angle}()$	returns l .direction().
bool	$l. {\tt horizontal}()$	returns true iff l is horizontal.
bool	$l. { m vertical}()$	returns true iff l is vertical.
real	$l.{ m slope}()$	returns the slope of l .

Precondition: l is not vertical.

real l.y.proj(real x) returns p.ycoord(), where $p \in l$ with p.xcoord()

= x. Precondition: l is not vertical.

real $l.x_proj(real y)$ returns p.xcoord(), where $p \in l$ with p.ycoord()

= y. Precondition: l is not horizontal.

real $l.y_abs()$ returns the y-abscissa of l $(l.y_proj(0))$.

Precondition: l is not vertical.

bool l.intersection(line g, point & p)

if l and g are not collinear and intersect the intersection point is assigned to p and true is returned, otherwise false is returned.

bool l.intersection(segment s, point& p)

if l and s are not collinear and intersect the intersection point is assigned to p and true is returned atherwise false is returned

returned, otherwise false is returned.

l.translate(vector v)

returns l + v, i.e., the line created by translating l by vector v. Precondition: v has dimension 2.

line l.translate(real alpha, real d)

returns the line created by a translation of

l in direction α by distance d.

line l.rotate(point q, real α) returns the line created by a rotation of l

about point q by angle α .

segment l.perpendicular(point p) returns the nromal of p with respect to l.

3. Operators

line& line = line assignment

bool line == line test for equality

bool line!= line test for inequality

6.1.4 Polygons (polygon)

An instance P of the data type polygon is a simple polygon in the two-dimensional plane defined by the sequence of its vertices in clockwise order. The number of vertices is called the size of P. A polygon with empty vertex sequence is called empty.

1. Creation of a polygon

- a) polygon P(list(point) pl);
- b) polygon P;

introduces a variable P of type polygon. P is initialized to the polygon with vertex sequence pl. Precondition: The vertices in pl are given in clockwise order and define a simple polygon. Variant b) creates the empty polygon and assings it to P.

2. Operations on a polygon P

list(point)	P.vertices()	returns the vertex sequence of P .
list (segment	$)P.{ m segments}()$	returns the sequence of bounding segments of Pin clockwise order.
list(point)	P.intersection(line l)	returns $P \cap l$ as a list of points.
list(point)	P.intersection(segment s)	returns $P \cap s$ as a list of points.
list(polygon)	P.intersection(polygon Q)	returns $P \cap Q$ as a list of points.
bool	P.inside(point p)	returns true if p lies inside of P , false otherwise.
bool	P.outside(point p)	returns $!P.inside(p)$.
polygon	$P.$ translate $(vector \ v)$	
		returns $P + v$, i.e., the polygon created by translating P by vector v . Precondition: v has dimension 2.
polygon	P.translate(real \alpha, real d	<i>(</i>)
		returns the polygon created by a translation of of P in direction α by distance d
polygon	$P.rotate(point q, real \alpha)$	returns the polygon created by a rotation of P about point q by angle α .
real	P.size()	returns the size of P.
real	P.empty()	returns true if P is empty, false otherwise.

3. Operators

polygon &	polygon = polygon	assignment
bool	polygon == polygon	test for equality
bool	polygon != polygon	test for inequality

6.1.5 Circles (circle)

An instance C of the data type circle is a circle in the two-dimensional plane, i.e., the set of points having a certain distance r from a given point p. r is called the radius and p is called the center of C. The circle with center (0,0) and radius 0 is called the empty circle.

1. Creation of a circle

- a) circle C(point p, real r);
- b) circle C;

introduces a variable C of type *circle*. C is initialized to the circle with center p and radius r. Variant b) creates the empty circle and assigns it to C.

2. Operations on a circle C

real	$C.\mathtt{radius}()$	returns the radius of C .
point	$C.\mathtt{center()}$	returns the center of C .
list(point)	C.intersection(line l)	returns $C \cap l$ as a list of points.
list(point)	C.intersection(segment s)	returns $C \cap s$ as a list of points.
list(point)	C.intersection(circle D)	returns $C \cap D$ as a list of points.
segment	$C.\operatorname{left_tangent}(point\ p)$	returns the line segment starting in p tangent to C and left of segment $[p, C.center()]$.
segment	C.right_tangent(point p)	returns the line segment starting in p tangent to C and right of segment $[p, C.center()]$.
real	$C. ext{distance}(point p)$	returns the distance between C and p (negative if p inside C).
real	$C. { m distance}(line \ l)$	returns the distance between C and l (negative if l intersects C).
real	$C. ext{distance}(circle \ D)$	returns the distance between C and D (negative if D intersects C).

bool C.inside(point p) returns true if P lies inside of C,

false otherwise.

bool C.outside(point p) returns !C.inside(p).

circle C.translate(vector v) returns C + v, i.e., the circle created by

translating C by vector v. Precondition:

 $v.\dim = 2.$

circle C.translate(real α , real d)

returns the circle created by a translation of C

in direction α by distance d.

circle C.rotate(point q, real α) returns the circle created by a rotation of C

about point q by angle α .

3. Operators

circle & circle = circle assignment

bool circle == circle test for equality

bool circle != circle test for inequality

6.1.6 Algorithms

Line segment intersection

void SEGMENT_INTERSECTION(list(segment) & L, list(point) & P);

SEGMENT_INTERSECTION takes a list of segments L as input and computes the list of intersection points between all segments in L.

Running Time: $O((n+k)\log n)$, where n is the number of segments, and k is the number of intersections.

Convex hull of point set

polygon CONVEX_HULL(list(point) L);

CONVEX_HULL takes as argument a list of points and returns the polygon representing the convex hull of L. It is based on a randomized incremental algorithm.

Running Time: $O(n \log n)$ (with high probability), where n is the number of segments.

• Voronoi Diagrams

VORONOI takes as input a list of points sites and a real number R. It computes a directed graph G representing the planar subdivision defined by the Voronoi-diagram of sites where all "infinite" edges have length R. Node_array P stores for each node of G the corresponding Voronoi vertex (point) and edge_array C gives for each edge e of G the site (point) whose Voronoi region is bound by e.

6.1.7 Predefined parameterized data types

```
list(point), list(segment), list(line), list(polygon), list(circle)
GRAPH(point, int), node_array(point), edge_array(point)
```

6.2 Two-dimensional dictionaries (d2_dictionary)

An instance D of the data type $d2_dictionary$ is a collection of items $(dic2_item)$. Every item in D contains a key from a linearly ordered data type K1, a key from a linearly ordered data type K2, and an information from a data type I. K1 and K2 are called the key types of D, and I is called the information type of D. The number of items in D is called the size of D. A two-dimensional dictionary of size zero is said to be empty. We use $k_1, k_2, i > 0$ to denote the item with first key $k_1, k_2, i > 0$ and information $k_1, k_2, k_3 > 0$. Additionally to the normal dictionary operations, the data type $k_1, k_2, k_3 > 0$. Additionally to the normal dictionary operations, the data type $k_1, k_2, k_3 > 0$. Additionally range queries on $k_1 \times k_2$.

1. Declaration of a two-dimensional dictionary type

 $declare2(d2_dictionary, K1, K2, I)$

introduces a new data type with name $d2_dictionary(K1, K2, I)$ consisting of all two-dimensional dictionaries with key types K1 and K2 and information type I. Precondition: K1 and K2 are linearly ordered.

2. Creation of a two-dimensional dictionary

 $d2_dictionary(K1, K2, I)$ D;

creates an instance D of type d2_dictionary(K1, K2, I) and initializes D to the empty dictionary.

3. Operations on a d2-dictionary D

K1	$D.\mathtt{key1}(dic2_item\ it)$	returns the first key of item it . Precondition: it is an item in D .
K2	$D.\mathtt{key2}(dic2_item\ it)$	returns the second key of item it . Precondition: it is an item in D .
I	$D.inf(dic2_item\ it)$	returns the information of item it . Precondition: it is an item in D .
$dic2_item$	$D.\mathtt{max_key1}()$	returns the item with maximal first key.
$dic2_item$	$D.\mathtt{max_key2()}$	returns the item with maximal second key.
$dic2_item$	$D.{ m min_key1}()$	returns the item with minimal first key.
$dic2_item$	$D.\min_{\mathbf{key2}}()$	returns the item with minimal second key.

 $D.insert(K1 \ k_1, \ K2 \ k_2, \ I \ i)$ $dic2_item$ associates the information i with the keys k_1 and k_2 . If there is an item $< k_1, k_2, j >$ in D then j is replaced by i, else a new item $\langle k_1, k_2, i \rangle$ is added to D. In both cases the item is returned. $D.\operatorname{lookup}(K1 \ k_1, \ K2 \ k_2)$ $dic2_item$ returns the item with keys k_1 and k_2 (nil if no such item exists in D). list(dic2_item) D.range_search(K1 a, K1 b, K2 c, K2 d) returns the list of all items $\langle k_1, k_2, i \rangle \in D$ with $a \leq k_1 \leq b$ and $c \leq k_2 \leq d$. returns the list of all items of D. $list(dic2_item)$ D.all_items() $D.\operatorname{del}(K1 \ k_1, \ K2 \ k_2)$ deletes the item with keys k_1 and k_2 voidfrom D. removes item it from D. $D.\text{del_item}(dic2_item\ it)$ voidPrecondition: it is an item in D. D.change_inf(dic2_item it, I i) voidmakes i the information of item it. Precondition: it is an item in D. D.clear()makes D the empty d2_dictionary. voidreturns true if D is empty, false otherwise. D.empty()bool

4. Iteration

int

for all_dic2_items(i, D) { "the items of D are successively assigned to i" }

5. Implementation

D.size()

Two-dimensional dictionaries are implemented by dynamic two-dimensional range trees based on BB[α] trees. Operations insert, lookup, del_item, del take time $O(\log^2 n)$, range_search takes time $O(k + \log^2 n)$, where k is the size of the returned list, key, inf, empty, size, change_inf take time O(1), and clear takes time $O(n \log n)$. Here n is the current size of the dictionary. The space requirement is $O(n \log n)$.

returns the size of D.

6.3 Sets of two-dimensional points (point_set)

An instance S of the data type $point_set$ is a collection of items (ps_item) . Every item in S contains a two-dimensional point as key (data type point), and an information from a data type I, called the information type of S. The number of items in S is called the size of S. A point set of size zero is said to be empty. We use < p, i > to denote the item with point p, and information i. For each point p there is at most one item $< p, i > \in S$. Beside the normal dictionary operations, the data type $point_set$ provides operations for rectangular range queries and nearest neighbor queries.

1. Declaration of a two-dimensional point set type

 $declare(point_set, I)$

introduces a new data type with name $point_set(I)$ consisting of all two-dimensional point sets with information type I.

2. Creation of a two-dimensional point set

 $point_set(I)$ S;

creates an instance S of type $point_set(I)$ and initializes S to the empty set.

3. Operations on a point_set S

point	$S.\text{key}(ps_item \ it)$	returns the point of item it. Precondition: it is an item in S.
I	$S.inf(ps_item\ it)$	returns the information of item it . Precondition: it is an item in S .
ps_item	S.insert(point p, I i)	associates the information i with point p . If there is an item $< p, j >$ in S then j is replaced by i , else a new item $< p, i >$ is added to S . In both cases the item is returned.
ps_item	S.lookup(point p)	returns the item with point p (nil if no such item exists in S).
$list(ps_item)$	$S.$ range_search $(real \ x_0, \ real \ x_1)$	$egin{aligned} x_0, \ real \ y_0, \ real \ y_1 \end{pmatrix} \ & ext{returns all items} < p, i > \in \ S \ ext{with} \ & ext{$x_0 \le p.xcoord}() \le x_1 \ ext{and} \ & ext{$y_0 \le p.ycoord}() \le y_1 \end{aligned}$

ps_item	$S.\mathtt{nearest_neighbor}(point\ q)$	returns the item $< p, i > \in S$ such that the distance between p and q is minimal.
void	S.del(point p)	deletes the item with point p from S
void	$S. ext{del_item}(ps_itemit)$	removes item it from S . Precondition: it is an item in S .
void	S.change_inf(ps_item it, I i)	makes i the information of item it .
		Precondition: it is an item in S.
$list(ps_item)$	$S.{ m all_items}()$	returns the list of all items in S .
list(point)	$S.\mathtt{all_points}()$	returns the list of all points in S .
void	S.clear()	makes S the empty point_set.
bool	$S.\mathtt{empty}()$	returns true iff S is empty.
int	S.size()	returns the size of S .

4. Iteration

 $forall_ps_items(i, S)$ { "the items of S are successively assigned to i" }

5. Implementation

Point sets are implemented by a combination of two-dimensional range trees and Voronoi diagrams. Operations insert, lookup, del_item, del take time $O(\log^2 n)$, key, inf, empty, size, change_inf take time O(1), and clear takes time $O(n \log n)$. A range_search operation takes time $O(k + \log^2 n)$, where k is the size of the returned list. A nearest_neighbor query takes time $O(n^2)$, if it follows any update operation (insert or delete) and $O(\log n)$ otherwise. Here n is the current size of the point set. The space requirement is $O(n^2)$.

6.4 Sets of intervals (interval_set)

An instance S of the data type $interval_set$ is a collection of items (is_item) . Every item in S contains a closed interval of the real numbers as key and an information from a data type I, called the information type of S. The number of items in S is called the size of S. An interval set of size zero is said to be empty. We use (x, y, i) to denote the item with interval [x, y] and information i, x (y) is called the left (right) boundary of the item. For each interval $[x, y] \subset \mathbb{R}$ there is at most one item $(x, y, i) \in S$.

1. Declaration of an interval set type

 $declare(interval_set, I)$

introduces a new data type with name $interval_set(I)$ consisting of all interval sets with information type I.

2. Creation of an interval set

 $interval_set(I)$ S;

creates an instance S of type $interval_set(I)$ and initializes S to the empty set.

3. Operations on a interval_set S

r	eal	$S.\operatorname{left}(is_item\ it)$	returns the left boundary of item it. Precondition: it is an item in S.
r	eal	$S.right(is_item\ it)$	returns the right boundary of item it. Precondition: it is an item in S.
Ι		$S.inf(is_item\ it)$	returns the information of item it. Precondition: it is an item in S.
i	s_item	S.insert(real x, real y, I i)	associates the information i with interval $[x,y]$. If there is an item $< x,y,j >$ in S then j is replaced by i , else a new item $< x,y,i >$ is added to S . In both cases the item is returned.
i	s_item	$S.lookup(real \ x, real \ y)$	returns the item with interval $[x, y]$ (nil if no such item exists in S).
l	$ist(is_item)$	S.intersection(real a, real b)	
			returns all items $< x, y, i > \in \ S$ with $[x,y] \cap [a,b] eq \emptyset.$

void	$S.del(real \ x, \ real \ y)$	deletes the item with interval $[x, y]$ from S .
void	$S. del_item(is_item it)$	removes item it from S .
		Precondition: it is an item in S .
void	$S.$ change_inf($is_item\ it,\ I\ i$)	makes i the information of item it .
		Precondition: it is an item in S .
void	$S.\mathtt{clear}()$	makes S the empty interval_set.
bool	$S. ext{empty}()$	returns true iff S is empty.
int	S.size()	returns the size of S .

4. Iteration

for all_is_items (i, S) { "the items of S are successively assigned to i"}

5. Implementation

Interval sets are implemented by two-dimensional range trees. Operations insert, lookup, del_item and del take time $O(\log^2 n)$, intersection takes time $O(k + \log^2 n)$, where k is the size of the returned list. Operations left, right, inf, empty, and size take time O(1), and clear $O(n \log n)$. Here n is always the current size of the interval set. The space requirement is $O(n \log n)$.

6.5 Sets of parallel segments (segment_set)

An instance S of the data type $segment_set$ is a collection of items (seg_item). Every item in S contains as key a line segment with a fixed direction α (see data type segment) and an information from a data type I, called the information type of S. α is called the orientation of S. We use < s, i > to denote the item with segment s and information i. For each segment s there is at most one item $< s, i > \in S$.

1. Declaration of a segment set type

 $declare(segment_set, I)$

introduces a new data type with name $segment_set(I)$ consisting of all segment sets with information type I.

2. Creation of a segment set

- a) $segment_set(I)$ $S(real \alpha);$
- b) $segment_set(I)$ S;

creates an empty instance S of type $segment_set(I)$ with orientation α . Variant b) creates a segment set of orientation zero, i.e., for horizontal segments.

3. Operations on a segment_set S

segment	$S.\text{key}(seg_item\ it)$	returns the segment of item it . Precondition: it is an item in S .
I	$S.inf(seg_item\ it)$	returns the information of item it . Precondition: it is an item in S .
seg_item	$S.insert(segment \ s, \ I \ i)$	associates the information i with segment s . If there is an item $< s, j >$ in S then j is replaced by i , else a new item $< s, i >$ is added to S . In both cases the item is returned.
ps_item	S.lookup(segment s)	returns the item with segment s (nil if no such item exists in S).
$list(seg_item)$	S. intersection (segment q)	returns all items $\langle s, i \rangle \in S$ with $s \cap q \neq \emptyset$. Precondition: q is orthogonal to the segments in S.

$list(seg_item)$	$S.intersection(line\ l)$	returns all items $\langle s,i \rangle \in S$ with $s \cap l \neq \emptyset$. Precondition: l is orthogonal to the segments in S .
void	$S.\mathrm{del}(segment \ s)$	deletes the item with segment s from S .
void	$S. ext{del_item}(seg_itemit)$	removes item it from S . Precondition: it is an item in S .
void	S.change_inf(seg_item it, I	i) makes i the information of item it. Precondition: it is an item in S.
void	$S.\mathtt{clear}()$	makes S the empty segment_set.
bool	$S.\mathtt{empty}()$	returns true iff S is empty.
int	$S.\mathrm{size}()$	returns the size of S .

4. Iteration

for all_seg_items (i, S) { "the items of S are successively assigned to i" }

5. Implementation

Segment sets are implemented by dynamic segment trees based on $BB[\alpha]$ trees. Operations key, inf, change_inf, empty, and size take time O(1), insert, lookup, del, and del_item take time $O(\log^2 n)$ and an intersection operation takes time $O(k + \log^2 n)$, where k is the size of the returned list. Here n is the current size of the set. The space requirement is $O(n \log n)$.

6.6 Planar Subdivisions (subdivision)

An instance S of the data type subdivision is a subdivision of the two-dimensional plane, i.e., an embedded planar graph with straight line edges (see also sections 5.3 and 5.6). With each node v of S is associated a point, called the position of v and with each face of S is associated an information from a data type I, called the information type of S.

1. Declaration

declare(subdivision, I)

introduces a new data type with name subdivision(I) consisting of all planar subdivisions with information type I. Precondition: The data type GRAPH(point, I) has been declared before.

2. Creation of a subdivision

```
subdivision(I) S(GRAPH(point, I) G);
```

creates an instance S of type subdivision(I) and initializes it to the subdivision represented by the parameterized directed graph G. The node entries of G (of type point) define the positions of the corresponding nodes of S. Every face f of S is assigned the information of one of its bounding edges in G. Precondition: G represents a planar subdivision, i.e., a straight line embedded planar map.

2. Operations on a subdivision S

point	$S.\mathtt{position}(node v)$	returns the position of node v .
ftype	S.inf(face f)	returns the information of face f .
face	$S.locate_point(point p)$	returns the face containing point p .

3. Implementation

Planar subdivisions are implemented by parameterized planar maps and an additional data structure for point location. Operations position and inf take constant time, a locate_point operation takes time $O(\log^2 n)$. Here n is the number of nodes. The space requirement and the initialization time is $O(n^2)$.

6.7 Graphic Windows (gwindow)

The data type gwindow provides an interface for the input and output of basic geometric objects in the plane (see section 5.1) through a graphic window on a SUN workstation. In the current implementation only the sunview (suntools) window system is supported, the include file is <LEDA/sunview.h>. Application programs must be started from a sunview (suntools) window and have to be linked with the libP.a, libG.a, libL.a, suntool, sunwindow, pixrect, and m libraries (see section 1.9).

An instance W of the data type gwindow is an iso-oriented rectangular window in the two-dimensional plane. Its size and position are defined by three real numbers: x_0 , the x-coordinate of the left side, x_1 , the x-coordinate of the right side, and y_0 , the y-coordinate of the bottom side. W is displayed on the screen as a sunview window, initially a 800×800 pixel square positioned in the upper right corner. The y-coordinate of the top side of W is determined by the current size and shape of the window on the screen, which can be changed interactively. A graphic window supports operations for drawing points, lines, segments, arrows, circles, polygons, graphs, ... and for graphical input of all these objects using the mouse input device. Most of the drawing operations have an optional color argument. Possible colors are black (default), white, blue, green, red, violet, and orange. On monochrome displays all colors different from white are turned to black. There are 6 parameters used by the drawing operations:

- 1. The line width parameter (default value 1 pixel) defines the width of all kinds of lines (segments, arrows, edges, circles, polygons).
- 2. The line style parameter defines the style of lines. Possible line styles are solid (default), dashed, and dotted.
- 3. The node width parameter (default value 10 pixels) defines the diameter of nodes created by the draw_node and draw_filled_node operations.
- 4. The text mode parameter defines how text is inserted into the window. Possible values are transparent (default) and opaque.
- 5. The drawing mode parameter defines the logical operation that is used for setting pixels in all drawing operations. Possible values are src_mode (default) and xor_mode . In src_mode pixels are set to the respective color value, in xor_mode the value is bitwise added to the current pixel value.
- 6. The redraw function parameter is used to redraw the entire window, whenever a redrawing is necessary, e.g., if the window shape on the screen has been changed. Its type is pointer to a void-function taking no arguments, i.e., void (*F)();

1. Creation of a graphic window

- a) $gwindow W(real x_0, real x_1, real y_0);$
- b) $gwindow W(real x_0, real x_1, real y_0, int d);$
- c) gwindow W;

creates a graphic window W with lower left corner (x_0, y_0) and lower right corner (x_1, y_0) . Variant b) takes an additional integer argument d to define a rectangular grid with integer coordinates of distance d. In this case the mouse cursor can only take grid point positions. Variant c) initializes W to a default sized window $(x_0 = 0, x_1 = 100, y_0 = 0)$. The *init* operation (see below) can always be used to change the window coordinates.

2. Operations

2.1 Initialization

void W.init(real x_0 , real x_1 , real y_0)

W is made a gwindow with lower left corner (x_0, y_0) and lower right corner (x_1, y_0) (like creation a).

void W.init(real x_0 , real x_1 , real y_0 , int d)

W is made a gwindow with lower left corner (x_0, y_0) and lower right corner (x_1, y_0) with a rectangular grid with integer coordinates of distance d (like creation b).

void W.clear() W is erased.

2.2 Setting parameters

int $W.set_line_width(int pix)$

Sets the line width parameter to pix pixels and returns its previous value.

line_style W.set_line_style(linestyle s)

Sets the line style parameter to s and returns its previous value.

int W.set_node_width(int pix)

Sets the node width parameter to pix pixels and returns its previous value.

text_mode W.set_text_mode(text_mode m)

Sets the text mode parameter to m and returns its previous value.

draw_mode W.set_mode(draw_mode m)

Sets the drawing mode parameter to m and returns its previous value.

void $W.set_redraw(void (*F)())$

Sets the redraw function parameter to F.

2.3 Reading parameters and window coordinates

int W.get_line_width() returns the current line width.

line_style W.get_line_style() returns the current line style.

int W.get_node_width() returns the current node width.

draw_mode W.get_text_mode() returns the current text mode.

draw_mode W.get_mode() returns the current drawing mode.

real W.xmin() returns x_0 , the minimal x-coordinate of W.

real W.ymin() returns y_0 , the minimal y-coordinate of W.

real W.xmax() returns x_1 , the maximal x-coordinate of W.

real W.ymax() returns y_1 , the maximal y-coordinate of W.

real W.scale() returns the number of pixels of a unit length

line segment.

2.4 Drawing points

void $W.draw_point(real x, real y, color c = black)$

draws the point (x, y) as a cross of a vertical and a horizontal segment intersecting at (x, y).

void $W.draw_point(point p, c = black)$

draws point (p.xcoord(), p.ycoord()).

void W.draw(point p, c = black)

same as $draw_point(p,c)$.

2.5 Drawing line segments

void W.draw_segment(real x_1 , real y_1 , real x_2 , real y_2 , color c = black)
draws a line segment from (x_1, y_1) to (x_2, y_2) .

void W.draw_segment(point p, point q, color c = black)

draws a line segment from point p to point q.

void W.draw_segment(segment s, color c = black)

draws line segment s.

void W.draw(segment s, c = black)
same as draw_segment(s,c).

2.6 Drawing lines

woid $W.draw_line(real \ x_1, \ real \ y_1, \ real \ x_2, \ real \ y_2, \ color \ c = black)$ draws a straight line passing through points (x_1, y_1) and (x_2, y_2) .

void W.draw_line(point p, point q, color c = black)

draws a straight line passing through points p and q.

void W.draw_hline(real y, color c = black)

draws a horizontal line with y-coordinate y.

void $W.draw_vline(real \ x, \ color \ c = black)$ draws a vertical line with x-coordinate x.

void W.draw_line(line l, color c = black)

draws line l.

void $W.draw(line \ l, \ c = black)$ same as $draw_line(l,c)$.

2.7 Drawing arrows

void W.draw_arrow(real x_1 , real y_1 , real x_2 , real y_2 , color c = black)

draws an arrow pointing from (x_1, y_1) to (x_2, y_2) .

void W.draw_arrow(point p, point q, color c = black)

draws an arrow pointing from point p to point q.

void W.draw_arrow(segment s, color c = black)

draws an arrow pointing from s.start() to s.end().

2.8 Drawing circles

void W.draw_circle(real x, real y, real r, color c = black)

draws the circle with center (x, y) and radius r.

woid $W.draw_circle(point p, real r, color c = black)$ draws the circle with center p and radius r.

woid $W.draw_circle(circle\ C,\ color\ c = black)$ draws circle C.

void $W.draw(circle\ C,\ c = black)$ same as $draw_circle(C,c)$.

2.9 Drawing discs

void W.draw_disc(real x, real y, real r, color c = black)

draws a filled circle with center (x, y) and radius r.

void W.draw_disc(point p, real r, color c = black)

draws a filled circle with center p and radius r.

void W.draw_disc(circle C, color c = black)

draws filled circle C.

2.10 Drawing polygons

void W.draw_polygon(list(point) lp, color c = black) draws the polygon with vertex sequence lp.

woid $W.draw_polygon(polygon P, color c = black)$ draws polygon P.

void W.draw(polygon P, c = black) same as $draw_polygon(P,c)$.

void $W.draw_filled_polygon(list(point) lp, color <math>c = black)$ draws the filled polygon with vertex sequence lp.

woid $W.draw_filled_polygon(polygon P, color c = black)$ draws filled polygon P.

2.11 Drawing functions

$$woid$$
 W.plot_xy(real x_0 , real x_1 , (real)(*F)(real), color $c = black$)

draws function F in range $[x_0, x_1]$, i.e., all points

 (x, y) with $y = F(x)$ and $x_0 \le x \le x_1$
 $woid$ W.plot_yx(real y_0 , real y_1 , (real)(*F)(real), color $c = black$)

draws function F in range $[y_0, y_1]$, i.e., all points

 (x, y) with $x = F(y)$ and $y_0 \le y \le y_1$

2.12 Drawing text

2.13 Drawing nodes

Nodes are circles of diameter node_width.

$$W.draw_node(real\ x_0,\ real\ y_0,\ color\ c = black)$$
 draws a node at position $(x_0,y_0).$
 $Void\ W.draw_node(point\ p,\ color\ c = black)$ draws a node at position $p.$
 $Void\ W.draw_filled_node(real\ x_0,\ real\ y_0,\ color\ c = black)$ draws a filled node at position $(x_0,y_0).$
 $Void\ W.draw_filled_node(point\ p,\ color\ c = black)$ draws a filled node at position $p.$
 $Void\ W.draw_text_node(real\ x,\ real\ y,\ string\ s,\ color\ c = black)$ draws a node filled with string s at position $(x_0,y_0).$
 $Void\ W.draw_text_node(point\ p,\ string\ s,\ color\ c = black)$

2.14 Drawing edges

Edges are straigth line segments or arrows with a clearance of node_width/2 at each end.

void W.draw_edge(real x_1 , real y_1 , real x_2 , real y_2 , color c = black)
draws an edge from (x_1, y_1) to (x_2, y_2) .

void W.draw_edge(point p, point q, color c = black)

draws an edge from p to q.

woid $W.draw_edge(segment s, color c = black)$ draws an edge from s.start() to s.end().

void W.draw_edge_arrow(real x_1 , real y_1 , real x_2 , real y_2 , color c = black)
draws a directed edge from (x_1, y_1) to (x_2, y_2) .

woid $W.draw_edge_arrow(point p, point q, color c = black)$ draws a directed edge from p to q.

void W.draw_edge_arrow(segment s, color c = black)

draws a directed edge from s.start() to s.end().

2.15 Mouse Input

int W.read_mouse() displays the mouse cursor until a button is pressed.

Returns integer 1 for the left, 2 for the middle, and
3 for the right button (-1,-2,-3, if the shift key is pressed simultaneously).

int W.read_mouse(real& x, real& y)

displays the mouse cursor on the screen until a button is pressed. When a button is pressed the current position of the cursor is assigned to to (x, y) and the pressed button is returned.

int W.read_mouse_seg(real x_0 , real y_0 , real& x, real& y)

displays a line segment from (x_0, y_0) to the current cursor position until a mouse button is pressed. When a button is pressed the current position is assigned to (x, y) and the pressed button is returned.

int W.read_mouse_rect(real x_0 , real y_0 , real& x, real& y)

displays a rectangle with diagonal from (x_0, y_0) to the current cursor position until a mouse button is pressed. When a button is pressed the current position is assigned to (x, y) and the pressed button is returned.

int W.read_mouse_circle(real x_0 , real y_0 , real& x, real& y)

displays a circle with center (x_0, y_0) passing

through the current cursor position until a mouse
button is pressed. When a button is pressed the

current position is assigned to (x, y) and the

pressed button is returned.

 $W.confirm(string \ s)$ displays string s and asks for confirmation. Returns true iff the answer was "yes".

bool W.acknowledge(string s)

displays string s and asks for acknowledgement.

bool W.message(string s) displays s (each call adds a new line).

bool W.del_message() deletes the text written by all previous message operations.

2.16 Input and output operators

For input and output of basic geometric objects in the plane such as points, lines, line segments, circles, and polygons the << and >> operators can be used. Similar to C++ input streams gwindows have an internal flag indicating whether there is more input to read or not. Its initial value is true and it is turned to false if an input sequence is terminated by clicking the right mouse button (similar to ending istream input by the eof-character ctrl-D). In conditional statements objects of type gwindow are automatically converted to boolean by simply returning this internal flag. Thus, they can be used in conditional statements exactly in the same way as C++ input streams. For example, to read a sequence of points terminated by a right button click, use "while $(W >> p) \{ \dots \}$ ".

2.16.1 Output

gwindow &	W <<	circle C	${\rm like}\ W.{\rm draw_circle}(C)$
gwindow &	W<<	polygon P	${\rm like}\ W.{\rm draw_polygon}(P)$

2.16.2 Input

gwindow &	W >> p	reads a point p : clicking the left button assigns the current cursor position to p .
gwindow&	W>>s	reads a segment s : use the left button to input the start and end point of s .
gwindow &	W>>l	reads a line l : use the left button to input two different points on l
gwindow &	W >> C	reads a circle C : use the left button to input the center of C and a point on C
gwindow &	W >> P	reads a polygon P : use the left button to input the sequence of vertices of P , end the sequence by clicking the middle button.

As long as an input operation has not been completed the last read point can be erased by simultaneously pressing the shift key and the left mouse button.

7. Miscellaneous

This section describes some additional useful data types, functions and macros of LEDA. They can be used in any program that includes the <LEDA/basic.h> header file.

7.1 File input streams (file_istream)

An instance I of the data type $file_istream$ is an C++ istream bound to a file F, i.e., all input operations or operators applied to I read from F.

1. Creation of a file input stream

```
file_istream in(string s);
```

creates an instance in of type file_istream bound to the file with name s.

2. Operations

All input operations and operators (>>) defined for C++ istreams can be applied to file input streams as well.

7.2 File output streams (file_ostream)

An instance O of the data type $file_ostream$ is an C++ ostream bound to a file F, i.e., all output operations or operators applied to O write to F.

1. Creation of a file output stream

```
file_ostream out(string s);
```

creates an instance out of type file_ostream bound to the file with name s.

2. Operations

All output operations and operators (<<) defined for C++ ostreams can be applied to file output streams as well.

7.3 Some useful functions

int	read_int(string $s = "")$	prints s and reads an integer
char	$read_char(string s = "")$	prints s and reads a character
real	read_real(string s = "")	prints s and reads a real number
string	read_string(string s = "")	prints s and reads a string
bool	Yes(string s = "")	returns $(read_char(s) == 'y')$
void	init_random()	initializes the random number generator.
real	random()	returns a real valued random number in $[0,1]$
int	random(int a, int b)	returns a random integer in $[ab]$
real	used_time()	returns the currently used cpu time in seconds.
real	$used_time(real\&\ T)$	returns the cpu time used by the program from
		T up to this moment and assings the current
		time to T .
void	print_statistics()	prints a summary of the currently used memory

7.4 Macros

newline	cout << "\n"
forever	for(;;)
loop(a,b,c)	for $(a = b; a <= c; a + +)$
$in_range(a,b,c)$	(b <= a && a <= c)
Max(a,b)	((a > b) ? a : b)
Min(a,b)	((a > b) ? b : a)

7.5 Error Handling

LEDA tests the preconditions of many (not all!) operations. Preconditions are never tested, if the test takes more than constant time. If the test of a precondition fails an error handling routine is called. It takes an integer error number i and a char* error message string s as arguments. It writes s to the diagnostic output (cerr) and terminates the program abnormally if $i \neq 0$.

Users can provide their own error handling function handler by calling

set_error_handler(handler).

After this statement handler is used instead of the default error handler. handler must be a function of type void handler(int, char*). The parameters are replaced by the error number and the error message respectively.

8. Programs

8.1 Graph and network algorithms

In this section we list the C++ sources for some of the graph algorithms in the library (cf. section 5.12).

Depth First Search

```
#include <LEDA/graph.h>
#include <LEDA/stack.h>
declare(stack, node)
list(node) DFS(graph&G, node v, node_array(bool)&reached)
{
  list(node) L;
  stack(node) S;
  node w;
  if (! reached[v])
     \{ reached[v] = true; 
       L.append(v);
       S.push(v);
  while (!S.empty())
     \{ v = S.pop();
       forall\_adj\_nodes(w, v)
          if (!reached[w])
          \{ reached[w] = true; 
            L.append(w);
            S.\operatorname{push}(w);
      }
  return L;
}
```

Breadth First Search

```
#include <LEDA/graph.h>
#include <LEDA/queue.h>
declare(queue, node)
void BFS(graph& G, node v, node_array(int)& dist)
{
  queue(node) Q;
  node w;
  forall_nodes(w, G) dist[w] = -1;
  dist[v] = 0;
  Q.append(v);
  while (!Q.empty())
    \{ v = Q.pop();
      forall\_adj\_nodes(w, v)
         if (dist[w] < 0)
         \{Q.append(w);
           dist[w] = dist[v] + 1;
          }
     }
}
Connected Components
#include <LEDA/graph.h>
int COMPONENTS (ugraph& G, node_array(int)& compnum)
{
  node v, w;
  list(node) S;
  int count = 0;
  node_array(bool) reached(G, false);
  for all_nodes (v, G)
     if (!reached[v])
       \{ S = \mathrm{DFS}(G, v, reached); \}
         forall (w, S) compnum[w] = count;
         count + +;
        }
  return count;
```

Depth First Search Numbering

```
#include <LEDA/graph.h>
int dfs_count1, dfs_count2;
void df_s(node v, node_array(bool)& S, node_array(int)& df snum,
                                        node_array(int)& compnum,
                                       list(edge) T)
{ // recursive DFS
  node w;
  edge e;
  S[v] = true;
  df snum[v] = + + df s\_count1;
  forall_adj_edges (e, v)
     \{ w = G.target(e); 
       if (!S[w])
         \{ T.append(e); 
           d.f.s(w, S, dfsnum, compnum, T);
      }
  compnum[v] = + + df s\_count2;
 }
list(edge) DFS_NUM(graph& G, node_array(int)& dfsnum, node_array(int)& compnum
{
  list(edge) T;
  node\_array(bool) reached(G, false);
  node v;
  dfs\_count1 = dfs\_count2 = 0;
  for all_nodes (v, G)
     if (!reached[v]) d.f.s(v,reached,dfsnum,compnum,T);
  return T;
 }
```

Topological Sorting

```
#include <LEDA/graph.h>
bool TOPSORT(graph& G, node_array(int)&ord)
  node\_array(int) INDEG(G);
  list(node) ZEROINDEG;
  int count = 0;
  node v, w;
  edge e;
  for all_nodes (v, G)
     if ((INDEG[v]=G.indeg(v))==0) ZEROINDEG.append(v);
  while (!ZEROINDEG.empty())
    \{ v = ZEROINDEG.pop(); 
      ord[v] = + + count;
      forall\_adj\_nodes(w, v)
         if (--INDEG[w]==0) ZEROINDEG.append(w);
     }
  return (count==G.number_of_nodes());
 }
//TOPSORT1 sorts node and edge lists according to the topological ordering:
node_array(int) ord;
int cmp_node_ord(node& v, node& w)
\{ \text{ return } ord[v] - ord[w]; \}
int cmp_edge_ord(edge& e1, edge& e2)
{ return cmp_node_ord(target(e1), target(e2)); }
bool TOPSORT1(graph& G)
{
   ord.init(G);
   if (TOPSORT(G,ord))
   { G.sort_nodes(cmp_node_ord);
     G.sort\_edges(cmp\_edge\_ord);
    return true;
   return false;
```

Strongly Connected Components

```
#include <LEDA/array.h>
declare(array,node)
int STRONG_COMPONENTS(graph& G, node_array(int)& compnum)
{
  node v, w;
  int n = G.number_of_nodes();
  int count = 0;
  int i;
  array(node) V(1,n);
  list(node) S;
  node\_array(int) \ dfs\_num(G), compl\_num(G);
  node_array(bool) reached(G, false);
  DFS_NUM(G, dfs_num, compl_num);
  for all_nodes (v, G) V[compl_num[v]] = v;
  G.rev();
  for (i = n; i > 0; i - -)
     if (!reached[V[i]])
       \{ S = \mathrm{DFS}(G, V[i], reached); \}
         forall (w, S) compnum[w] = count;
         count + +;
        }
  return count;
 }
```

Dijkstra's Algorithm

```
#include <LEDA/graph.h>
void DIJKSTRA(graph& G, node s, edge_array(int)& cost,
                  node_array(int)& dist, node_array(edge)& pred)
\{ \text{ node-pq(int) } PQ(G); 
  int c;
  node u, v;
  edge e;
  forall_nodes(v, G)
     \{ pred[v] = 0;
       dist[v] = infinity;
       PQ.insert(v, dist[v]);
      }
  dist[s] = 0;
  PQ.decrease\_inf(s, 0);
  while (! PQ.empty())
     \{ u = PQ. delete\_min() \}
       forall_adj_edges(e, u)
         \{ v = G.target(e); 
            c = dist[u] + cost[e];
           if (c < dist[v])
              \{ dist[v] = c;
                pred[v] = e;
                PQ.decrease_inf(v, c);
           } /* forall_adj_edges */
      } /* while */
}
```

Bellman/Ford Algorithm

```
#include <LEDA/graph.h>
#include <LEDA/queue.h>
declare(queue, node)
bool BELLMAN_FORD(graph& G, node s, edge_array(int)& cost,
                        node_array(int)& dist, node_array(edge)& pred)
{ node_array(bool) in_Q(G, false);
  node\_array(int) count(G, 0);
  int n = G.number_of_nodes();
   queue(node) Q(n);
  node u, v;
  edge e;
  int c;
   forall_nodes (v,G) { pred[v] = 0;
                          dist[v] = infinity;
   dist[s] = 0;
   Q.append(s);
   in_{-}Q[s] = true;
   while (!Q.empty())
     \{ u = Q.pop();
       in_{-}Q[u] = false;
       if (+ + count[u] > n) return false; //negative cycle
       forall_adj_edges (e, u)
          \{ v = G.target(e); 
          c = dist[u] + cost[e];
          if (c < dist[v])
            \{ dist[v] = c;
              pred[v] = e;
              if (!in_Q[v])
                \{Q.append(v);
                  in_{-}Q[v] = true;
             }
           } /* forall_adj_edges */
      } /* while */
  return true;
```

All Pairs Shortest Paths

```
#include <LEDA/graph.h>
void all_pairs_shortest_paths(graph& G, edge_array(real)& cost,
                                       node_matrix(real)& DIST)
{
  // computes for every node pair (v, w) DIST(v, w) = cost of the least cost
  // path from v to w, the single source shortest paths algorithms BELLMAN_FORD
  // and DIJKSTRA are used as subroutines
  edge e;
  node v;
  real C=0;
 forall_edges(e, G) C += fabs(cost[e]);
  node s = G.new\_node();
                                         // add s to G
  forall\_nodes(v, G) G.new_edge(s, v);
                                         // add edges (s, v) to G
  node\_array(real) \ dist1(G);
  node\_array(edge) pred(G);
  edge\_array(real) cost1(G);
  forall\_edges(e, G) cost1[e] = (G.source(e) == s) ? C : cost[e];
  BELLMAN_FORD(G, s, cost1, dist1, pred);
  G.del\_node(s);
                                          // delete s from G
  edge\_array(real) cost2(G);
  forall\_edges(e, G) \ cost2[e] = dist1[G.source(e)] + cost[e] - dist1[G.target(e)];
  forall_nodes(v, G) DIJKSTRA(G, v, cost2, DIST[v], pred);
  for all nodes(v, G)
    forall_nodes(w, G) DIST(v, w) = DIST(v, w) - dist1[v] + dist1[w];
}
```

Minimum Spanning Tree

```
#include <LEDA/graph.h>
edge\_array(real)* C;
int cmp_edges(edge& e1, edge& e2)
{ return (*C)[e1] - (*C)[e2]; }
void MIN_SPANNING_TREE(graph& G, edge_array(real)& cost, list(edge)& EL)
  node v, w;
  edge e;
  node_partition Q(G);
  list(edge) OEL = G.all\_edges();
  C = \& cost;
  OEL.sort(cmp_edges);
   EL.clear();
  forall(e, OEL)
     \{ v = G.source(e); 
       w = G.target(e);
       if (!(Q.same\_block(v, w))
         \{Q.union\_blocks(v, w);
           EL.append(e);
          }
      }
}
```

9. Tables

9.1 Data Types

array	21	node_partition	64
array2	22	node_pq	65
b_priority_queue	39	node_set	63
b_queue	26	partition	34
b_stack	25	planar_map	54
bool	15	point	73
circle	79	point_set	84
d2_dictionary	82	polygon	78
	42		37
d_array		priority_queue	
dictionary	40	PLANAR_MAP	59
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