CS335 Class Notes

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1 Lecture 1: C++ basics

1.1 C++ classes

```
class IntCell {
    public:
        // This is a constructor
        IntCell() { _value = 0; }
        // This is another constructor. It takes an initial value.
        IntCell( int initialValue ) { _value = initialValue; }
        // This is an accessor (or getter) method.
        int read() { return _value; }
        // This is a setter method.
        void write( int newValue ) { _value = newValue; }
    private:
        int _value;
};
int main() {
    // Using the first constructor w/o parameters
    IntCell x;
    int xValue = x.read(); // 0
    x.write(5);
    xValue = x.read(); // 5
    // Using the first constructor with a single parameter
    IntCell y(12);
    int yValue = y.read(); // 12
    y.write(y.read() * 2);
    yValue = y.read();
                         // 24
}
```

The keywords **public** and **private** determine the visibility of class members. A member that is **public** may be accessed by any method in any class while a member that is **private** may only be accessed by methods in the class it's declared. Date members are usually declared **private** to hide a class's internal details (i.e., **information hiding**).

1.2 Initialization list

Instead of initializing data members inside a constructor's body, an **initialization list** can be used to do so right in the constructor's signature:

By default C++ does behind-the-scenes type conversions in all one-parameter constructors. Thus, to avoid this the keyword explicit is used.

With implicit type conversion,

```
IntCell obj;
obj = 37;
```

means

```
IntCell temp = 37;
obj = temp;
```

With **explicit type conversion** (by using explicit), a "type mismatch" compiler error is thrown instead. The use of explicit means a one-parameter constructor cannot be used to generate an implicit temporary object.

1.3 Constant member functions

A member function that examines but does not change the state of its object is an **accessor**. In the IntCell, read is an accessor and thus it's marked explicitly as an accessor by using the keyword **const** after the closing parenthesis that ends the parameter list. This signals that this method doesn't change the state of an IntCell and if it tries the compiler complains.

1.4 Interface and implementation

An **interface** lists the class and its members (both data and methods) and the **implementation** provides the implementations of the methods.

In C++, this is done by placing the interface in a .h file and the implementation in a .cpp file. Source code that requires knowledge of the interface must include the interface file (e.g., #include Interf

.h). To avoid including files multiple times, a few preprocessor commands are used in the interface file.

```
#ifndef INTCELL_H // if INTCELL_H is not defined...
#define INTCELL_H // ...define it.

// Here's the IntCell class's interface
#endif // end the definition
```

1.4.1 IntCell interface

```
// IntCell.h
#ifndef INTCELL_H
#define INTCELL_H

class IntCell {
    public:
        explicit IntCell( int initialValue = 0 ) {}
        int read() const { return _value; }
        void write( int newValue ) { _value = newValue; }

private:
    int _value;
};
#endif
```

1.4.2 IntCell implementation

```
// IntCell.cpp
#include "IntCell.h" // we're making use IntCell's interf. so we include
   it.

IntCell::IntCell( int initialValue = ) : _value { initialValue } { }

// Return the stored value.
int IntCell::read() const { return _value; }

// Change the stored value.
void IntCell::write( int newValue ) { _value = newValue; }

};
```

Notice that we must use the **scope resolution operator** (::) to identify the class each member function belongs to. Otherwise, it's assumed that the function is in the **global scope**. Also note that the signature of an implemented member function *must* match exactly the signature in the class interface.

1.4.3 Using IntCell

```
// main.cpp
#include "IntCell.h"

int main() {
    IntCell m{};
    m.write(5);
    int val = m.read(); // 5

    return 0;
}
```

To compile the program, run:

```
g++ -o prog main.cpp IntCell.cpp
```

The header files (e.g., IntCell.h) aren't listed since they're included in the implementation files and thus are compiled too.

1.5 C++ Objects

In C++ objects are declared much like primitive types (e.g., char, int, float, etc.):

```
IntCell obj1;  // Zero parameter constructor
IntCell obj2(12); // One parameter constructor
```

However, due to an effort to standardize the uniform initialization syntax using braces, it's recommended to declare objects as follows:

```
IntCell obj1;  // Zero parameter constructor
IntCell obj1{};  // same as before
IntCell obj2{12};  // One parameter constructor
```

1.6 vectors and strings

The C++ standard defines the classes vector and string that intend to replace the built-in C++ array. Unlike arrays, these classes behave like first-class objects. Thus, they can be copied with =, they remember how many items they can store, and their indexing operator check that the provided valid (to access some element) is valid.

1.6.1 vector

```
vector<int> numbers = {1, 2, 3, 4, 5}; // Or...
vector<int> numbers {1, 2, 3, 4, 5};
```

However, there's some ambiguity with the declaration. For instance, vector<int> a(12) declares a vector that stores 12 integers but vector<int> a{12} declares a vector of size 1 with the value 12 at position 0. This is because C++ gives precedence to the initializer list. If the intention is to declare a vector of size 12, use the old C++ syntax using parentheses: vector<int> a(12).

Instead of using the regular for loop, vectors can be looped over using the following syntax:

```
int sum = 0;
vector<int> numbers = {1, 2, 3, 4, 5};
for( int num : numbers ) {
    sum += num;
}
```

The keyword auto can be used to let the compiler determine the type:

```
int sum = 0;
vector<int> numbers = {1, 2, 3, 4, 5};
for( int num : numbers ) {
    sum += num;
}
```

However, keep in mind that this range syntax is only appropriate when 1) accessing elements sequentially and 2) when the index isn't needed.

1.7 C++ details

1.7.1 Pointers

A **pointer** is a variable that stores the address of a memory location (i.e., where an object resides). The syntax for declaring a pointer is as follows:

```
char* ptr_c;  // ptr_c is a pointer-to-char
int *ptr_i;  // ptr_i is a pointer-to-int
float * ptr_f;  // ptr_f is a pointer-to-float
void *u_ptr;  // ptr_u is a untyped pointer
```

Note that the position of the asterisk * (known as the **indirection operator** in this context) doesn't matter.

In order to obtain an object's address, the **address-of** operator & is used and a pointer variable is used to store it:

```
int num = 5;
int *ptr_num = # // get the address of num
int val = *ptr_num; // get the value stored in the address ptr_num points
    to
*ptr_num = 6; // now num is also 6. after all ptr_num to the same
    memory location
```

Thus, the indirection operator has two jobs: 1) it declares a pointer and 2) it dereferences a pointer.

1.7.2 Dynamic object allocation and garbage collection

Objects are created dynamically by allocating memory with **new** which returns a pointer to the newly created object:

```
IntCell *m;
m = new IntCell(); // OK
m = new IntCell{}; // OK
m = new IntCell; // OK
```

If a pointer variable points at a class type, then a (visible) member of the object being pointed at can be accessed via the -> operator:

```
int x = m->read();
m->write(6);
```

When an object that is allocated by **new** is no longer referenced, the delete operation must be applied to the object (through a pointer). Otherwise, the memory that it consumes is lost (until the program terminates). This is known as a **memory leak**.

```
delete m;
```

1.7.3 Lvalues, and Rvalues

An **lvalue** is an expression that identifies a non-temporary object:

```
vector<int> arr(3);
const int x = 2;
int y;
int z = x + y;
vector<string> *ptr = &arr;
```

The expressions arr, x, y, z, ptr, *ptr, and z are all lvalues.

An **rvalue** is an expression that identifies 1) a temporary object or 2) a value (e.g., literal constant) not associated with any object:

```
vector<int> arr(3);
const int x = 2;
int y;
int z = x + y;
vector<string> *ptr = &arr;
```

Here, 2, x+y, and &arr are all rvalues.

1.7.4 References

A reference type allows us to define a new name for an existing value. In classic C++, a reference can generally only be a name for an Ivalue, since having a reference to a temporary would lead to the ability to access an object that has theoretically been declared as no longer needed, and thus may have had its resources reclaimed for another object. However, in C++11, we can two types of references:

• An **Ivalue reference** is declared by placing an & after some type. An Ivalue reference then becomes a synonym for the object it references.

```
string str = "hello";
string &str_r = str;
str_r += " world"; // now str is "hello world"

string &lit_r = "hi"; // ILLEGAL: "hi" is not modifiable
string &bad_r = str + "!"; // ILLEGAL: str + "!" isn't an lvalue
```

• An **rvalue reference** is declared by placing an && after some type. An rvalue reference has the same characteristics as an Ivalue reference except that, unlike an Ivalue reference, *an rvalue reference can also reference an rvalue* (i.e., a temporary).

1.7.5 Uses of Ivalue references

 Aliasing complicated names. We can rename objects that are too long and complicated to simpler names.

```
auto &whichList = theLists[ myhash( x, theLists.size( ) ) ];
if( find( begin( whichList ), end( whichList ), x ) != end( whichList ) )
    {
    return false;
}
whichList.push_back( x );
```

Note that simply writing auto whichList = theLists[myhash(x, theLists.size())]; wouldn't work because this would create a copy, and whichList.push_back(x); would be applied to the copy, not the original.

• **Range for loops.** By default, a range **for** loop cannot change the elements it iterates over, however taking a lvalue reference allows to modify those elements.

```
vector<int> numbers = {1, 2, 3};

for( auto &number : numbers ) {
    number++;
}
```

• **Avoiding a copy.** Given a function that returns an element of an array/vector, we could return a non-modifiable reference to that element instead of a copy. For instance, instead of

```
// @arr: A non-empty vector of strings.
// @return the maximum string in the @arr.
// Will abort() if @arris empty.
string FindMax1(constvector<string>&arr) {
    if (arr.empty()) abort();
    int max_index=0;
    for (inti =1; i < arr.size(); ++i) {
        if(arr[max_index]<arr[i]) { max_index = i; }
    }
    return arr[max_index];
}</pre>
```

we could have

```
// @arr: A non-empty vector of strings.
// @return the maximum string in the @arr.
// Will abort() if @arris empty.
const string &FindMax1(constvector<string>&arr) {
    if (arr.empty()) abort();
    int max_index=0;
    for (inti =1; i < arr.size(); ++i) {
        if(arr[max_index]<arr[i]) { max_index = i; }
    }
    return arr[max_index];
}</pre>
```

Syntax is needed in function declarations and returns to enable the passing and returning using references instead of copies. Notice the **const** keyword and & in the function's header.

2 Lecture 2: C++ Basics (cont.)

2.1 Parameter passing

Many languages, including C++, pass all parameters using **call-by-value**: the actual argument is copied into the formal parameter. However, this might be utterly inefficient if large complex objects are being passed since they're copied in their entirety. Historically C++ has had three ways to pass parameters:

• call-by-value: Useful to pass small objects that shouldn't be mutated by the function.

```
double average( double a, double b );
double x = 2.5, y = 3.5;
double z = average(x, y); // x and y remain unchanged

void swap( double a, double b ); // swap a and b
swap(x, y); // Not what's expected; x and y remain unchanged
```

• call-by-reference (call-by-lvalue-reference): Useful for all type of objects that may be changed by the function.

```
double x = 2.5, y = 3.5;
swap( &a, &b );
swap(x, y); // now x = 3.5 and y = 2.5
```

• call-by-constant-reference: Useful for large objects that are expensive to copy and that must not be changed by the function. For this, the parameter is declared as a reference and the keyword const is used to signal that it cannot be modified.

```
string randomItem( const vector<string> &arr ); // return a random item in
    arr
```

There's still another fourth way to pass parameters:

• **call-by-rvalue-reference:** Instead of copying a temporary object stored in an rvalue, a move is used; moving an object's state is easier than copying, as *it may involve just a simpler pointer change*. Functions know if a value is temporary or not based on their signature so the primary use for this type of parameter passing is overloading a function based on whether a parameter is an lyalue or rvalue.

This idiom is particularly useful for defining the behavor of = and in writing constructors.

2.2 Return passing

In C++, there are several mechanism for returning from a function:

• **return-by-value:** A copy of the object is returned which can be inefficient, however in C++11 return-by-value may still be efficient for large objects if the returned objects are rvalues.

The following are two versions of the function randomItem. The second version avoids the creation of a temporary LargeType object, but only if the caller accesses it with a constant reference:

• return-by-constant-reference: This avoid creating an immediate copy of an object. However, the caller must also use a constant reference to access the return value; otherwise, there will be still a copy. What does the constant reference mean? It means that we don't want to allow changes to be made by the caller by using the return value.

```
const LargeType & randomItem2( const vector<LargeType> &arr ) {
   return arr[ randomInt(0, arr.size() -1) ];
}

vector<LargeType> vec;
const LargeType &item3 = randomItem2(vec); // no copy
```

• **return-by-reference:** A reference is returned and the caller can modify the returned value. This is used in a few places to allow the caller of a function to have modifiable access to the internal data representation of a class.

2.3 C++11 std::swap and std::move

Given that copying large objects is expensive, C++11 allows the programmer to easily replace expensive copies with moves provided the object's class supports moves.

Take the following example of a swap function that swap its arguments by three copies:

```
void swap( vector<string> &x, vector<string> &y ) {
    vector<string> tmp = x;
    x = y;
    y = tmp;
}
```

In C++11, if the right-hand side of the assignment operator (or constructor) is an rvalue, then if the object supports moving, we can automatically avoid copies. In the example above, we know that vector supports moving so instead of copy operations we could do move operations. These could be done either by casting the right-hand side of an assignment to an rvalue reference or by using std:: move.

```
// Using type-casting
void swap( vector<string> &x, vector<string> &y ) {
    vector<string> tmp = static_cast<vector<string> &&>( x );
    x = static_cast<vector<string> &&>( y );
    y = static_cast<vector<string> &&>( tmp );
}

// Using std::move, equivalent to casting but more succint
void swap( vector<string> &x, vector<string> &y ) {
    vector<string> tmp = std::move(x);
    x = std::move(y);
    y = std::move(tmp);
}
```

NOTE: std::move doesn't move anything; rather, it makes a value (either Ivalue or rvalue) subject to

be moved.

It's worth noting that std::swap is already part of STL and works for any part:

```
vector<string> x;
vector<string> y;
std::swap(x, y); // x contains y's contents and y contains x's contents
```

2.4 The big five

In C++, classes come with five special functions already written for each class. These are the **destructor**, **copy constructor**, **move constructor**, **copy assignment operator**, and **move assignment operator**. In many cases, you accept the default behavior provided by the compiler can be relied on. Sometimes you cannot.

Let's assume the following interface for the IntCell class:

```
#ifndef INTCELL_H
#define INTCELL_H

class IntCell {
   public:
        explicit IntCell( int initialValue = 0 ) { value_ = new int{
            initialValue}; }
        int read() const {}
        void write( int x ) {}
        private:
        int *value_;
};

#endif
```

2.4.1 Destructor

This function is called whenever an object goes out of scope or is subjected to a delete operation. Typically, the only responsibility of the destructor is to free up any resources that were acquired during the use of the object. This includes calling delete for any corresponding news, closing any files that were opened, and so on.

```
IntCell::~IntCell() {
    delete value_;
}
```

2.4.2 Copy constructor and move constructor

These two constructors are required to construct a new object, initialized to the same state as another object of the same type.

- If the existing object is an **lvalue**, then it's a **copy constructor**.
- If the existing object is an **rvalue**, then it's a **move constructor**.

```
// Copy constructor, the parameter is an lvalue of the same type.
IntCell::IntCell( const IntCell &rhs ) {
    stored_ = new int{*rhs.value_};
}

// Move constructor, the parameter is an rvalue of the same type.
IntCell::IntCell( IntCell &&rhs ) : value_{rhs.value_} {
    rhs.value_ = nullptr;
}
```

When is either constructor called?

• During a declaration with initialization.

```
IntCell B = C;  // Copy constructor if C is lvalue; Move constructor is C
   is rvalue
IntCell B { C }; // same as above
```

- An object passed using call-by-value (rarely done).
- An object returned by value.

2.4.3 Copy assignment and move assignment (operator =)

The assignment operator is called when = is applied to two objects that have both been previously constructed. Given the expression lhs = rhs, then the state of rhs (right hand side) is copied into lhs (left hand side).

- If rhs is an **lvalue**, this is done using the copy assignment operator.
- If rhs is an **rvalue**, this is done using the move assignment operator.

```
// Copy assignment operator
IntCell & IntCell::operator=( const IntCell &rhs ) {
    if (this &= &rhs) {
        *value_ = *rhs.value_;
    }
    return *this;
}

// Move assignment operator
IntCell & IntCell::operator=( IntCell &&rhs ) {
    std::swap(value_, rhs.value_);
    return *this;
}
```

In C++11, the copy assignment operator could be implemented more idiomatically with a copy-and-swap idiom:

```
// Copy assignment operator
IntCell & IntCell::operator=( const IntCell &rhs ) {
    IntCell copy = rhs; // calls the copy constructor
    std::swap(*this, copy);
    return *this;
}
```

2.4.4 Defaults

It's often the case that the defaults "big five" are perfectly acceptable, so nothing need to be done. If a class consists of data members that are exclusively primitive types and objects for which the defaults make sense, the class defaults will usually make sense. Thus a class whose data members are **int**, **double**, string, and even vector<string> can accept defaults.

When do defaults fail? The defaults fail in a class that contains a data member that is a pointer:

- The default destructor does nothing to data members that are pointers. It's imperative that we delete them ourselves.
- The copy constructor and copy assignment operator both copy the value of the pointer (i.e., a
 memory address) rather than the object being pointed at. This means we end up with two class
 instances that contain pointers that point to the same objects being pointed at. This is known as
 shallow copy. However, we would typically expect a deep copy, in which a clone of the entire
 object is created.

Thus, as a result, when a class contains pointers as data members, and deep semantics are important, we typically must implement the destructor, copy assignment, and copy constructors ourselves. Doing so removes the move defaults, so we also must implement move assignment and the move constructor. As a general rule, either you accept the default for all five operations, or you should declare all five, and explicitly define, default (use the keyword default), or disallow each (use the keyword delete). Generally we will defined all five.

NOTE: If you write any of the big-five, it would be good practice to explicitly consider all the others, as the defaults may be invalid or inappropriate. Changing a default method is an all or nothing situation.

When don't the defaults work? The most common situation in which the defaults do not work occurs when a data member is a pointer type and the pointer is allocated by some object member function (such as the constructor). A problem that might arisse is the default copy assignment and copy constructor doing shallow copies. Another problem is **memory leak**; an object allocated by a constructor might remain allocated and it's never reclaimed.

2.5 Templates

An algorithm is **type independent** if the logic of the algorithm does not depend on the type of items it handles. When we write C++ code for a type-independent algorithm or data structure, it's preferable to write the code once rather than recode it for each different type. This is accomplished by using **templates**.

2.5.1 Function templates

A **function template** isn't an actual function, but instead a pattern for what could become a function. For example, the following is a function template:

```
/*
Return the maximum item in array a.
Assumes a.size() > 0.
Comparable object must provide operator< and operator=.
*/

template <typename Comparable>
const Comparable & findMax( const vector<Comparable> &a ) {
   int maxIndex = 0;
   for (int i = 1; i < a.size(); i++) {
      if (a[maxIndex] < a[i]) {
         maxIndex = i;
      }
   }
   return a[maxIndex];
}</pre>
```

Using the function template would look as follows:

```
vector<int> v1 = { 37, 12, 1, 89 };
vector<double> v2 = { 78.1, 89.8, 12.4, 1.1 };
vector<string> v3 = { 'hi', 'ha', 'ho', 'he' };
vector<IntCell> v4(75);

findMax(v1); // OK: Comparable = int
findMax(v2); // OK: Comparable = double
findMax(v3); // OK: Comparable = string
findMax(v4); // Illegal; IntCell doesn't implement operator< and thus not
a Comparable</pre>
```

Thus, one of the limitations of function templates is that object passed to the function need to implement whatever operator the function uses internally. However, instead of relying on "hardcoded" conditions, a function that perform the required operation could be passed alongside the object. This type of functions are known as **function objects**.

2.6 Function objects

Previously we implemented findMax as a function template but it was only limited to objects that have an operator < function defined. Instead, we need to rewrite findMax to accept as parameters an array of object and a comparison function that explains how to decide which of two objects is the larger and which is the smaller. Instead of relying on the array objects knowing how to compare themselves, we completely decouple this information from the object in the arrays.

How do we pass a functions as parameters though? One way is to define a class with no data and one member function, and pass an instance of the class. In fact, the function is passed by placing it

inside an object, which is known as a function object.

The following is the simplest implementation of the function object idea:

```
#include <iostream>
#include <vector>
#include <string>
// Generic findMax, with a function object.
// Precondition: a.size() > 0.
// Comparator object is assumed to implement the isLessThan method.
template <typename Object, typename Comparator>
const Object & findMax( const std::vector<Object> &arr, Comparator cmp ) {
    int maxIndex = 0;
    for (int i = 1; i < arr.size(); i++) {</pre>
        if (cmp.isLessThan(arr[maxIndex], arr[i])) {
            maxIndex = i;
        }
    }
    return arr[maxIndex];
};
class StringComparisonByLength {
    public:
        bool isLessThan( const std::string &lhs, const std::string &rhs )
            return lhs.length() < rhs.length();</pre>
        }
};
class StringComparisonCaseInsentitive {
    public:
        bool isLessThan( const std::string &lhs, const std::string &rhs )
            return std::strcasecmp(lhs.c_str(), rhs.c_str()) < 0;</pre>
        }
};
int main() {
    std::vector<std::string> greetings = {"hi", "hello", "bonjour", "HOLA"
       };
    std::cout << findMax( greetings, StringComparisonByLength{} ) << "\n";</pre>
    std::cout << findMax( greetings, StringComparisonCaseInsentitive{} )</pre>
       << "\n";
    return 0;
}
```

C++ function objects are implemented using this basic idea, but with some fancy syntax. First, instead of using a function with a name, we use operator overloading. Instead of the function being isLessThan, it is operator(). Second, when invoking operator(), cmp.operator()(x,y) can be shortened to cmp(x,y). Third, we can provide a version of findMax that works without a function object that uses a default ordering; the implementation uses the Standard Library function object template less (defined in header file functional) to generate a function object that imposes the normal default ordering.

The following is the more idiomatic implementation:

```
#include <iostream>
#include <vector>
#include <string>
// Generic findMax, with a function object.
// Precondition: a.size() > 0.
template <typename Object, typename Comparator>
const Object & findMax( const std::vector<Object> &arr, Comparator
   isLessThan ) {
    int maxIndex = 0;
    for (int i = 1; i < arr.size(); i++) {</pre>
        if (isLessThan(arr[maxIndex], arr[i])) {
            maxIndex = i;
        }
    }
    return arr[maxIndex];
};
// Generic findMax, using default ordering.
const Object &findMax( const vector<Object> &arr ) {
    return findMax(arr, less<0bject>{} );
class StringComparisonByLength {
    public:
        bool operator()( const std::string &lhs, const std::string &rhs )
            return lhs.length() < rhs.length();</pre>
        }
};
class StringComparisonCaseInsentitive {
    public:
        bool operator()( const std::string &lhs, const std::string &rhs )
            return std::strcasecmp(lhs.c_str(), rhs.c_str()) < 0;</pre>
        }
};
int main() {
    std::vector<std::string> greetings = {"hi", "hello", "bonjour", "HOLA"
       };
    std::cout << findMax( greetings, StringComparisonByLength{} ) << "\n";</pre>
    std::cout << findMax( greetings, StringComparisonCaseInsentitive{} )</pre>
        << "\n";
    std::cout << findMax( greetings ) << "\n";</pre>
    return 0;
}
```

2.6.1 Class templates

In its most simplest form, a class template works much like a function template. The following MultiCell is an implementation that is like IntCell, but works for any type Object, provided that Object has a zero-parameter constructor, a copy constructor, and a copy assignment operator.

Notice that

- Object is passed by constant reference.
- the default parameter for the constructor is not 0, because 0 might not be a valid Object. Instead, the default parameter is the result of constructing an Object with its zero-parameter constructor.

If we implement class templates as a single unit, then there is very little syntax baggage. Many class templates are, in fact, implemented this way because, currently, separate compilation of templates does not work well on many platforms. **Therefore, in many cases, the entire class, with its implementation, must be placed in a .h file**. Popular implementations of the STL follow this strategy.

2.7 Using matrices

This will be implemented by using a vector of vectors (e.g., a vector of **int** vectors).

```
#ifndef MATRIX H
#define MATRIX_H
#include <vector>
template <typename Object>
class Matrix {
    public:
        /*
         * Create _array as having rows entries each of type vector<0bject
         * We have a rows zero-length vectors of Object so each row is
         * to have cols columns. Thus this creates a two-dimensional array
         */
        Matrix( int rows, int cols ) : _array(rows) {
            for (auto &row : array) {
                row.resize(cols);
            }
        }
        Matrix( std::vector<std::vector<Object>> v ) : _array{ v } { }
        Matrix( std::vector<std::vector<Object>> &&v ) : _array{ std::move
           (v) } { }
         * Array indexing operator. This returns the row (a vector<0bject>
         * the index row.
        const std::vector<Object> & operator[]( int row ) const {
            return _array[row]
        int numrows() const {
            return _array.size();
        int numcols() const {
            return numrows() ? _array[0].size() : 0;
    private:
        /*
        A matrix is represented by _array, a vector of vector<Object>.
        */
        std::vector<std::vector<Object>> _array;
};
#endif
```

2.8 Class problems

1. What's 1 + 2 + 3 + ... + n? Prove it.

$$1 + 2 + 3 + \dots + n = \frac{n(n+1)}{2}$$

Proof by induction:

- Base case: When n = 1, then 1 = 1(1+1)/2 = 1.
- Inductive step: Since 1=(1*2)/2, the statement P(1) is true. Assume that P(k) is true for an arbitrary positive integer k. We show that P(k+1) is true. In other words,

$$1+2+3+\ldots+(k+1)=\frac{(k+1)(k+2)}{2}$$

Thus

$$\begin{aligned} 1+2+3+\cdots +(k+1) &= (1+2+3+\cdots +k) +(k+1) \\ &= \frac{k(k+1)}{2} +(k+1) \\ &= \frac{k(k+1)}{2} +(k+1) \\ &= \frac{(k+1)(k+2)}{2} \end{aligned}$$

Therefore, by the principle of mathematical induction, P(n) is true for every positive integer n.

2. What's $1 + 2 + 4 + \dots + 2^n$? Prove it.

$$n = 0 \Rightarrow 1 \Rightarrow 1$$

$$n = 1 \Rightarrow 1 + 2 \Rightarrow 3$$

$$n = 2 \Rightarrow 1 + 2 + 4 \Rightarrow 7$$

$$n = 3 \Rightarrow 1 + 2 + 4 + 8 \Rightarrow 15$$
...
$$n = n \Rightarrow 1 + 2 + \dots + 2^{n} \Rightarrow 2^{(n+1)} - 1$$

Thus,
$$1 + 2 + 4 + \dots + 2^n = 2^{(n+1)} - 1$$
.

Proof by induction:

- Base case: When n=0, then $1=2^{(0+1)}-1=1$. When n=1, then $2^0+2^1=3=2^{(1+1)}-1=3$.
- Inductive step: Since $1=2^{(0+1)}-1$, the statement P(1) is true. Assume that P(k) is true for an arbitrary positive integer k. We show that P(k+1) is true. In other words, we show that

$$1 + 2 + 4 + \dots + 2^{(k+1)} = 2^{(k+2)} - 1.$$

Thus,

$$\begin{split} 1+2+4+\cdots+2^{(k+1)} &= (1+2+4+\cdots+2^k)+2^{(k+1)}\\ &= 2^{(k+1)}-1+2^{(k+1)}\\ &= 2^{(k+1)}+2^{(k+1)}-1\\ &= 2\cdot 2^{(k+1)}-1\\ &= 2^{(k+2)}-1 \end{split}$$

Therefore, by the principle of mathematical induction, P(n) is true for every non-negative integer n.

3. If $A_0=1$ and $A_n=2A_{(n-1)}+1$, what's a closed formula for A_n ? Prove it.

$$\begin{split} A_1 &= 2 \cdot A_0 + 1 = 3 \\ A_2 &= 2 \cdot A_1 + 1 = 7 \\ A_3 &= 2 \cdot A_2 + 1 = 15 \\ A_4 &= 2 \cdot A_3 + 1 = 31 \\ \dots \\ A_n &= 2 \cdot A_{(n-1)} + 1 = 2^{(n+1)} - 1 \end{split}$$

3 Lecture 3: Algorithm analysis

An **algorithm** is a clearly defined set of simple instructions which must are to be followed in order to solve a particular problem.

3.1 Mathematical background

Algorithm analysis is grounded on mathematics and the definitions below set up a formal framework to study algorithms. These definitions try to establish a **relative order among functions**. Given two functions, there are usually points where one function is smaller than the other so it doesn't make sense to claim, for instance, f(N) < g(N). Instead, we compare their **relative rates of growth**.

Big-Oh T(N) = O(f(N)) if there are positive constants c and n_0 such that $T(N) \le c \cdot f(N)$ when $N \ge n_0$.

Informally, the growth rate T(N) is less than or equal to that f(N).

Big-Omega $T(N)=\Omega(g(N))$ if there are positive constants c and n_0 such that $T(N)\geq c\cdot g(N)$ when $N\geq n_0$.

Informally, the growth rate T(N) is greather than or equal to that g(N).

Big-Theta
$$T(N) = \Theta(h(N))$$
 if and only if $T(N) = O(h(N))$ and $T(N) = \Omega(h(N))$.

Informally, the growth rate T(N) equals the growth rate of h(N). This means that T(N) is eventually sandwiched between two different constant multiples of h(N).

Little-Oh T(N) = o(p(N)) if, for all positive constants c, there exist an n_0 such that $T(N) < c \cdot p(N)$ when $N > n_0$.

Informally, the growth rate T(N) is less than the growth rate of f(N). This is different from Big-Oh, given that Bog-Oh allows the possibility that the growth rates are the same.

NOTE: In the above definitions, both c and n_0 are constants and they cannot depend on N. If you see yourself saying "take $n_0=N$ " or "take $c=log_2N$ " in an alleged Big-Oh proof, then you need to start with choices of c and n_0 that are indepedent of N.

3.2 Notation

3.2.1 Big-Oh notation

When we say that T(N) = O(f(N)), we're guaranteeing that the function T(N) grows at a rate no faster than f(N); thus f(N) is an **upper bound** on T(N). Alternatively, we could say that T(N) is

bounded above by a constant multiple of f(N).

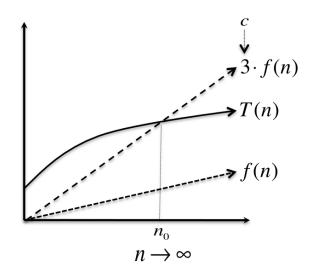


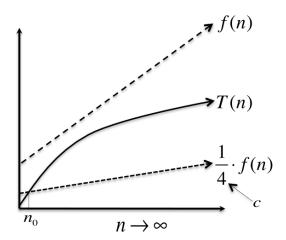
Figure 2.1: A picture illustrating when T(n) = O(f(n)). The constant c quantifies the "constant multiple" of f(n), and the constant n_0 quantifies "eventually."

Figure 1: Big-Oh

For example, $T(N)=O(N^2)$ means that the function T(N) grows at a rate no faster than N^2 ; its growth could equal that of N^2 but it could never surpass it.

3.2.2 Big-Omega notation

When we say that $T(N) = \Omega(g(N))$, we're guaranteeing that the function T(N) grows at a rate no lower than g(N); thus g(N) is a **lower bound** on T(N). Alternatively, we could say that T(N) is **bounded below** by constant mutiple of g(N).



T(n) again corresponds to the function with the solid line. The function f(n) is the upper dashed line. This function does not bound T(n) from below, but if we multiply it by the constant $c = \frac{1}{4}$, the result (the lower dashed line) does bound T(n) from below for all n past the crossover point at n_0 . Thus $T(n) = \Omega(f(n))$.

Figure 2: Big-Omega

For example, $T(N)=\Omega(N^2)$ means that the function T(N) grows at a rate no lower than N^2 ; its growth could equal that of N^2 but it could never drop below it.

Whenever talking about Big-Oh, there's always an implication about Big-Omega. For instance, T(N)=O(f(N)) (i.e., f(N) is an **upper bound** on T(N)) implies that $f(N)=\Omega(T(N))$ (i.e., T(N) is a **lower bound** on f(N)). As an example, N^3 grows faster than N^2 , so we can say that $N^2=O(N^3)$ or $N^3=\Omega(N^2)$.

3.2.3 Big-Theta notation

When we say that $T(N) = \Theta(h(N))$, we're guaranteeing that the function T(N) grows at the same rate as g(N). However, when two functions grow at the same rate, then the decision of whether or not to signify this with $\Theta()$ can depend on the context.

3.3 Typical growth rates

Function	Name
c	Constant
log_2N	Logarithmic
log^2N	Log-squared
N	Linear
NlogN	
N^2	Quadratic
N^3	Cubic
2^N	Exponential

3.4 Rules

Rule 1 If $T_1(N) = O(f(N))$ and $T_2(N) = O(g(N))$, then

1.
$$T_1(N)+T_2(N)=O(f(N)+g(N)).$$
 Less formally it is $O(\max(f(N),g(N))),$ and

2. $T_1(N)\cdot T_2(N) = O(f(N)\cdot g(N))$

Rule 2 If T(N) is a polynomial of degree k, then $T(N) = \Theta(N^k)$.

Rule 3 $log^k N = O(N)$ for any constant k. In other words, logarithms run at a lower rate than linear functions which means logarithms grow very slowly.

3.5 Few points

• In simple terms, the goal of asymptotic notation is to suppress constants factors and lower-order. Thus, they aren't included in Big-Oh. For instance, don't say $T(N) = O(2N^2)$ or $T(N) = O(N^2 + N)$. In both cases, the correct form is $T(N) = O(N^2)$.

When analyzing the running time of an algorithm, why would we want to throw away information like constant factors and lower-order terms? 1) Lower-order terms become increasingly irrelevant as you focus in large inputs, which are the inputs that require algorithmic ingenuity and 2) constant factors are generally highly dependent on the details of the environment (e.g., programming language, architecture, compiler, etc.) and thus ditching them allows to generalize by not committing ourselves to a specific programming language, architecture, etc.

• The relative growth rates of two functions f(N) and g(N) can always be determined by computing $\lim_{N\to\infty}\frac{f(N)}{g(N)}$, using L'Hopital's rule if necessary. The limit can have four possible values:

- The limit is 0, meaning that f(N) = O(g(N)).
- The limit is $c \neq 0$, meaning that $f(N) = \Theta(g(N))$.
- The limit if ∞ , meaning that g(N) = o(f(N)).
- The limit doesn't exist and thus no relation exist.
- Stylistically it's bad to say that $f(N) \leq O(g(N))$ since the inequality is already implied by the definition of Big-Oh.
- It's wrong to write $f(N) \ge O(g(N))$, because it doesn't make sense.

3.6 Model of computation

The model of computation is basically a normal computer with the following characteristics:

- Instructions are executed sequentially.
- The model has the standard repertoires of simple instructions, such as addition, multiplication, comparison, and assignment. Unlike real computers, this computer takes exactly one time unit to do anything.
- The computer has fixed-size (e.g., 32-bit) integers and no fancy operations (e.g., matrix invertion, sorting, etc.).
- The computer has infinite memory.

3.7 What to analyze

The most important resource to analyze is generally the *running time* and typically, the size of the input is the main consideration. We define two functions, $T_{avg}(N)$ (for the average-case running time) and $T_{worst}(N)$ (for the worst-case running time) used by an algorithm on input of size N. It's worth noting that $T_{avg}(N) \leq T_{worst}(N)$ (i.e., $T_{avg}(N)$ has a lower rate growth than that of $T_{worst}(N)$).

3.7.1 Types of performance

Best-case performance Although it can be ocassionally analyzed, it's often of little interest since it doesn't represent typical behavior.

Average-case performance It often reflects typical behavior, however it doesn't provide a bound for all input and it can be difficult to compute. Furthermore, it's also hard to define what's the average input

Worst-case performance It represents a guarantee for performance on any possible input. This is the quantity generally required because it provides a bound for all input.

3.8 Running-time calculations

When computing a Big-Oh running time, there are several general rules:

For loops The running time of a **for** loop is *at most* the running time of the statements inside the **for** loop times the number of iterations.

Nested loops Analyze these inside out. The total running tome of a statement inside a group of nested loops is the running time of the statement multiplied by the product of the sizes of all the loops.

For example, the following program is $O(N^2)$ because the inner loop does N iterations and the outer loops does N too. This amounts to $N\times N=N^2$ iterations. The first assignment counts as one operation (thus O(N)) while the innermost statement counts for 2 operations (1 multiplication and 1 assignment). Thus, to be more precise the program's running time is $1+2N^2$, however Big-Oh suppresses constant factors and lower-order terms.

```
k ← 0
for [1, n] → i:
    for [1, n] → j:
        k ← i * j
```

Consecutive statements They just add (which means that the maximum is the one that counts).

For example, the following program fragment, which has O(N) work followed by $O(N^2)$ work, is ultimately $O(N^2)$ which dominates O(N):

```
for [0, n) → i:
    a[i] = 0

for [0, n) → i:
    for [0, n) → j:
        a[i] += a[j] + i + j
```

If/else For the fragment **if** condition { S1 } **else** { S2 }, the running time of an **if/else** statement is never more than the running time of the test plus the larger of the running times of S1 and S2.

3.9 Sample problems

3.9.1 Sum of cubes

```
Input: a positive integer n.
Output: the sum of all cubes from 1 to n^3.

SumOfCubes(n):
    sum \( \chi \)
    for [1, n] \( \rightarrow \) i:  # O(n)
        sum += i \( \rightarrow \) i  # O(4), 1 assignment, 1 addition and 2 products
    return sum  # O(1)
```

Thus, for SumOfCubes(N) = 1 + 4n + 1 = 4n + 2 = O(n). We discard both the constant factors and the lower-order terms.

3.9.2 Factorial

The factorial is defined as

$$n! = n \times (n-1) \times (n-2) \times \cdots \times 2 \times 1$$

Implemented recursively, the algorithm is as follows:

```
Input: a non-negative integer n
Output: the factorial of n

Factorial(n):
    if n ≤ 1:
        return 1
    else:
        return n * Factorial(n-1)
```

However, this is a thinly veiled **for** loop. In this case, the analysis involves the recurrence relation T(n) = 1 + T(n-1) for n > 1, T(1) = 2 that needs to be solved:

$$T(N) = 1 + T(n-1)$$

$$= 1 + (1 + T(-2))$$

$$= 1 + (1 + (1 + T(n-3)))$$

$$= \dots$$

$$= 1 + (1 + (1 + T(n-k))) k 1's$$

$$T(N) = k + T(n-k)$$

$$= (n-1) + T(n-(n-1))$$

$$= (n-1) + T(1) = (n-1) + 2 = n + 1$$

Thus, T(n) = O(n).

3.9.3 Maximum subsequence sum problem

Given (possibly negative) integers A_1,A_2,\ldots,A_N , find the maximum value of $\sum_{k=1}^i A_k$.

In other words, given a one-dimensional array of numbers, find a contiguous subarray with the largest sum. For example, with the input -2, 11, -4, 13, -5, -2 the answer is 11 - 4 + 13 = 20.

This problem is interesting mainly because there are many algorithms to solve it, and the performance of these algorithms varies drastically.

	Algorithm Time			
Input Size	$\frac{1}{O(N^3)}$	2 O(N ²)	3 O(N log N)	4 O(N)
N = 100	0.000159	0.000006	0.000005	0.000002
N = 1,000	0.095857	0.000371	0.000060	0.000022
N = 10,000	86.67	0.033322	0.000619	0.000222
N = 100,000	NA	3.33	0.006700	0.002205
N = 1,000,000	NA	NA	0.074870	0.022711
Linear ————————————————————————————————————	80 90 100	Running Time	1000 2000 3	Linear O(N log N) Quadratic Cubic

Figure 3: Plot (N vs. time) of various algorithms

For algorithm 4 (linear), as the problem size increases by a factor of 10, so does the running time. For algorithm 3 (quadratic), a tenfold increase in input size yields roughly hundrefold (10^2) increase in running time.

3.9.4 Algorithm 1 (brute force)

implementations/max-subsequence-sum:

For this algorithm, there's N starting places, average $\frac{N}{2}$ lengths to check, and average $\frac{N}{4}$ numbers to add. Thus we have $O(N^3)$.

3.9.5 Algorithm 2 (brute force)

implementations/max-subsequence-sum:

For this algorithm, the innermost **for** loop has been removed. There's N starting places and an average $\frac{N}{4}$ numbers to add. Thus we have $O(N^2)$.

3.9.6 Algorithm 3

implementations/max-subsequence-sum:

```
Input: array A of integers.
Output: find the maximum sum in subarray spanning A[LEFT..RIGHT]. It does
not attempt to mantain actual best sequence.
MaxSumRec( A, LEFT, RIGHT ):
    # Base case
    if LEFT = RIGHT:
        if A[LEFT] > 0:
            return A[LEFT]
        else
            return 0
    center ← (LEFT + RIGHT) div 2
    maxLeftSum ← MaxSumRec(A, LEFT, center)
    maxRightSum ← MaxSumRec(A, center + 1, RIGHT)
    maxLeftBorderSum ← 0
    leftBorderSum ← 0
    for [center, LEFT] → i:
        leftBorderSum += A[i]
        if leftBorderSum > maxLeftBorderSum:
            maxLeftBorderSum ← leftBorderSum
    maxRightBorderSum ← 0
    rightBorderSum ← 0
    for [center + 1, RIGHT] \rightarrow j:
        rightBorderSum += A[j]
        if rightBorderSum > maxRightBorderSum:
            maxRightBorderSum ← RightBorderSum
    return max(maxLeftSum, maxRightSum, maxLeftBorderSum +
       maxRightBorderSum)
# Driver for divide-and-conquer maximum contiguous subsequence sum
   algorithm.
MaxSubsequenceSum( A ):
    return MaxSumRec(A, 0, A.size - 1)
```

3.9.7 Algorithm 4

implementations/max-subsequence-sum:

```
Input: array A of integers.
Output: the maximum positive subsequence sum.

MaxSubsequenceSum( A ):
    maxSum \in 0
    currentSum \in 0

for [0, A.size] \rightarrow j:
    currentSum += A[j]

    if currentSum > maxSum:
        maxSum \in currentSum
    else if currentSum < 0:
        currentSum = 0</pre>
```

4 Lecture 4: Algorithm analysis (cont.) and lists/stacks/queues

4.1 Binary search

Given an integer X and integers A_0, A_1, \dots, A_{N-1} , which are **presorted** already in memory, find index i such that $A_i = X$, or return -1 if X is not in the group of integers.

implementations/binary-search

```
Input: array A of elements and item x which we are searching for.
Output: item where item is found or -1 if not found.

BinarySearch( A, x ):
    low \in 0
    high \in A.size - 1
    while low \le high:
        mid \in (low + high) / 2
        if A[mid] \le x:
            low = mid + 1
        else if A[mid] > x:
            high = mid - 1
        else:
            return mid
    return -1
```

4.1.1 Running time

We have the following recurrence relation T(N)=1+T(N/2) and T(1)=1. We must solve it to find out the algorithm's running time:

$$T(N) = 1 + T(N/2)$$

$$= 1 + 1 + T(N/2^{2})$$

$$= 1 + 1 + 1 + T(N/2^{3})$$
...
$$= k + T(N/2^{k})$$

If N is a power of 2 (i.e., $N=2^k$ with k=log(N)) we'll have:

$$T(N) = k + T(N/2^k)$$
$$= k + T(1) = log(N) + 1$$

This means that T(N) = O(log N).

4.2 List/stacks/queue ADTs

An **abstract data type** (ADT) is a set of objects (lists, sets, graphs, etc.) together with a set of related operations (add, remove, size, etc.). They provide a template for the objects, not how the objects and their respective set of operations are implemented.

4.2.1 The List ADT

We usually deal with a general list of the form $A_0, A_1, A_2, \dots, A_{N-1}$. We say that the size of this list is N. We will call the special list of size 0 an **empty list**. The **position** of the element A_i in a list is i.

The List ADT could describe the followin operations:

find(x) Return the position of the occurrence of item x.

insert(x, i) Insert some element x at position i.

remove(i) Remove some element at position *i*.

4.2.2 Vector in the STL

Pros:

- · Constant tome indexing
- Fast to add data at the end (not the front).

Cons:

- · Slow to add data in the middle.
- · Inefficient for searches.

Operations

- void push_back(const T &x) add x at the end of the list.
- void pop_back() remove element at the end of the list.
- const T &back() const return the element at the end of the list.
- const T &front() const return the element at the front of the list.
- void push_front(const T &x) add x to the front of the list.
- **void** pop_front() remove the element at the front of the list.

4.2.3 List in the STL

Pros:

- Implemented as a doubly linked list.
- Fast insertion/removal of items in any position.

Cons:

- · No indexing.
- Inefficient for searches.

Operations

- T & operator[](int idx) return element at index idx with no bounds checking.
- T &at(int idx) return element at index idx with bounds checking.
- int capacity()conts return internal capacity of vector'.
- void reserve(int new_capacity) set new capacity and possibly void expansion of vector.

4.3 Containers

A **container** is a holder object that stores a collection of other objects. This is usually implemented as a class templates which provides it with great flexibility for the types supported as elements.

A container

- manages the storage space for its elements and provides member functions to access them either directly or through **iterators**.
- replicates structures commonly used in programming such as dynamic arrays (e.g., vectors), queues, stacks, heaps (e.g., priority queues), linked lists, trees (e.g., sets), associative arrays (e.g., maps), etc.

4.4 Iterators

Some operations on lists require the notion of a position. In the STL, a position is represented by some nested type known as an **iterator**. In particular, for a list<string>, the position is represented by the type list<string>::iterator; for a vector<**int**>, Vector<**int**>::iterator.

4.4.1 How to get an iterator

The STL lists (and all other STL containers) define a pair of methods:

- iterator begin(): returns an appropriate iterator representing the first item in the container.
- iterator end(): returns an appropriate iterator representing the endmarker in the container. This endmarker is "out-of-bounds".

4.4.2 Iterator methods

- itr++ and ++itr: advances the iterator itr to the next location.
- *itr: returns a reference, which may or may not be modifiable, to the object stored at iterator itr's location.
- itr1 == itr2: returns true if iterators itr1 and itr2 refer to the same location and false otherwise.
- itrl != itrl: returns true if iterators itrl and itrl refer to a different location and false otherwise.

For example the code:

```
for (int i = 0; i <!= v.size(); i++) {
   std::cout << v[i] << "\n";
}</pre>
```

could be rewritten as follows using iterators:

```
for (vector<int>::iterator itr = v.begin(); itr != v.end(); itr++) {
    std::cout << *itr << "\n";
}</pre>
```

Alternatively:

```
vector<int>::iterator itr = v.begin();
while (itr != v.end()) {
    std::cout << *itr++ << "\n";
}</pre>
```

4.4.3 Container operations that require iterators

- iterator insert(iterator pos, **const** T & x): adds x into the list, prior to the position by the iterator pos. This is a constant operator for list, but not for vector. The return value is an iterator representing the position of the inserted item.
- iterator erase(iterator pos): removes the object at position and return the position of the element that followed pos prior to the call. It invalidates pos, making it stale.
- iterator erase(iterator start, iterator pos): removes all items beginning at position up to, but not including end. An entire list c can be erased by c.erase(c.begin(), c.end()).

4.4.4 Example: Using erase on a list

4.4.5 const_iterators

The result of *itr is both the value of the item that the iterator is viewing but also the item itself. This is very powerful but also introduces some complications.

Let's analyze the following routine that works for both vector and list and runs in linear time:

The potential problem arises if Container c was passed to a routine using call-by-constant reference, meaning we would expect that not changes would be allowed to c, and the compiler would ensure this by nota allowing calls to any c's mutators. For example, consider the following code that prints a list of integers but also tries to do some changes:

```
void print( const list<int> &lst, ostream &out = cout ) {
   typename Container::iterator itr = lst.begin();
   while (itr != lst.end()) {
      out << *itr << "\n";
      *itr = 0; // <= This is the suspect.
      itr++;
   }
}</pre>
```

The solution provided by the STL is that every collection contains not only an iterator nested type but also const_iterator nested type. The main difference between them is that operator* for const_iterator returns a constant reference, and thus itr* for a const_iterator cannot appear on the left-hand side of an assignment statement.

Further the compiler will force you to use a const_iterator to traverse a *constant* collection and does so by providing two versions versions of begin and two versions of end:

```
iterator begin()
const_iterator begin()const
iterator end)
const_iterator end()const
```

Note: The two versions of begin can be in the same class only because of the const-ness of a method (whether an accessor or mutator) is considered to be part of the signature. The trick is overloading operator[].

Using auto to declare your iterators means the compiler will deduce for you whether an iterator or const_iterator is substituted. This exempt the programmer from having to keep track of the correct iterator type and is precisely one of the intended uses of auto.

4.4.6 Printing a container

```
* @brief Prints out the container on the output stream.
* @param c a given container.
* @param out an output stream.
 */
template <typename Container>
void print( const Container &c, ostream &out = cout ) {
    if (c.empty()) {
        out << "(empty)";
    }
    else {
        typename Container::iterator itr = begin(c);
        out << "[" << *itr++; // print first item</pre>
        while (itr != end(c)) {
            out << ", " << *itr++;
        out << "]" << "\n";
   }
}
```

5 Lecture 5: Lists, Stacks, Queues, and the STL

5.1 Implementation of a vector

The main details of the implementation of the Vector class is as follows:

• It'll maintain the primitive array (via a pointer variable to the block of allocated memory), the array capacity, the current number of stored items.

- It'll implement the "big-five" to provide **deep-copy** semantincs.
- It'll provide the resize and reserve methods that will change the size of the Vector and the capacity of the Vector respectively.
- It'll overload the operator[].
- It'll provide several basic methods, such as size, empty, clear, back, pop_back, and push_back.
- It'll provide support for the nested types iterator and const_iterator, and associated begin and end methods.

Implementation: implementations/Vector

Why create a class Vector? Is it better than a simple array? There are few reasons why creating Vector class template makes sense. The main one is that Vector will be a first-clas type, which means that 1) it can be copied, and 2) the memory it uses can be reclaimed back (via its destructor). This is not the case for primitie array in C++.

5.2 Implementation of a list

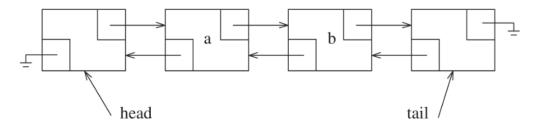
The main details of the implementation of the List class template is as follows:

- The underlying data structure is a **doubly linked list** and thus we must mantain pointers to both ends of the list. This is in order to mantain constant time cost per operation, so as long as the operation occurs at a known position. The known position can be either end or at a position specified by an iterator.
- Four classes will be provided to better compose the List class template:
 - The List class itself, which contains links to both ends, the size of the list, and several methods.
 - The Node class, which is likely to be a private nested class. A node contains the data and pointers to the previous and next nodes, along with appropriate constructors.

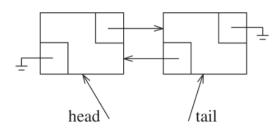
 The const_iterator class, which abstracts the notion of a position, and is a public nested class. It stores a pointer to the curren node, a provodes implementation of the basic iterator operations.

- The iterator class, which abstracts the notion of a position, and is a public nested class. It
 has similar functionality to const_iterator, except that operator* returns a reference
 to the item being viewed, rather than a constant reference.
- Extra nodes are at both the front and end of the list. The node at the front represents
 the beginning marker while the one at the end represents the endmarker. These are the
 sentinel nodes; the one at the front is the header node and the one at the end is the tail
 node. Using the sentinels simplify the coding by removing a host of special cases (e.g.,
 removing the first node).

Implementation: implementations/List



A doubly linked list with header and tail nodes

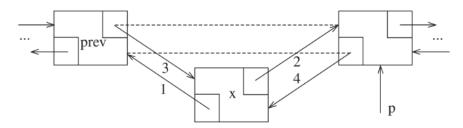


An empty doubly linked list with header and tail nodes

Figure 4: Doubly linked list

5.2.1 Insertion

```
Node *newNode = new Node{ x, p->prev, p }; // Steps 1 and 2
p->prev->next = newNode; // Step 3
p->prev = newNode; // Step 4
```

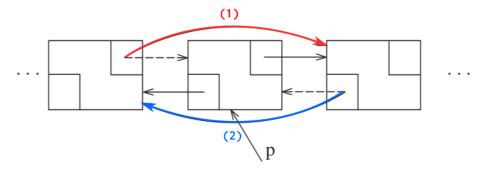


Insertion in a doubly linked list by getting a new node and then changing pointers in the order indicated

Figure 5: Adding a node

5.2.2 Erasure

```
p->prev->next = p->next; (1)
p->next->prev = p->prev; (2)
delete p;
```



Removing node specified by p from a doubly linked list

Figure 6: Removing a node

5.3 The Stack ADT

A **stack** (sometimes known as a LIFO (*last in, first out*) list) is a special list where insertions and deletions can be performed only from the end of the list (a.k.a., the **top**).

The fundamental operations on a stack are:

push Push an item to the top of the stack. Equivalent to insert.

pop Pop the most recently inserted item. The topmost item can be examined by using the top routine.

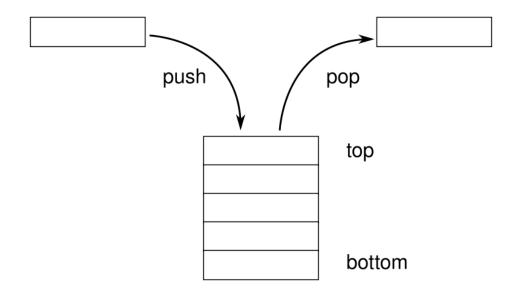


Figure 7: Stack model

5.3.1 Implementations

Two population implementations of stacks are:

Linked list implementation

This implementation of a stack uses a singly linked list. A push operation is performed by inserting at the front of the list, a pop operation is performed by deleting the element at the front of the list, and a top operation merely examines the element at the front of the list by returning its value.

Implementation: implementations/LLStack

· Array implementation

This implementation of a stack avoids pointers and is usually the most popular solution. It uses the back, push_back, and pop_back implementation from vector. Associated with each stack are array to stores the items and top_of_stack which is -1 for an empty stack.

Implementation: implementations/ArrayStack

5.3.2 Applications

After the array, the stack is probably the most fundamental data structure in computer science and for this reason, the application of stacks can be found in many places. Some of them are:

Balancing symbols

Given that compiler check programs for syntax error, a useful tool to check for the lack of symbols that might cause errors is a program that checks whether everything is balanced (i.e., every right brace, bracket, and parenthesis must correspond to its left counterpart). For instance, the sequence [()] is legal, but [(]) is wrong. This algorithm uses a stack and could be described as follows:

Make an empty stack. Read characters until end of file. If the character is an opening symbol, push it onto the stack. If it is a closing symbol and the stack is empty, report an error. Otherwise, pop the stack. If the symbol popped is not the corresponding opening symbol, then report an error. At end of file, if the stack is not empty, report an error.

Postfix expressions

This notation, also known as **reverse Polish notation**, is a mathematical notation in which operators follow their operands, in contrast to Polish notation, in which operators precede their operands. For example, the infix expression $((15 \div (7-(1+1))) \times 3) - (2+(1+1))$ can be written in reverse Polish notation as $15\,71\,1 + - \div 3 \times 21\,1 + +-$ which yields 5 regardless of the direction of the evaluation. An algorithm that uses a stack and that processes the expression from left to right is described below:

```
Input: EXPR is an expression in reverse Polish notation.
Output: EXPR

EvaluatePostfixExpression( EXPR ):
    stack \( \) []
    for EXPR \( \) token:
        if token = operator:
            operand_2 \( \) stack.pop()
            operand_1 \( \) stack.pop()
            result \( \) evaluate operator_1 and operand_2 with operator stack.push(result)
    else if token = operand:
            stack.push(token)
    return stack.pop()
```

5.4 The Queue ADT

A **queue** (sometimes known as a FIFO (*first in, first out*) list) is a speciallist where insertions are performed at one end and deletions are performed at the other end.

The fundamental operations on a queue are:

enqueue Insert an element at the end of the list (called the rear).

dequeue Delete (and return) the element at the start of the list (called the front).

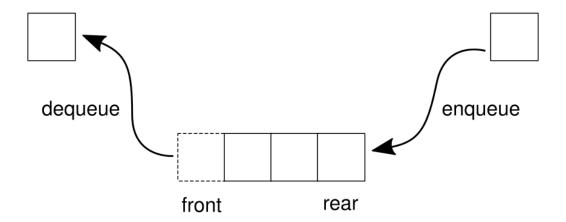


Figure 8: Queue model

5.4.1 Implementations

Two population implementations of stacks are:

• Linked list implementation.

This is implementation is similar to its counterpart stack implementation.

Implementation: implementations/LLQueue

Array implementation

For this implementation, we keep an array, array, and the positions front and back, which represents the ends of the queue. Likewise, we keep track of the number of items currently in the queue, current_size. To enqueue an element x, we increment current_size and back, then

set _array[back] = x. To dequeue an element, we set the return value to array[front], decrement current_size, and increment front.

However there's one potential problem with this implementation. After a certain number of operations, a queue might appear full since back is now at the last array index, and the next enqueue would be in a nonexistent position. A simple solution would be to wrap around to the beginning whenever front or back gets to the end of the array. This is known as a **circular array** implementation for which modular arithmetic is used.

Implementation: implementations/ArrayQueue

5.4.2 Applications

There are many algorithms that used queues to give efficient running times. Some examples are:

- Jobs submitted to a printer are arranged in order of arrival. Thus, they're essentially placed in a queue.
- Virtually every real-life line is supposed to be a queue. For instance, lines at ticket counters are queues, because the service is first-come first-served.
- There are many network setups of personal computers in which the disk is attached to one machine, known as the file server. Users on other machines are given access to files on a first-come first-served basis, so the data structure is a queue.

5.5 Trees

Recursively, a **tree** is either empty or consists a root node r and zero or more non-empty subtrees whose roots are connected by an edge.

- Used to implement file systems of several popular OSs
- Used to evaluate arithmetic expressions
- Support O(logN) search
- Set and map classes.

6 Lecture 6: Trees

6.1 Preliminaries

A **tree** is a widely used abstract data type (ADT) that simulates a hierarchical tree structure, with a root value and subtrees of children with a parent node, represented as a set of linked nodes.

Trees (more specifically binary trees) are really important so a certain terminoly has developed for them:

- A **node** is a structure which may contain a value or condition, or represent a separate data structure.
- An edge is connection between one node and another.
- A path is a sequence of nodes and edges connecting a node with a descendant.
- An **ancestor** is a node reachable by repeated proceeding from child to parent. A **descendant** is a node reachable by repeated proceeding from parent to child. For example, if there's a path from node u to node v, then u is an ancestor of v and v is a descendant of u. If $u \neq v$, then u is a **proper ancestor** of v and v is a **proper descendant** of u.
- The root r of a tree is its top node known as the prime ancestor.
- The root of each subtree is said to be a **child** of the root r, and r is the **parent** of each subtree root.
- Two nodes \boldsymbol{u} and \boldsymbol{v} are **siblings** if they have the same parent.
- The **subtree** of a node, u, is the tree rooted at u and contains all of u's descendants.
- The **depth** of a node, u, is the length of the path from u to the root of the tree. This translates into the number of edges from node to the tree's root node. Thus, a root node will have a depth 0.
- The **height** of a node, u, is the length of the longest path from u to one of its descendants. Thus, the height of a tree is equal to the height of the root. Since leaves has no children, all leaves have height 0.
- A node, u, is a **leaf** if it has no children.

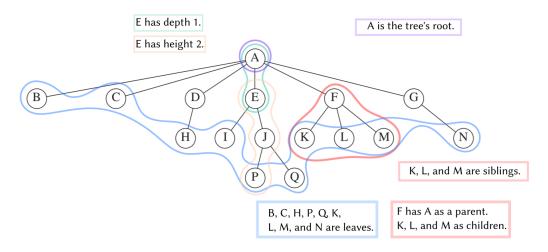


Figure 9: A tree

An m-ary tree is a rooted tree in which each node has no more than m children. When m=2, the tree

is known as a **binary tree** (i.e., each node has at most two children, the left child and the right child). When m=3, the tree is a **ternary tree**.

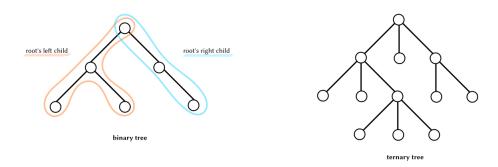
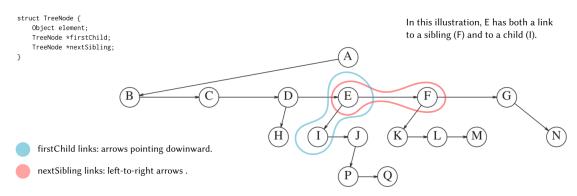


Figure 10: Binary and ternary tree

6.2 Implementation of trees

An obvious way to implement a tree would be to have in each node, besides its data, a link to each child of the node. However, the number of children per node can vary and is not known in advance which might translate into too much wasted space. The solution is simple: Keep the children of each in a linked list of nodes. In this way, each node can have as many children as the tree it composes allows.

```
template < typename Object >
struct TreeNode {
    Object element;
    TreeNode* firstChild;
    TreeNode* nextSibling;
}
```



First child/next sibling representation of the tree

Figure 11: First child/next-sibling representation

6.3 Tree traversals

A **tree traversal** refers to the process of visiting each node in a tree data structure, exactly once. Such traversals are classified by the order in which the nodes are visited.

Trees can be traversed either:

• by deepening the search as much as possible on each child before going to the next sibling. This is known as **depth-first search** (DFS).

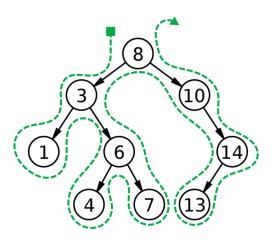


Figure 12: Depth-first search

• by visiting every node on a level before going to a lower level. This is known as **breadth-first**

search (BFS).

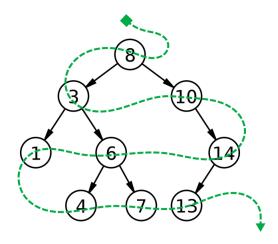


Figure 13: Breadth-first search

6.3.1 Depth-first search

Depending on when the work at a node is performed, a traversal of this type can be classified into:

- **pre-order traversal**. In a pre-order traversal, work at a node is performed before (*pre*) its children are processed.
- 1. Check if the current node is empty or null.
- 2. Display the data part of the root (or current node).
- 3. Traverse the left subtree by recursively calling the pre-order function.
- 4. Traverse the right subtree by recursively calling the pre-order function.

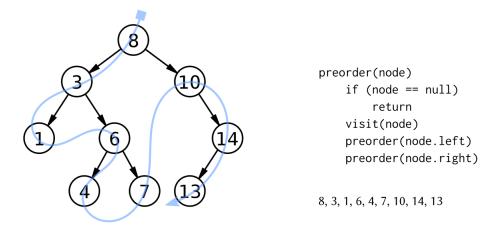


Figure 14: Pre-order traversal

- in-order traversal. In an in-order traversal, work at a node is performed after its left child is visited, followed by the right child.
 - 1. Check if the current node is empty or null.
- 2. Traverse the left subtree by recursively calling the in-order function.
- 3. Display the data part of the root (or current node).
- 4. Traverse the right subtree by recursively calling the in-order function.

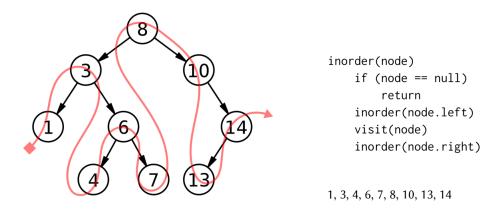
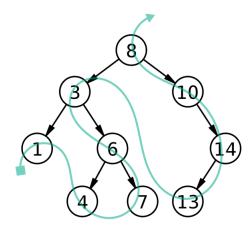


Figure 15: In-order traversal

• **post-order traversal**. In a post-order traversal, work at a node is performed after (*post*) its children are evaluated.

- 1. Check if the current node is empty or null.
- 2. Traverse the left subtree by recursively calling the post-order function.
- 3. Traverse the right subtree by recursively calling the post-order function.

4. Display the data part of the root (or current node).



```
postorder(node)
   if (node == null)
      return
   postorder(node.left)
   postorder(node.right)
   visit(node)
```

1, 4, 7, 6, 3, 13, 14, 10, 8

Figure 16: Post-order traversal

6.4 Binary trees

As stated earlier, a **binary tree** is a tree in which no node can have more than two children.

A property of a binary tree that is sometimes important is that the depth of an average binary tree is considerably smaller than N. An analysis shows that the average depth is $O(\sqrt{N})$, and that for a special type of binary tree, namely the **binary search tree**, the average value of the depth is $O(\log N)$. Unfortunately, the depth can be as large as N-1. In its worst case, we end up with a linked list (which is also a type of tree).

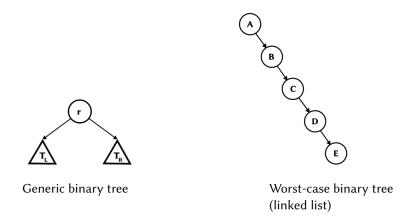


Figure 17: Generic binary tree and linked list

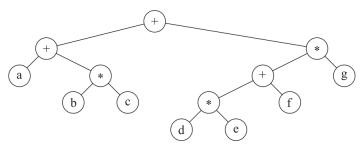
6.4.1 Implementation

Given that a binary tree has at most two children per node, we can keep direct links to them. Thus, a binary node is just a structure consisting of the data plus two pointers to current node's left and right nodes. For example:

Binary trees have many important uses not associated with searching. One of the principal uses of binary trees is in the area of compiler design (e.g., expression trees).

6.4.2 Expression trees

A **binary expression tree** is a specific kind of a binary tree used to represent expressions. The leaves of an expression are **operands**, such as constants or variable names, and the other nodes contain **operators**. We can evaluated an expression, T, by applying the operator at the root to the values obtained by recursively evaluating the left and right subtrees.



Expression tree for (a + b * c) + ((d * e + f) * g)

Figure 18: Expression tree

- To produce an (overly parenthesized) *infix expression*, recursively produce parenthesized left expression, then printing out the operator at the root, and finally recursively producing a parenthesized right expression. This is an **in-order traversal**.
- To produce a *postfix expression*, recursively print out the left subtree, the right subtree, and then the operator. This is an **post-order traversal**.
- To produce a *prefix expression*, print out the operator first and then recursively print out the left and right subtrees. This is an **pre-order traversal**.

6.4.3 Constructing an expression tree

This algorithm describes the steps to convert a postfix expression into an expression tree:

Implementation: implementations/ExpressionTree

6.5 Binary search trees (BTSs)

A binary search tree is a special type of binary trees with the following property:

For every node, X, in the tree, the values of all the items in its left subtree are smaller than the item in X, and the value of all the items in its right are larger than the item in X.

This implies that all the elements in the tree can be ordered in some consistent manner.

Binary search trees keep their keys in sorted order, so that lookup and other operations can use the principle of binary search: when looking for a key in a tree (or a place to insert a new key), they traverse the tree from root to leaf, making comparisons to keys stored in the nodes of the tree and deciding, on the basis of the comparison, to continue searching in the left or right subtrees. On average, this means that *each comparison allows the operations to skip about half of the tree*, so that each lookup, insertion or deletion takes time proportional to the logarithm of the number of items stored in the tree.

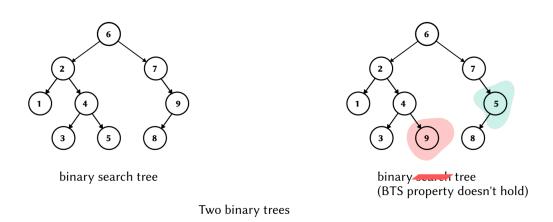


Figure 19: Binary search tree vs Non-binary search tree

Complete implementation: implementations/BinarySearchTree

6.5.1 Operations on a BTS

contains For this operation, we're required to return **true** if there is a node in the tree T that has item X. Otherwise, return **false**. If T is empty, then we can return **false** right away. Otherwise, if the item stored at T is X, we can return **true**. Otherwise, we can make a recursive call on a subtree of X, either left or right, depending on the relationship between X and the item stored in T. It's implied that the objects being compared have implemented the used relational operators appropriately.

```
Input: X is an item to search for and T is the node that roots
the subtree.
Output: Return true or false depending on whether the tree
contains the node storing the item.

contains( X, T ):
    if T = null:
        return false
    else if X < T.element():
        return contains X, T.left()
    else if X > T.element():
        return contains X, T.right()
    else:
        return true # Match
```

findMin and findMax These operations can either return the node (or its value) containing the smallest and largest element in the tree, respectively. To perform a findMin, start at the root and go left as long as there's a left child. The stopping point is the smallest element. The findMax routine is similar, except that branching is to the right child.

Recursive algorithm of findMin:

```
Input: the root of the tree.
Output: the smallest element in the tree.

findMin( T ):
    if T = null:
        return T
    if T.left() = null:
        return T;
    return findMin T.left()
```

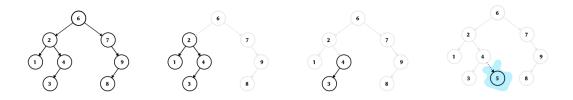
Iterative algorithm of findMax:

insert To insert X into tree T, proceed down the tree as you would with the contains operation. If X is found, do nothing. Otherwise, insert X at the last spot on the path traversed.

Duplicates can be handled by keeping an extra field in the node object indicating the frequency of ocurrence for a particular element X. This adds some space extra space to the entire tree but is better than putting duplicates in the tree (which tends to make the tree very deep).

```
Input: X is an item to inserted in the tree T.
Output: void.

insert( X, T ):
    if T ← null:
        T ← BinaryNode(X, null, null)
    else if X < T.element()
        insert X, T.left()
    else if X > T.element()
        insert X, T.right()
    else:
        # duplicate; do nothing
```

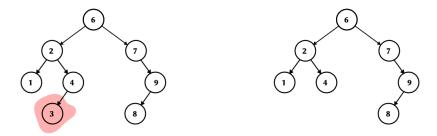


Progression of inserting 5 into the tree

Figure 20: Progression of an insertion

remove As is common with many data structures, the hardes operation is deletion. After finding the item to be deleted several possibilities must be considered:

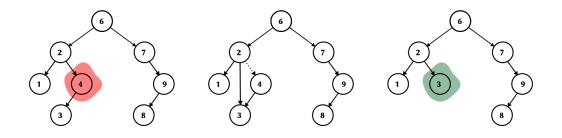
• If the node is a leaf, it can be deleted immediately.



Removing a childless node (3) from the tree

Figure 21: Removing childless node from tree

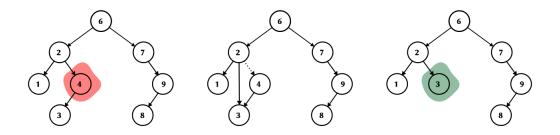
• If the node has one child, the node can be deleted after ist parent adjusts a link to bypass the node.



Progression of removing node (4) with one child

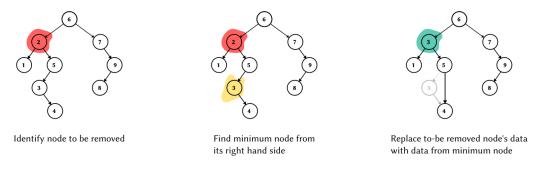
Figure 22: Removing single child node from tree

• If the node has two children, replace the data of this node with the smallest data of the right subtree and recursively delete that node (which is now empty). Because the smallest node in the right subtree cannot have a left child, the second remove is an easy one.



Progression of removing node (4) with one child

Figure 23: Removing single child node from tree



Progression of removing node (2) with two children

Figure 24: Removing node with two children from tree

```
Input: X is an item to be removed and T is the tree.
Output: void
remove( X, T ):
    if T = null:
       return; # Item not found; do nothing
    if X < T.element():</pre>
        remove X, T.left()
    else if X > T.element():
        remove X, T.right()
    else if T.left() ≠ null AND T.right() ≠ null:
       min ← findMin T.right()
        T.element ← min.element()
        remove T.element(), T.right()
    else:
        old-node ← T
        T ← T.left ≠ null ? T.left() : T.right()
```

The algorithm above is inefficient because it makes two passes down the tree to find and delete the smallest node in the right subtree when this is appropriate. It's easy to remove this inefficiency by writing a special removeMin method.

If the numder of deletions is expected to be small, then a popular strategy to use is **lazy evaluation**: When an element is to be deleted, it's left in the tree and merely *marked* as being deleted.

Props of lazy evaluation (in this context):

- Easy to handle if a deleted item is reinserted, do not need to allocate a new node.
- Do not need to handle finding a replacement node.

Cons of lazy evaluation (in this context):

• The depth of the tree will increase. However this increase is usually a small amount relative to the size of the tree.

6.5.2 Average-case analysis

The running time of all the operations (except make Empty and copying) is O(d), where d is the depth of the node containing the accessed item.

The sum of the depths of all nodes in a tree is known as the **internal path length**.

Let D(N) be the internal path length for some tree T of N nodes. An N-node tree consists of an i-node left subtree and an (N-i-1)-node right subtree, plus a root at depth zero for $0 \le i \le N$. D(i) is the internal path length of the left subtree with respect to its root. The same holds for the right subtree. We get the following recurrence D(N) = D(i) + D(N-i-1) + N - 1.

In a BTS all subtree sizes are equally likely, so

$$D(N) = \frac{2}{N} \left[\sum_{j=0}^{N-1} D(j) \right] + N - 1$$

By solving the above recurrence we show that $D(N)=O(Nlog_N)$. Thus the expected depth of any node is $O(log_N)$.

Although it's tempting to say that this result implies the average running time of all the operatuons is log_N , but this is not entirely true. The reason for this is that because of deletions, it's not clear that all BSTs are equally likely. In particular, the deletion algorithm favors making the left subtrees deeper than the right, because we're always replacing a deleted node with a node from the right subtree.

6.5.3 Balanced trees

If the input comes into a tree presorted, then a series of inserts will take *quadratic time* and give a very expensive implementation of a linked list, since the tree will consist only of nodes with no left children. One solution to the problem is to insist on an extra structural condition called **balance** where no node is allowed to get too deep.

There are quite a few general algorithms to implement **balanced trees**. Most are quite a bit more complicated than a standard binary search tree, and all take longer on average for updates. They do, however, provide protection against the embarrassingly simple cases. Below, we will sketch one of the oldest forms of balanced search trees, the AVL tree.

6.6 AVLs