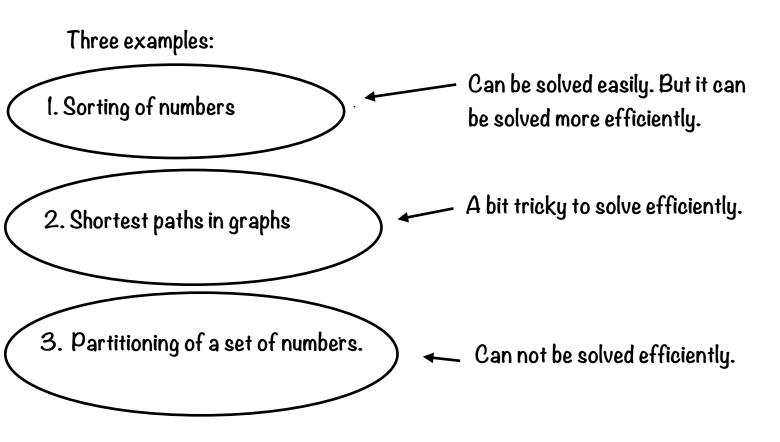
What is the course about?

How to solve problems in an algorithmic way.



There is a set of natural questions concerning algorithms:

How do we measure efficiency?

How do we find algorithms?

How do we describe algorithms?

How do we know if an algorithm works?

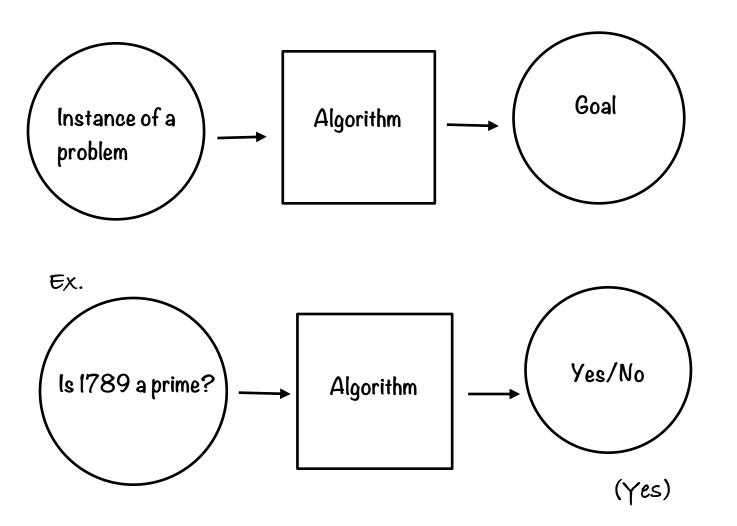
How do we know if a problem can be solved efficiently?

How do we know if a problem can be solved algorithmically at all?

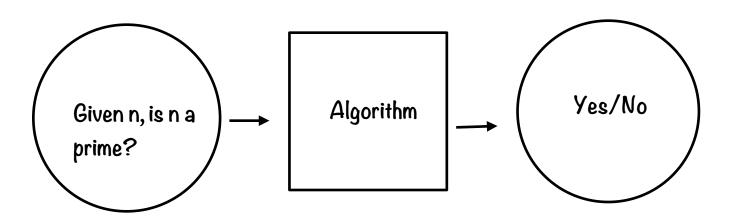
By the way, what is a problem?

We will try to provide some answers.

A general picture:



But an algorithm is usually not designed to solve just one instance. Normally it solves an infinite numbers of instances:



So what is a problem?

A problem consist of Input and Goal

Ex:

PRIME NUMBER

Input: An integer n

Goal: Is n a prime number?

SORTING

Input: A list $L = \{l_1, l_2, ...\}$ of numbers

Goal: A sorting $L' = \{ l'_1, l'_2, ... \}$ such that $l'_1, l'_2, ...$ is increasing

SHORTEST PATH

Input: A graph G. Two nodes a,b

Goal: A path from a to b with as few edges as possible

How do we desribe an algorithm?

An answer would be to just give a program code.

But that can be inconvenient.

Take for instance the most famous sorting algorithm: Insertion-Sort.

Informal description:

We start with allist A[1], A[2], ...

Take A[1] and set $L_1 = A[1]$.

Take A[2] and sort into L_1 . This gives us a new list L_2 .

Take A[3] and sort into L_2 . This gives us a new list L_3 .

And so on ...

We describe the algorithm with so called pseudocode.

```
INSERTION-SORT(A)

1 for j = 2 to A.length

2  key = A[j]

3  // Insert A[j] into the sorted sequence A[1..j-1].

4  i = j-1

5  while i > 0 and A[i] > key

6  A[i+1] = A[i]

7  i = i-1

8  A[i+1] = key
```

How do we know that Insertion-Sort works correctly?

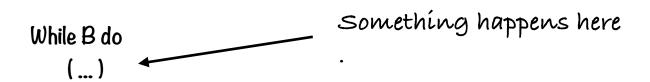
Think like this: At step k we know that

A[1], A[2], ..., A[k] is a part of the start list and A[1], A[2], ..., A[k] is sorted.

When the algorithm stops A[1], A[2], ..., A[k] must be the sorted list.

Usually, the problem is to decide if a loop works correctly. Sometimes we can define a loop-invariant I that is always true for each step in the lopp

Suppose the loop has the form



When the loop ends we have I and not B. If we have choosen I wisely, we will have reached our goal.

What about efficiency? How do we measure it?

Basic idea:

We measure the number of steps needed as a function of the size of the input. This type of measure is called time-complexity.

Insertion-Sort: If we have n elements we need $O(n^2)$ steps.

Problem: How do we define size of input?

Basic idea: We use a string to represent the input. We measure the size of the string.

In practice, we use something that is a convenient approximation of the size of the string.

Ex: If the input is n elements in al list we normally use n as a measure of the input size.

Ex: If the input is just an integer n we use log2n as the size of the input.

But there is a problem with finding good representations.

A bit more about complexity

Time-complexity is not the only measure of complexity.

We can measure the size of the memory space the algorithm needs to complete the computation. This measure is called space-complexity.

Time-complexity more important but space-complexity is also important.

Later we will show that high space-complexity implies high time-complexity

The opposite is not necessarily true.

Unit cost and bit cost

An interesting question is what counts as a step in a computation.

Unit cost: Each "algebraic operation" counts as one step

Bit cost: Each bit operation counts as a step

Ex:

We want to compute 521*394. What is the cost?

Unit cost: Just one operation! The cost is 1.

Bit cost: 521 has 9 bits. 394 has 8 bits. The cost is 9*8 = 72

(The answer is 205274.)

Unit cost is easier to work with and we normally use it.

What about efficiency? How do we measure it?

Basic idea:

We measure the number of steps needed as a function of the size of the input.

Insertion–Sort: If we have n elements we need $O(n^2)$ steps.

Ex: How do we represent a graph?

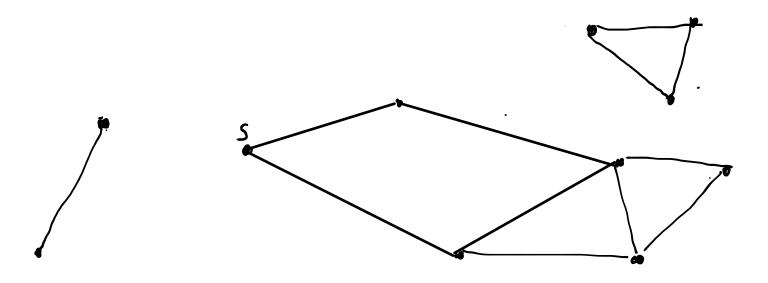
The natural measure of the input size could be |V| or |E| or |V| + |E|

Usually, the complexity is given as a function of I V I and I \in I.

We will study a special problem in graph theory

Input: A graph G and a node s.

Goal: Which nodes can be reached by paths from s?



A simple algorithm

R ←{s}

While there is (u,v) such that $u \in R$ and $v \not\in R$ Add v to R

End while

When the algorithm stops, \boldsymbol{R} is the set of nodes reachable from s.

But there is an uncertainty. How shall (u,v) be choosen?

Data structures for graphs

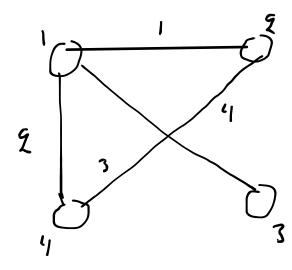
There are basically 3 ways to describe a graph in a form suitable for computation.

Adjacency matrix

Incidence matrix

Adjacency lists

Ex:



adjacency matrix A(i,j) = O/1

$$A = \begin{pmatrix} 0 & 1 & 1 & 1 \\ 1 & 0 & 0 & 1 \\ 1 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 \end{pmatrix}$$

 $\Delta(i,j)=1$ There is an edge (i,j)

Incidence matrix

$$(i,j) = 0/1$$

```
Adjacency lists
```

```
jis in Li <=> There is an edge (i,j)
```

L1: 2,3,4 L2:1, 4 L3:1 L4:1,2

> We now present an algorithm which is amore detailed solution to our graph problem.

When the algorithm stops,T is a tree containing all nodes that can be reached from s. The algorithm is known as Breadth-First-Search (BFS)

The complexity is O(V + E)

```
For all v \in V
    set vis(v) = 0
End for
Set vis(s) = 1
Set count = 0
Set T = \emptyset
Set L[O] = \{s\}
While L[i] is not empty
    Set L[i+1] = \emptyset
    For each u \in L[i]
         For each edge (u,v)
              If vis(v) = O
                  Set vis(v) = 1
                  Add (u,v) to T
                  Add v to L[i+1]
              End if
         End for
    End for
```

Set i = i+1

End while

We describe one more search algorithm for graphs: Depth-First Search (DFS)

```
Set = \emptyset
                                               DFS(u):
For all v \in V
     Set vis(v) = O
                                               Set vis(u) = 1
End for
                                               Add u to R
DFS(s)
                                               For each v such that v is adjacent to u
                                                    If vis(v) = O
                                                        DFS(v)
                                                    End if
                                               End for
                                             Set R = \emptyset
                                             Let S' be a stack and set S = \{s\}
The complexity is O(|V| + |E|).
                                             For all v \in V
The algorithm is defined
                                                 Set vis(v) = 0
recursively. A non-recursive
                                             End for
definition is:
                                             Set vis(s) = 1
                                             While S \neq \emptyset
                                                 Take the top node u from S
                                                 If vis(s) = 0
                                                      Add u to R
                                                      For each v adjacent to u
                                                           Add v to the top of S
                                                      End for
                                                 End if
                                             End while
```

Efficient algorithms

What is an efficient algorithm? We can start by asking what an inefficient algorithm is. If the size of the input is n and the algorithm works in time $O(2^n)$ we say that the algorithm has exponential complexity. Such an algorithm is clearly very inefficient.

In several situations where we have a problem with an input of size n, we have a set of possible solutions of size exponential in n. Just one (or a few) of these possible solutions are real solutions. If we just test all possible solutions to see if any of them are real solutions we get an algorithm withexponential complexity. In order to get a better algorithm we must find a way of "zooming in" on the real solutions.

It seems natural to say that an efficient algorithm is an algorithm which is not exponential. But we don't do this. Instead we use the following definition:

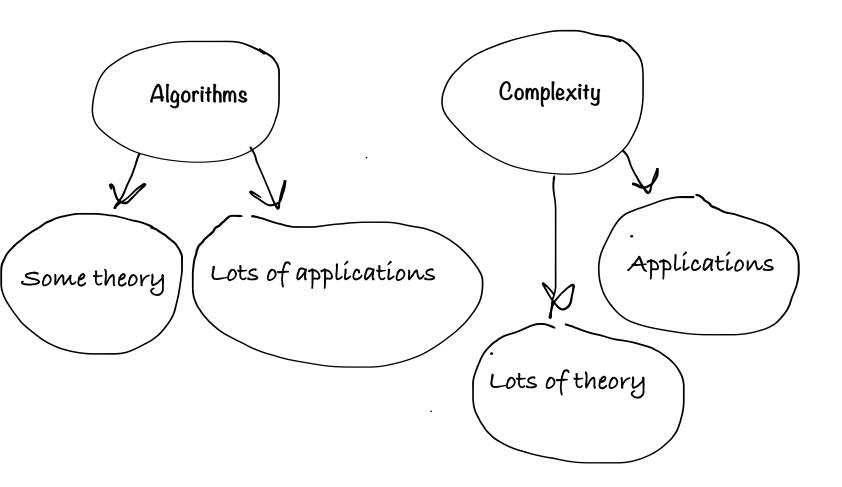
We say that an algorithm has polynomial time-complexity if there is an integer k such that the algorithm, if started with input of size n, runs in $O(n^k)$ steps.

Efficient algorithms:

The standard definition is to say that an algorithm is efficient if and only if it has polynomial time-complexity.

An outline of the course

We have two parts:



Algorithms

We describe several types of algorithms and ways to construct algorithms.

Complexity
Not all problems can be solved efficiently.
Some problems can not be solved at all!
Some of the things we will do in this part are:

Study Turing Machines and formally define computing

Study uncomputable problems.

Study NP-problems which (probably) can not be solved efficiently.

Study so called approximation algorithms.