

Computer Science/Mathematics 3804

Design and Analysis of Algorithms I

Course Outline

Class Schedule

Classroom:	Tory Building 447
Class Times:	Wednesdays and Fridays 4:05 pm - 5:25 pm
Course Website:	http://scs.carleton.ca/~sack/teaching/3804

Instructor Information

Professor	Office	Telephone	Email	Office Hours
Dr. J.-R. Sack	HP 5350	N/A	sack@scs.carleton.ca	Wednesdays 10:30-12:00

Teaching Assistants

T.A.	Office Hours	Room Number
Amin Gheibi	Thursdays 3-5 p.m.	1170 HP
Sarah Liske	Every Wednesday 3-4 after an assignment has been handed back	1170 HP

Course Outline cont'd

Course Description

An introduction to the design and analysis of algorithms.

Topics Covered

Topics include: recurrence relations, sorting and searching, divide-and-conquer, dynamic programming, greedy algorithms, NP-completeness.

Prerequisites

COMP 2002 or COMP 2402, and either COMP 1805 or both of MATH 2007 and MATH 2108, or equivalents.

Textbook (s)

Algorithms by S. Dasgupta, C.H. Papadimitriou, and U.V. Vazirani, (2007) Published by McGrawHill also available online from the first author's web-page

alternate reference – more detailed:

Introduction to Algorithms (3rd Edition) by Thomas H. Cormen, Charles E. Leiserson, Ronald L. Rivest and Clifford Stein (2009). ISBN 0-262-03384-4. Published by MIT Press.

The Appendices A,B, and C provide a useful summary of material (much of it) should be known.

Course Outline cont'd

Component	Weight	Due Date
Assignment 1	10%	Sept 28 th before class
Assignment 2	10%	Oct 19 th before class
Assignment 3	10%	Nov. 9 th before class
Assignment 4	10%	Nov. 30 th before class
Midterm Exam	20%	October 31 st during class
Final Exam	40%	tba

Course Outline cont'd

Excerpts from Calendar; refer to Calendar

September 6 classes begin. **September 19** Last day for registration. Last day to change courses or sections for fall/winter and fall term courses. **September 30** Last day to withdraw from fall term and fall/winter courses with a full fee adjustment. **October 5, 2012** University Day at Carleton. Undergraduate classes suspended. **November 9, 2012** Last day to submit, to the Paul Menton Centre for Students with Disabilities, Formal Examination Accommodation Forms for December examinations. **December 3, 2012** Fall term ends. Last day for academic withdrawal from fall term courses

Final Exam Note

The final exam will be scheduled by the University.

Assignments

Late assignments will not be accepted.

Collaboration Policy

Students are encouraged to collaborate on assignments, but at the level of discussion only. When writing down the solutions, students must do so in their own words.

Computer Science



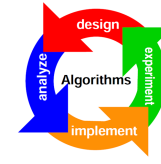
Hardware

- Hardware
- Communication Infra-structure



Software

- Programs
- Software Infra-structure



Efficient algs+data str.

- Algorithms
- Data Structures



Introduction

- “Algorithms change/d the world”
- Precise instructions of how to add, subtract, multiply, divide, extract square roots, get digits of π , ... [Al Khwarizmi, AD 600, Baghdad]
- Unambiguous, precise, mechanical, efficient and correct

Three questions to ask - algorithm

After we understand what it does or is supposed to do.

1. Is it correct?
2. How much time does it take, as a function of the input size
3. Can we do better

A Sequence

- The first 21 numbers are:
0, 1, 1, 2, 3, 5, 8, 13, 21, 34, 55, 89, 144, 233,
377, 610, 987, 1597, 2584, 4181, 6765

What can we say about this sequence?

Many things

- Non-negative
- Increasing
- Rapidly increasing almost as fast as 2^n , in fact a good approximation is: $2^{0.694n}$
- If we call F_n the n^{th} number then

$$F_n = F_{n-2} + F_{n-1} \text{ where } n > 1 \text{ and}$$

$$F_1 = 0 \text{ and } F_0 = 1.$$

Fibonacci Numbers



F_n is called the n^{th} Fibonacci number.

Fibonacci was living between 1170 and 1250.

Fibonacci Occurrences in Nature

Number of Petals	Flower
3 petals (or 2 sets of 3)	lily (usually in 2 sets of 3 for 6 total), iris
5 petals	buttercup, wild rose, larkspur, columbine (aquilegia), vinca
8 petals	delphinium, coreopsis
13 petals	ragwort, marigold, cineraria
21 petals	aster, black-eyed susan, chicory
34 petals	plantain, daisy, pyrethrum
55 petals	daisy, the asteraceae family
89 petals	daisy, the asteraceae family

0, 1, 1, 2, 3, 5, 8, 13, 21, 34, 55, 89, 144, ...

Pineapple

The number of spirals going in each direction is a Fibonacci Number.

Here, there are 13 spirals that turn clockwise and 21 curving counterclockwise.



In sunflowers, the number of clockwise and counterclockwise spirals will always be consecutive Fibonacci Numbers like 21 and 34 or 55 and 34.

Algorithm I for Fibonacci

- Function fib1(n)

```
if n = 0 return 0
```

```
if n = 1 return 1
```

```
return fib1(n-1) + fib1(n-2)
```

The Three Questions

1. Is the algorithm correct?
2. How much time does it take to compute?
3. Can we do better?

1. Is easy that is simply the definition of the Fibonacci sequence.

2. How much time does it take to compute?

- The time is a function of n , let us call it $T(n)$
- $T(n) \leq 2$, for $n \leq 1$ why?
- For larger values of n , i.e., $n > 1$

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

so, we can see that $T(n) \geq F(n)$

bad news?!

- to compute $T(200) \geq F(200) \geq 2^{138}$ steps would be required
- this is huge the NEC Earth Simulator can do 40 trillion steps per second. Thus, to compute $F(200)$ would take at least 2^{92} seconds that is about 78509642268225537474 years.
- The earth is expected to live to 5 billion (5000000000) years only.

So, can we do better?

3. Can we do better?

Why is fib1 so bad?

Function fib1(n)

```
if n = 0 return 0
if n = 1 return 1
return fib1(n-1) + fib1(n-2)
```

a second attempt fib2

Function fib2(n)

```
if n = 0 return 0
```

```
create an array f[0 .. n]
```

```
f[0] = 0, f[1] = 1
```

```
for i = 2 ... n
```

```
    f[i] = f[i-1] + f[i-2]
```

```
return f[n]
```

The three questions

1. Is the algorithm correct?
2. How much time does it take to compute?
3. Can we do better?

1. Is the algorithm correct?

again, here the correctness is self-evident
as it follows from the definition of Fibonacci.

2. How much time does it take to compute?

- Except of the small number of constant operations the main work is carried out in the loop.
 - the loop body is one addition
 - and is executed $n-1$ times

So, the total number of operations taken by fib2 is linear in n .

Careful

- We get that fib2 executes a linear number of additions. However, the numbers are huge.
- Our standard model of analysis assumes that the numbers have a constant number of bits (32 or 64 ...).
- Here the numbers are about $0.694n$ bits long!
- Such numbers cannot be added in one step!
- Adding 2 n -bit numbers takes time proportional to n .

Conclusion

- Lemma: The running time of $\text{fib2}(n)$ is proportional to n^2 .
 - What is the running time of $\text{fib1}(n)$ then?
3. Can we do better? YES, see Exercise 0.4 in book

Recall Big-O

- Let $f(n)$ and $g(n)$ be functions from the positive integers to the reals. We say that $f = O(g)$ if there is a constant $c > 0$ such that $f(n) \leq c * g(n)$.
- This means that f grows no faster than g .
- Also, the constant c allows us to ignore small values of n .

Some examples

- $f(x) = x^2 + \log(x) + 7x + 1000000000$

Course requirement

- I will assume that you are familiar with big-O.

Divide&Conquer Algorithm

- Recall D&C
- Break problem into subproblems that are smaller instances of the same type of problem
- Recursively solve the subproblems
- Combine the answers to the subproblems into an answer to the original problem

Multiplication "x"

Gauss

Multiplication of complex numbers

$$(a+bi)(c+di) = \underset{\uparrow}{ac} - \underset{\uparrow}{bd} + (\underset{\uparrow}{bc} + \underset{\uparrow}{ad})i$$

4 "x"

but, he saw it can be done with 3 "x"

$$\underset{\uparrow}{ac}, \underset{\uparrow}{bd} \text{ and } (a+b)\underset{\uparrow}{(c+d)}$$

Since

$$bc + ad = (a+b)(c+d) - \underset{\checkmark}{ac} - \underset{\checkmark}{bd}$$

Seems insignificant

3 "x" instead of 4 "x"

but, when applied recursively,
it becomes significant!

$$X = x_1 \dots x_{\lfloor n/2 \rfloor} \dots x_n \quad \text{2 } n\text{-bit numbers}$$

$$Y = y_1 \dots y_{\lfloor n/2 \rfloor} \dots y_n$$

assume n is even

$$X = \boxed{x_1 \dots x_{n/2}} \boxed{x_{n/2+1} \dots x_n} = 2^{n/2} X_L + X_R$$

similarly,

$$Y = \boxed{y_1 \dots y_{n/2}} \boxed{y_{n/2+1} \dots y_n} = 2^{n/2} Y_L + Y_R$$

$$XY = (2^{n/2} X_L + X_R)(2^{n/2} Y_L + Y_R)$$

$$= 2^n \underset{\uparrow}{X_L Y_L} + 2^{n/2} (\underset{\uparrow}{X_L Y_R} + \underset{\uparrow}{X_R Y_L}) + \underset{\uparrow}{X_R Y_R}$$

$$T(n) = 4T(n/2) + O(n)$$

↑

4 "x" of 2 $\frac{n}{2}$ -bit numbers

$$O(n) \leftarrow \begin{cases} 2^n \text{ or } 2^{n/2} \cdot \text{ is a shift each} \\ \Rightarrow \text{linear} \\ \text{addition linear} \end{cases}$$

$$T(n) = 4T\left(\frac{n}{2}\right) + O(n) \stackrel{\text{see later}}{=} O(n^2)$$

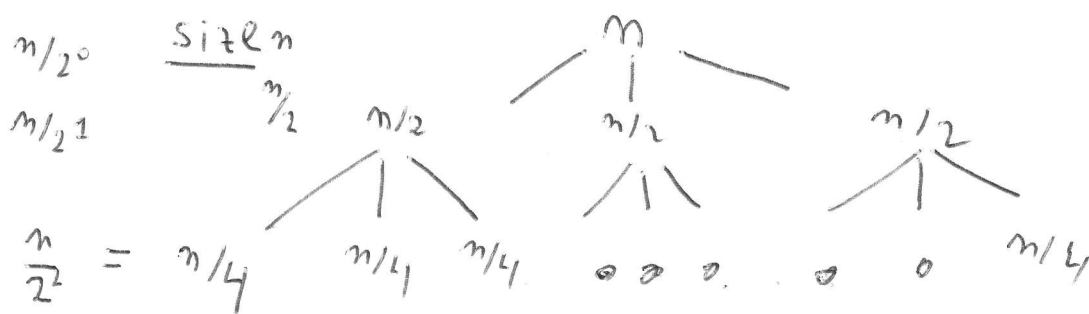
Now, Gauss \Rightarrow 3 "x" only

$$X_L Y_L, X_R Y_R \text{ and } (X_L + X_R)(Y_L + Y_R)$$

\Rightarrow

$$T(n) = 3T\left(\frac{n}{2}\right) + O(n) = O(n^{1.59})$$

Recursion tree



of problems

$$1 = 3^0$$

$$3 = 3^1$$

$$9 = 3^2$$

<u>level k:</u>	problem size	$\frac{n}{2^k}$	(root at level 0)
	# of problems	3^k	
# of levels	$\log_2 n$		

How much work is done?

level 0	$O(n)$
1	$\frac{3}{2} n$
⋮	
k	$3^k * O(\frac{n}{2^k}) = (\frac{3}{2})^k * O(n)$
⋮	
$\log_2 n$	$3^{\log_2 n} * O(1) = n^{\log_2 3}$

geometrically increasing series
 $\rightarrow O(n^{\log_2 3}) = O(n^{1.59})$

Thus, the run-time of multiplying 2 n -bit numbers is not quadratic, but using Gauss + D & C $O(n^{1.59})$

can still be improved using Fast Fourier transform (D & C)

Note:

$$3^{\log_2 n} = n^{\log_2 3}$$

Why?

$$\log_3 (3^{\log_2 n}) = \log_3 (n^{\log_2 3})$$

$$\log_2 n = \log_2 3 \log_3 n$$

$$\frac{\log_2 n}{\log_2 3} = \log_3 n$$

The last equation holds
base transform

$$\frac{\log_a x}{\log_a b} = \log_b x$$

Proof:

$$b^{\log_b x} = x$$

$$\log_a b \cdot \log_b x = \log_a x \quad \square$$

Recurrence Relations

Many D & C algorithms can be analyzed as follows:

Let a = # of subproblems
 m/b = size of subproblems
 n = total size

$$T(n) = a \cdot T(n/b) + O(n^d)$$

then,

$$T(n) = \begin{cases} \text{(A)} O(n^d) & \text{if } d > \log_b a \\ \text{(B)} O(n^d \log n) & \text{if } d = \log_b a \\ \text{(C)} O(n^{\log_b a}) & \text{if } d < \log_b a \end{cases}$$

Master Theorem

Proof (sketch)

assume w.l.o.g. $n = b^j$, for some j .

work on level k of recurrence tree

$$a^k \cdot O\left(\frac{n}{b^k}\right)^d$$

\uparrow # of sub-problems on level k

\uparrow their size

Work due to $O(n^d)$

$$a^k \cdot O\left(\frac{n}{b^k}\right)^d = O(n^d) \left(\frac{a}{b^d}\right)^k$$

$$k = 0, \dots, \log_b n$$

\Rightarrow geometric series with ratio $\frac{a}{b^d}$

Cases :

① $\frac{a}{b^d} < 1 \Rightarrow$ sum is the first term i.e. $O(n^d)$

② $\frac{a}{b^d} = 1 \Rightarrow$ all $O(\log n)$ terms are equal to $O(n^d)$

③ $\frac{a}{b^d} > 1 \Rightarrow$ Series is increasing
 and its sum = last term
 $O(m^{\log_b a})$

$$\begin{aligned} m^d \left(\frac{a}{b^d}\right)^{\log_b m} &= m^d \left(\frac{a^{\log_b m}}{(b^{\log_b m})^d}\right) \\ &= a^{\log_b m} = a^{\log_a m \log_b a} \\ &= m^{\log_b a} \end{aligned}$$

□

Recall Geometric Series

$$1 + r + r^2 + \dots$$

$$= \lim_{n \rightarrow \infty} (1 + r + \dots + r^n)$$

$$= \lim_{n \rightarrow \infty} \frac{1 - r^{n+1}}{1 - r}$$

$$r^{n+1} \rightarrow 0, \text{ for } |r| < 1$$

$$a + ar + ar^2 + \dots + ar^{n-1} = a \frac{1 - r^n}{1 - r}$$

Examples Master Theorem

I. Binary Search

$$T(n) = T(n/2) + O(1)$$

$$a = 1 \quad b = 2 \quad d = 0$$

$$\log_b a = \log_2 1 = 0$$

$$d = 0 = \log_2 1 \Rightarrow$$

Case B:

$$\begin{aligned} T(n) &= O(n^d \log n) \\ &= O(\log n) \quad d = 0 \end{aligned}$$

Binary Search : $O(\log n)$

II Merge Sort

function mergesort ($a[1..n]$)

input: An array of numbers $a[1..n]$

output: A sorted array of those numbers

if $n > 1$

return merge(mergesort($a[1..L^{n/2}]$),
mergesort($a[L^{n/2}+1, \dots, n]$))

else

return a

function merge ($x[1..k], y[1..l]$)

if $k = 0$ return $y[1..l]$

if $l = 0$ return $x[1..k]$

if $x[1] \leq y[1]$

return $x[1] \circ \text{merge}(x[2..k], y[1..l])$

else return $y[1] \circ \text{merge}(x[1..k], y[2..l])$

\circ = concatenation

Observation 1) function merge is linear in $k+l$, i.e. the run time is $O(k+l)$

2) function mergesort's run time is given by

$$T(n) = 2T\left(\frac{n}{2}\right) + O(n)$$
$$= O(n \log n)$$

Theorem Merge Sort sorts n numbers in $O(n \log n)$ time.

"The three questions"

① Is the algorithm correct?

< see class >

② Time see above $O(n \log n)$

③ Can we do better?

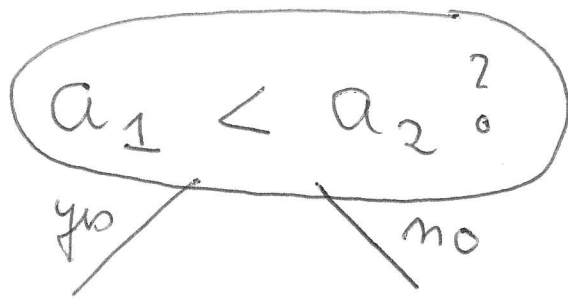
A lower bound for sorting

We will show that no algorithm can exist that sorts n numbers with "fewer than $n \log n$ comparisons".

So, $\Omega(n \log n)$ is a lower bound on the number of comparisons required to sort n numbers.

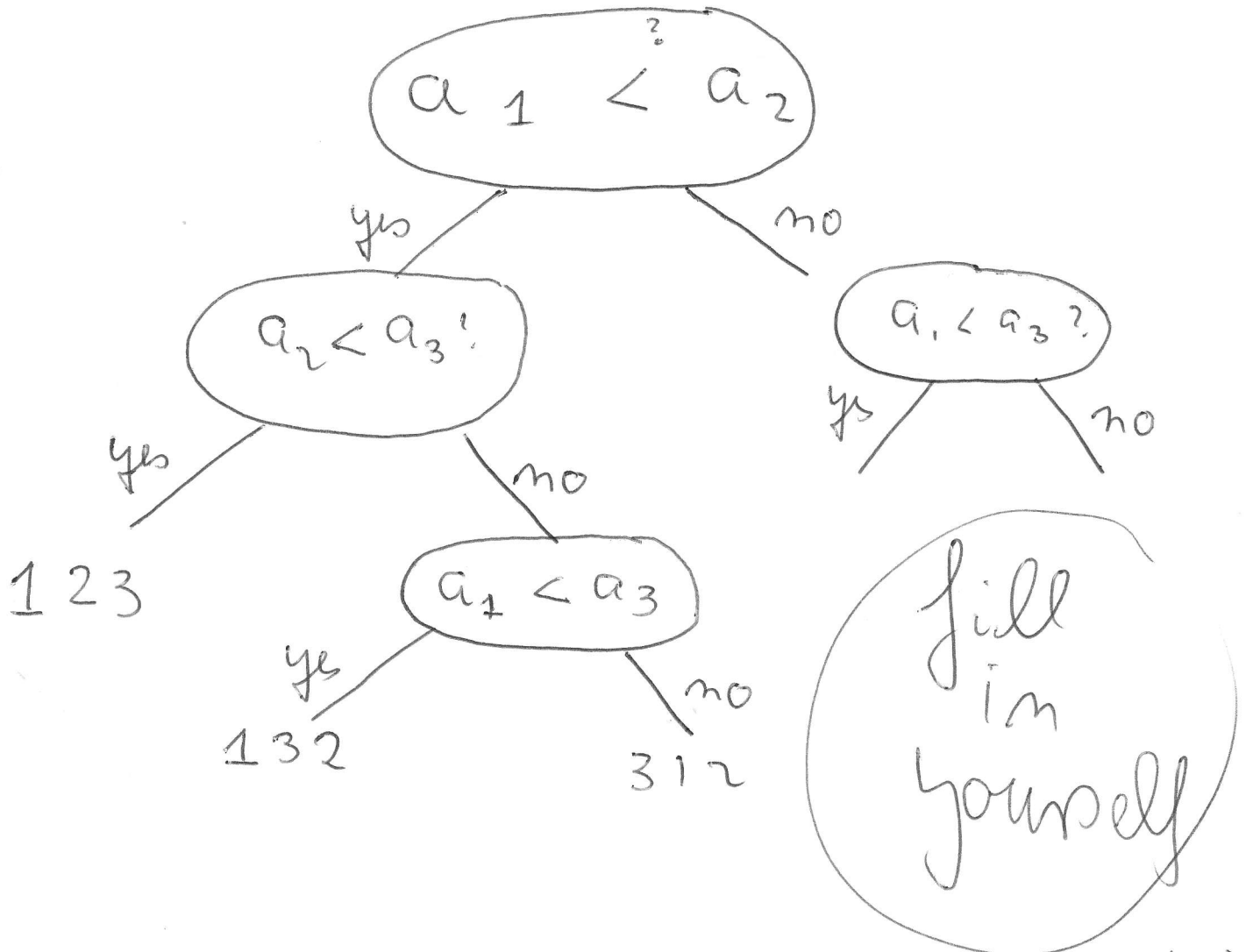
(numbers or elements we can compare)

Comparison



($a_1 \geq a_2$)
if elements are distinct $a_1 > a_2$

Example Sorting 3 elements
1, 2, 3 stored somehow in a_1, a_2, a_3



Consider now sorting n comparable elements.

Any algorithm that is based on comparisons that sorts these elements has an execution that can be described by such a comparison tree.

Every permutation of the n elements is possible & the algorithm must be able to sort them.

- ! o Any permutation must appear at a leaf.
- o There are $n!$ leaves.

Comparison trees are binary trees.

Binary tree of depth d has at most 2^d leaves.

$$\Rightarrow \text{depth of tree} \geq \log(n!)$$

$$\log(n!) \geq c \cdot n \log n, \text{ for some } c > 0$$

Theorem: Any comparison-based sorting algorithm has an $\Omega(n \log n)$ lower bound, to sort n element.

Corollary: Mergesort is an optimal comparison-based sorting algorithm.

Why is $\log_{n/2} (n!) \geq c \cdot n \log n$?

$$n! \geq \binom{n}{n/2}$$

$$\log n! \geq n/2 \log n/2$$

a better bound is derived from Stirling's approximation

$$n! \approx \sqrt{\pi (2n + \frac{1}{3})} \cdot n^n \cdot e^{-n}$$

Median(s)

Median (informal def)
of a list of numbers is a
number s. t. half the numbers
are larger than it, and half
are smaller.

More precisely (and correctly)

The median (lower median)
of a set of n numbers, is
the element in position $\lfloor n/2 \rfloor$
after sorting the elements in
increasing order.

Motivation:

see class

Algorithms for Median

① $\langle \text{brute force} \rangle$ see class

② A randomized D&C algorithm
for selection

Selection is more general than
median finding

Selection (S, k)

input. S : set of n elements (orderable)

k : input parameter

output: k^{th} smallest element of S

$$\text{Selection}(S, k) = \begin{cases} \text{Selection}(S_L, k) & \text{if } k \leq |S_L| \\ \checkmark & \text{if } |S_L| < k \leq |S_L| + |S_v| \\ \text{selection}(S_R, k - |S_L| - |S_v|) & \text{if } k > |S_L| + |S_v| \end{cases}$$

$$S_L = \{x \in S \mid x < v\}$$

$$S_v = \{x \in S \mid x = v\}$$

$$S_R = \{x \in S \mid x > v\}$$

(duplicates of v in S)

Example.

see class

Runtime. S_L, S_V, S_R can be
computed in linear time + space.

After the split the runtime is
proportional to $\max\{S_L, S_R\}$

① we are lucky

$$|S_L| = |S_R|$$

$$T(n) = T(n/2) + O(n)$$

$$= \quad ?$$

(see class)

② we are unlucky

$$\text{one of } |S_L|, |S_R| = 1 \quad (\text{or } 0)$$

e.g. for medic

$$n + (n-1) + \dots + (n/2) = \Theta(n^2)$$

(3) we are sort of lucky

v is called good if it is between the $n/4$... and $3n/4$ element every time (n changes)

then $T(n) \leq T(\frac{3}{4}n) + O(n)$
still good! $O(\log n)$

Lemma On average a fair coin needs to be tossed two times before a "heads" is seen.

Proof. Let $E = \text{exp. \# of } \text{exp}$ tosses before heads is seen

$$E = 1 + \frac{1}{2} E \Rightarrow E = 2$$

\uparrow
1 toss
if heads
done

\uparrow
in half
the cases
we recurse

Therefore, after 2 split operations on average, the array will shrink to at most $\frac{3}{4}$ of its size

(a randomly chosen v has a probability of 0.5 to be good)
so 2 random choices of v on average suffice

Theorem. The expected run-time of the Selection Algorithm is $O(n)$

NOT in text book

Instead of an expected running time, we can achieve a guaranteed linear running time for selection.

Select(S, k)

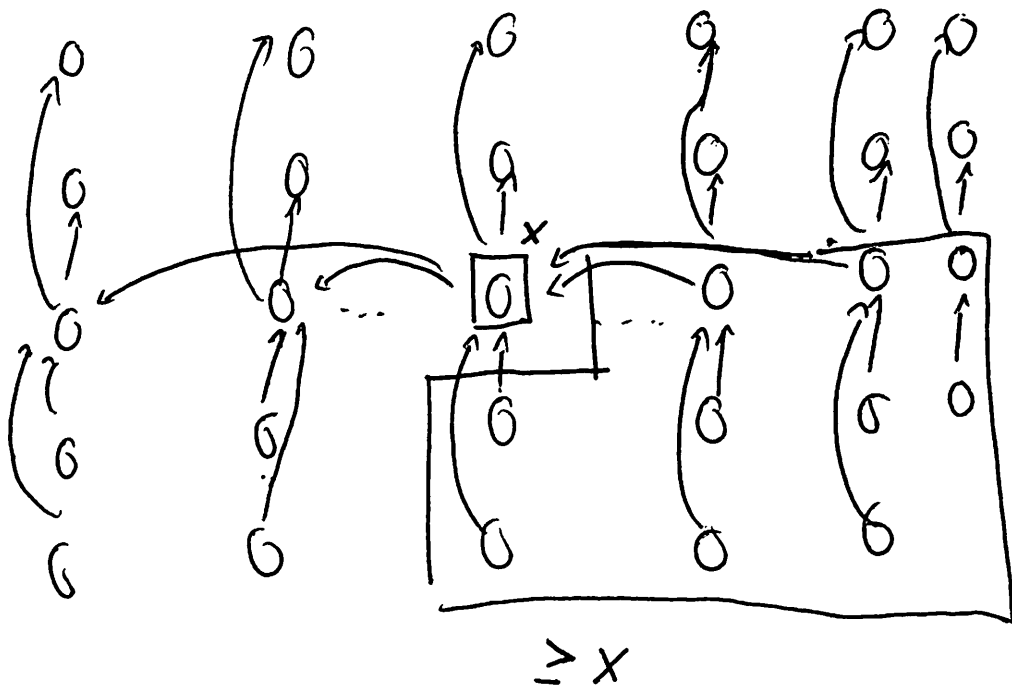
1. Divide the n -element input array into $\lfloor n/5 \rfloor$ groups of 5 elements each + one group of the remaining $n \bmod 5$ elements. illustration see class
2. Find the median of each of the 5 element group by sorting them (say insertion sort; $O(n)$)

3. Recursively find the median x of the $\lceil n/5 \rceil$ medians (from 2.)

4. Partition the input array around x (from 3.)

Let $j := \#$ of elements $\leq x$; call the set S_L
 $n-j :=$ the remaining elements; call this set S_R

5. if $j < k$ recursively call
Select ($S_R, k-j$)
 $j \geq k$ (S_L, k)



$$\underset{\substack{\uparrow \\ \text{elements}}}{3} \cdot \left(\underbrace{\left\lceil \frac{1}{2} \left\lceil \frac{n}{5} \right\rceil \right\rceil}_{\substack{\text{half of} \\ \text{the group}}} - 2 \right) \geq \frac{3n}{10} - 6$$

\uparrow two groups not counted that have fewer than 5 elements or contain x

of elements guaranteed to be $\geq x$ is $\frac{3n}{10} - 6$

analogously, for the # of elements guaranteed to be $\leq x$

Analysis :

Steps 1, 2 and 4 : $O(n)$

Step 3 : $\Theta(T(\lceil n/5 \rceil))$

Step 5 : $T(\underbrace{\frac{7n}{10} + 6}_{\text{why?}})$

Notes:

$$\frac{7n}{10} + 6 < n \quad \text{for } n > 20$$

Sorting say 80 numbers $O(1)$

$$T(n) = \begin{cases} \Theta(1) & \text{if } n \leq 80 \\ T(\lceil n/5 \rceil) + T(\frac{7n}{10} + 6) + O(n), & \text{if } n > 80 \end{cases}$$

$$\begin{aligned}
T(n) &\leq c \lceil n/5 \rceil + c(7n/10 + 6) + O(n) \\
&\leq cn/5 + c + 7cn/10 + 6c + O(n) \\
&\leq 9cn/10 + 7c + O(n) \\
&\leq cn
\end{aligned}$$

by def. of "O"

$$T(n) = O(n)$$

Theorem:

Selecting the k^{th} element in a set of n elements takes $O(n)$ time.

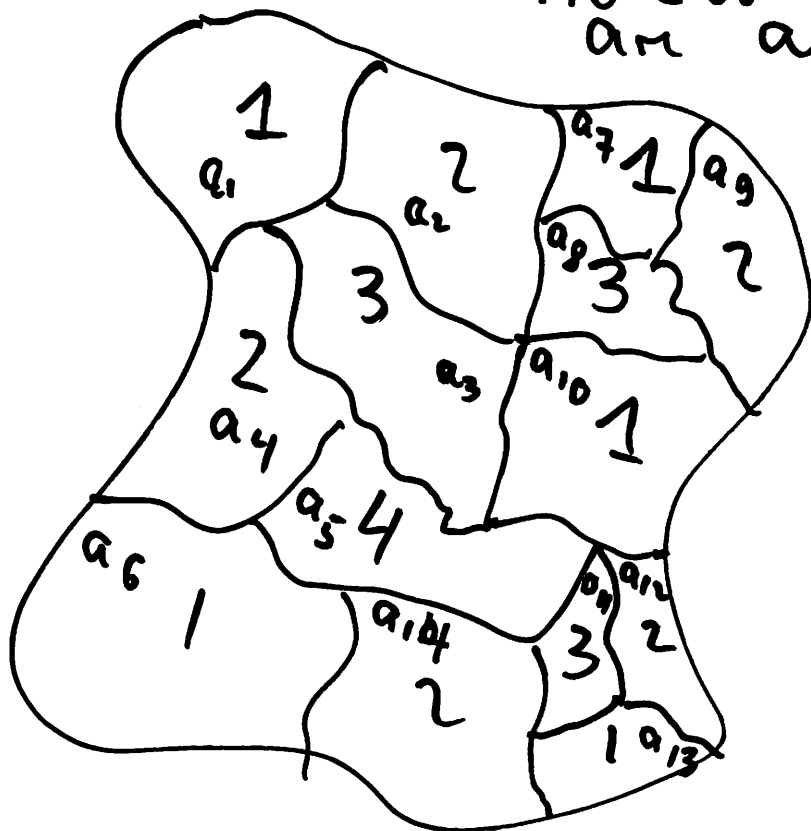
Question. Why does this method not work with groups of 3 elements?

Chapter 3 [in your book]

Graphs have many... many applications.

1. Colouring a map
2. Scheduling an examination set
3. Describing industrial processes
- ⋮

1. Colouring a map



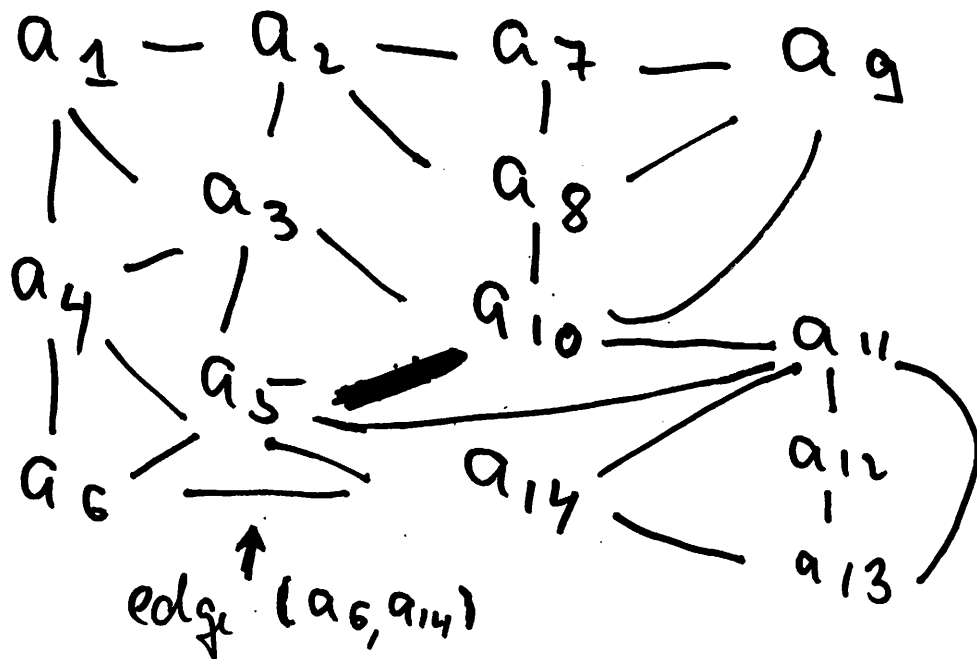
no two countries
are allowed

to share
a border
(except at
a point)
and have
the same
color

This is best do with graph theory.

1) Vertices: countries represent a_1, \dots, a_{13}

2) edges: 2 countries sharing a boundary represent



Graph $G = (V, E)$

V : = set of vertices

E : = set of edges

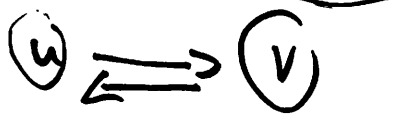
each edge connect a pair of vertices

Directed Graphs

- $(u) \rightarrow (v)$ directed edges
 (u, v) from u to v

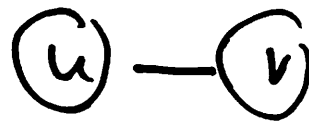
e.g., travel in one-way street

- Undirected edges

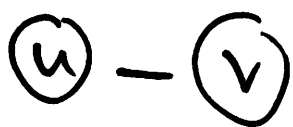


directed
in
both
directions

\cong



same
as
undirected



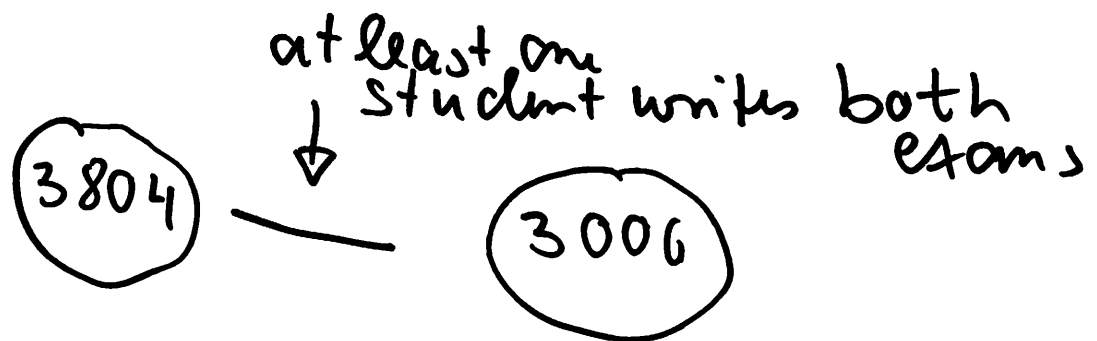
: you can go
from u to v
or from v to u

2. Scheduling Examinations

constraint: no student can write an exam at the same time

Vertices: exams

Edges: a student is taking both exams



Assigning time slots for the exams corresponds to graph colouring

3. Describing industrial Processes

< see class >

Representations of Graph

It is critical to represent a graph correctly.

Size of the graph can be huge.

Ex: Facebook

— June 2012 955M
active users

to represent the face book
graph

Vertices: users

$$|V| = 955M$$

Edges :

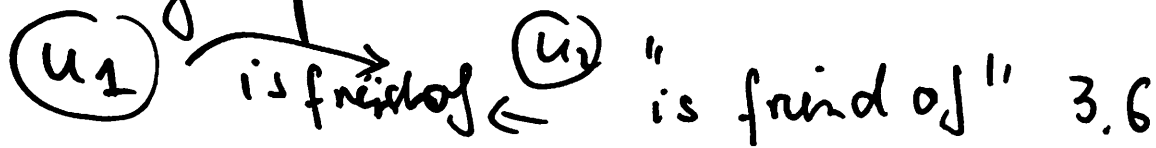
different possibilities
of defining this
say friends!

$$|E| \leq \frac{|V| \cdot |V-1|}{2}$$

if friends are mutual
i.e. undirected graph

$$|E| \leq |V| \cdot |V-1|$$

if it is a directed
graph



So, how many edges does the facebook graph have maximally?

$$|E| \leq |V|(|V|-1) = 955000000 * 954999999$$

$$\approx 9.1 * 10^{17}$$

$$(9.12024999045000000)$$

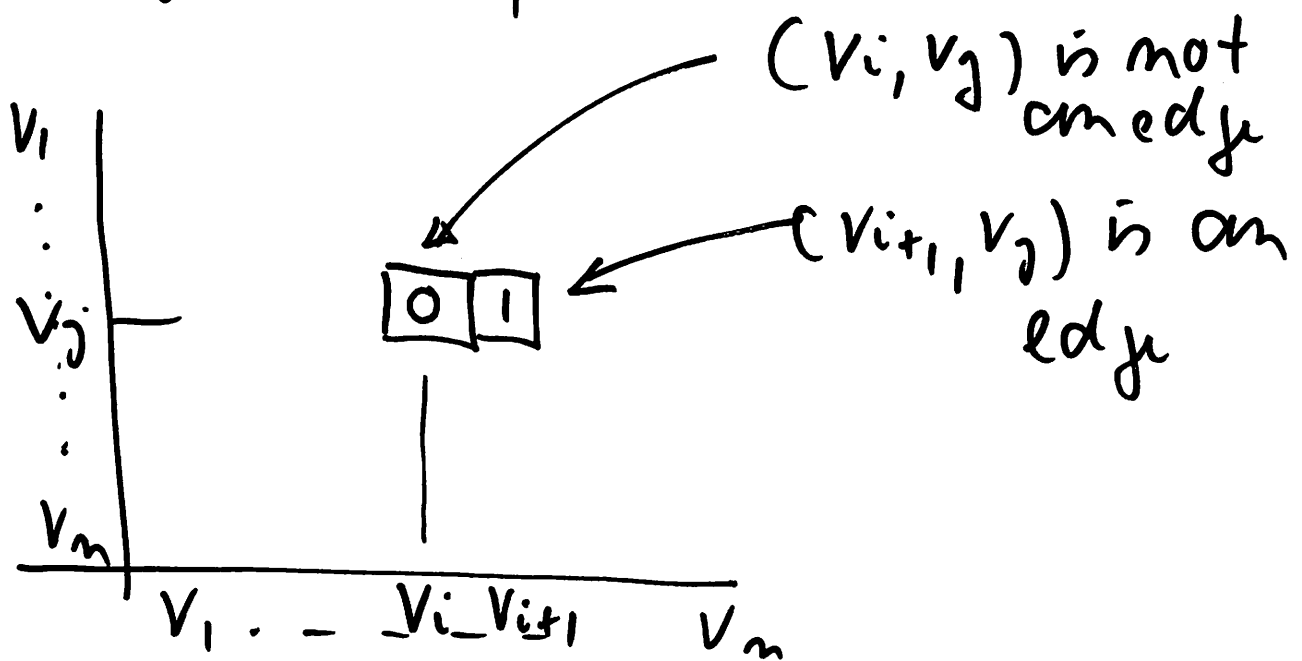
How to store graphs?

For this, we need to know what operations we want to carry out using the graph.

Why?

because the cost of an operation could depend critically on the representation.

I. Bit-array



Storage cost: $|V|^2$ bits

access to an edge $O(1)$

For small graphs this is an efficient representation

For large graph storage cost is prohibitive, esp. if the number of edges is small compared to $|V|^2$.

Operations (Part I)

- is (v_i, v_j) an edge of the graph?
- give me all edges $(v_i, -)$
(so to which vertices is v_i adjacent?)

these need to be executed fast

Representation as Lists

For each vertex v_i : list of
all vertices v_j s.t.
 $(v_i, v_j) \in E$

v_1 : $(v_5) \rightarrow (v_6) \rightarrow (v_{100}) \rightarrow (v_{205})$

.

.

.

.

v_i :

.

.

v_m :

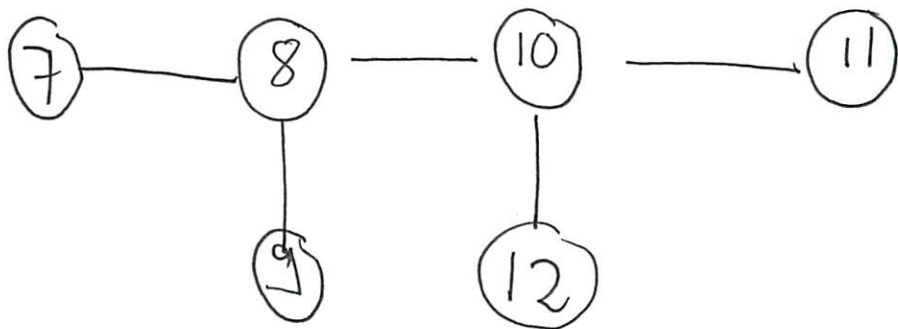
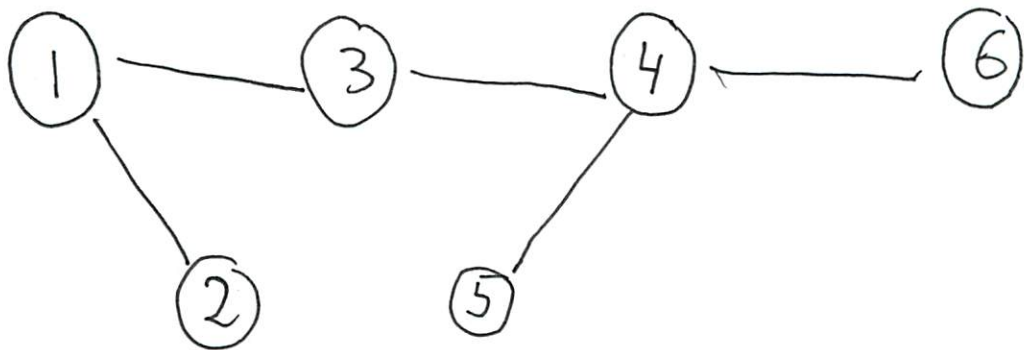
Storage: minimal, as required

however to see if say

$(v_1, v_{100}) \in E$ takes
linear time in the length
of the list for v_1 .

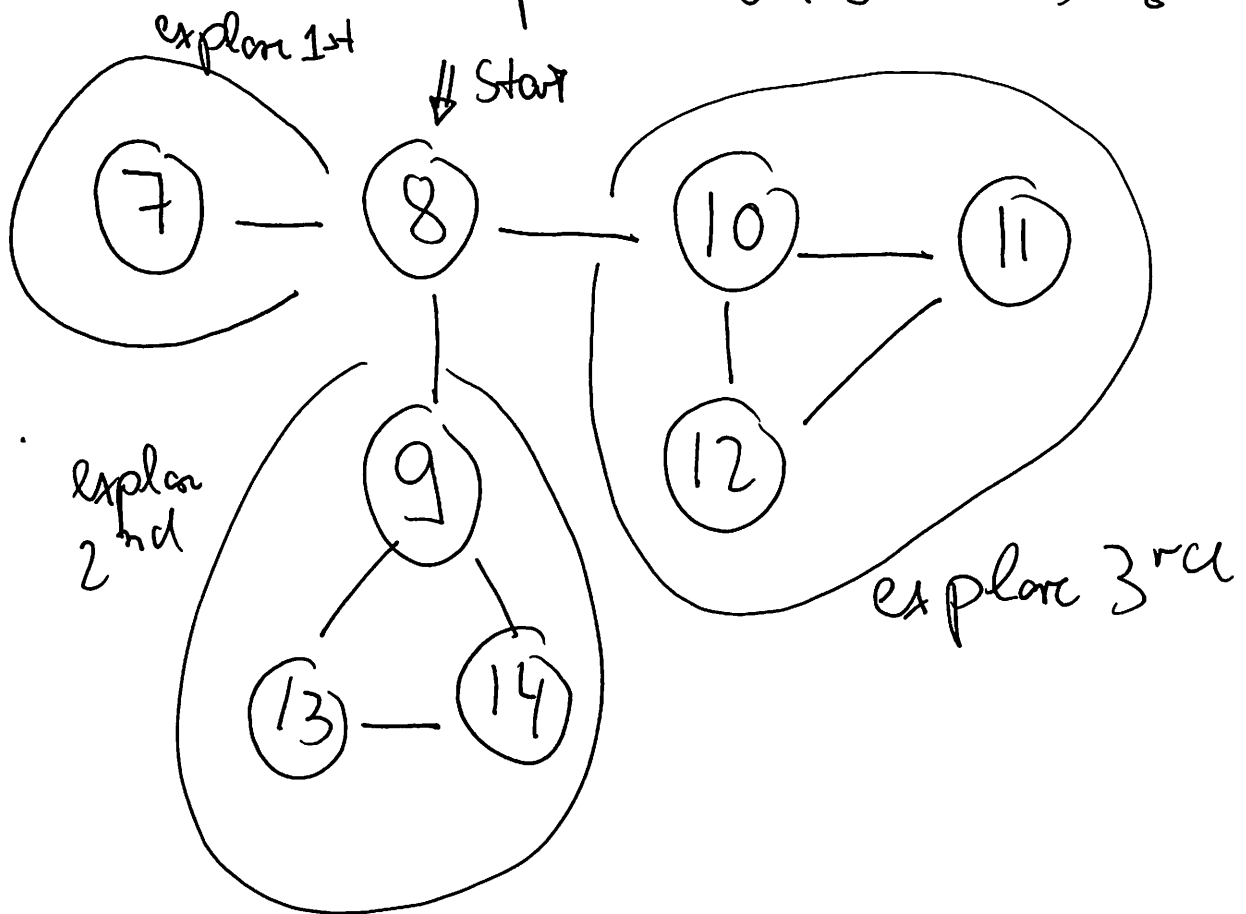
Depth-first Search in Undirected Graph

Q: Which parts of a graph are reachable from a given vertex



reachable from say 8
are nodes 7, 10, 11, 12 and 9

How to implement this?

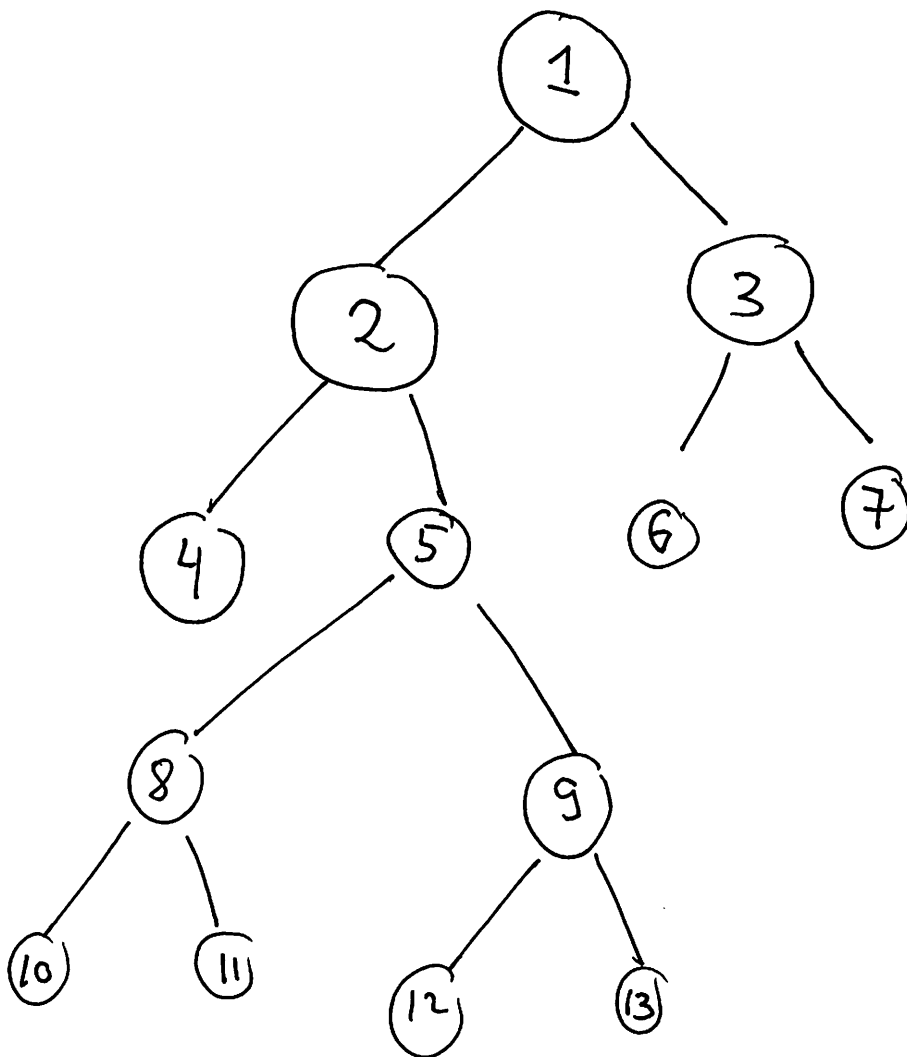


① we could explicitly use a stack

② we can use recursion

One difficulty are cycles because we need to know that we already visited nodes. 3.12

(A) Assume we deal with binary trees, i.e. graphs without cycles and outdegree at most 2



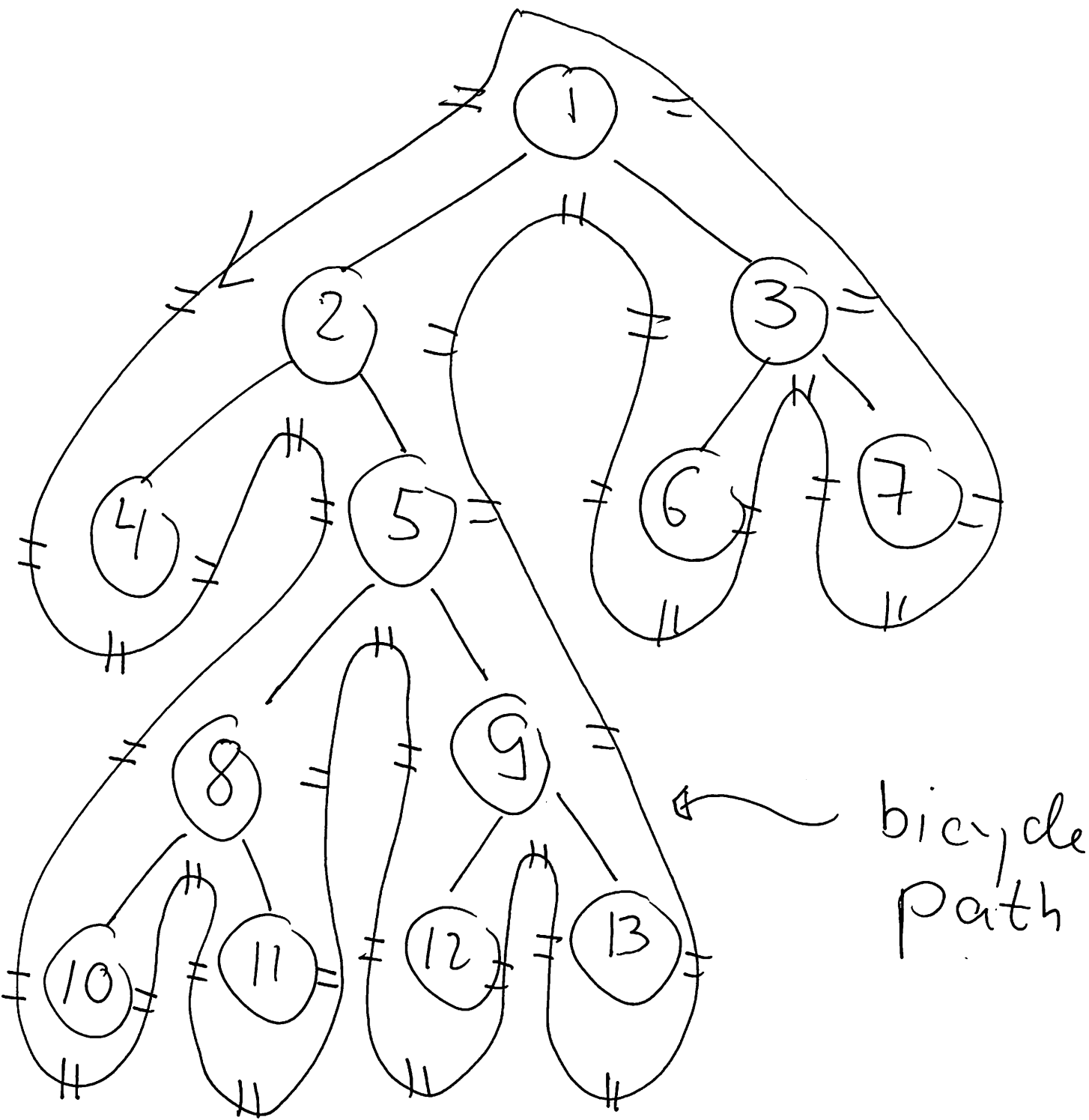
Assume a travel on
the bicycle path

3 methods

① explore each west-
window and print
the elements

② explore each south
window and print
the elements

③ explore each east
window and print
the elements



each node has

3 windows



① produces

1, 2, 4, 5, 8, 10, 11, 9, 12, 13, 3, 6, 7
| | | | | | | | | | | | |

② 4, 2, 10, 8, 11, 5, 12, 9, 13, 1, 6, 3, 7
| | | | | | | | | | | | |

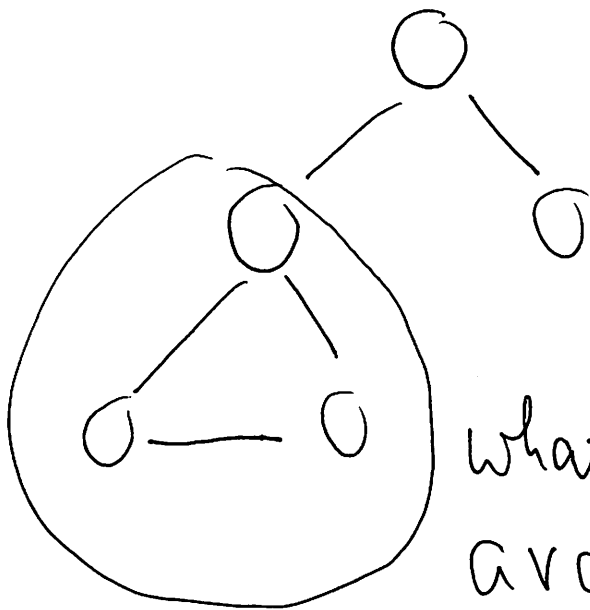
③ 4, 10, 11, 8, 12, 13, 9, 5, 6, 7, 3, 1
| | | | | | | | | | | | |

Algorithms

- ① preorder traversal (T, a)
Visit (a)
preorder traversal ($T, \text{leftchild}(a)$)
preorder traversal ($T, \text{rightchild}(a)$)
- ② inorder traversal (T, a)
inorder traversal ($T, \text{leftchild}(a)$)
Visit (a)
inorder traversal ($T, \text{rightchild}(a)$)
- ③ postorder traversal (T, a)
postorder traversal ($T, \text{leftchild}(a)$)
postorder traversal ($T, \text{rightchild}(a)$)
Visit (a)

This generalizes to trees that have a larger than 2 Outdegree. [How?]

Problem: cycles



what to do here to avoid multiple visits

Idea: Mark what you have already seen

Procedure $\text{Explore}(G, v)$

input: a Graph $G = (V, E)$ and a node (vertex) $v \in V$

output: $\text{visited}(u)$ is set to true for all nodes u reachable from v

$\text{visited}(v) := \text{true}$

$\text{previsit}(v)$

for each edge $(v, u) \in E$:

if not $\text{visited}(u)$: $\text{Explore}(u)$

$\text{postvisit}(v)$

Note: at this point previsit and postvisit are not required

Why is the algorithm correct?

assume that a vertex w is reachable from v but the algorithm does not set $\text{visited}(w)$ to true

Reachability from v means that there is a sequence of edges

$(v, x_1) (x_1, x_2) \dots (x_m, w)$

assume that w is the first node in that which has $\text{visited}(w) \neq \text{true}$

(otherwise, an inductive argument can be made)

then x_m is reachable + $\text{visited}(x_m) = \text{true}$

But $(x_m, w) \in E$ and the loop body would then $\text{explore}(w)$

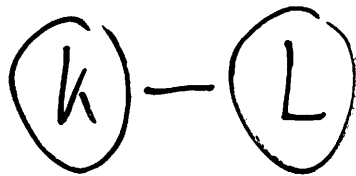
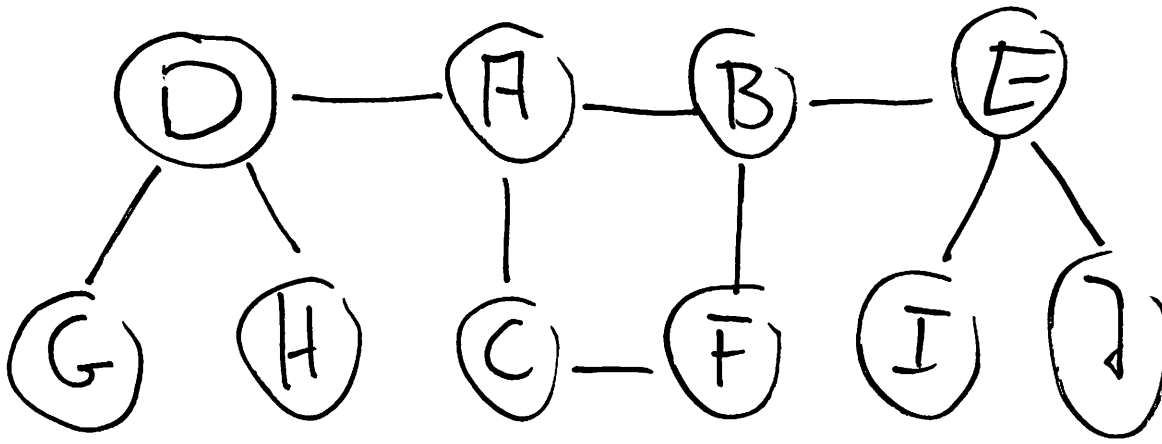
How fast is the algorithm?

$$O(|E|)$$

Can it be improved?

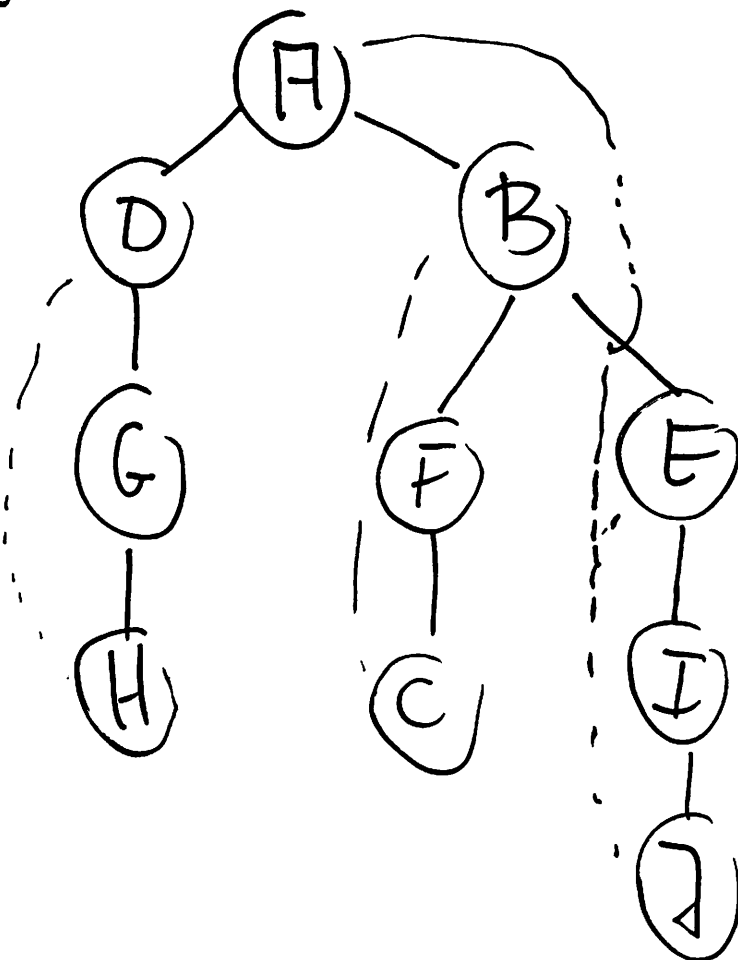
no we need to examine

all edges



Explor (G, A)

yields



Depth-first search

Task: visit all vertices (nodes)
of a graph

Procedure $dfs(G)$

for all $v \in V$

visited(v) = false

for all $v \in V$

if not visited(v): Explore(v)

Is the algorithm correct? Yes
Each vertex is explored and
it is explored only once
(due to visited array)

Complexity!

The loop goes over all
vertices : $O(V)$

Each edge is examined twice
once as (x, y) and
once as (y, x)
(again as in explore)

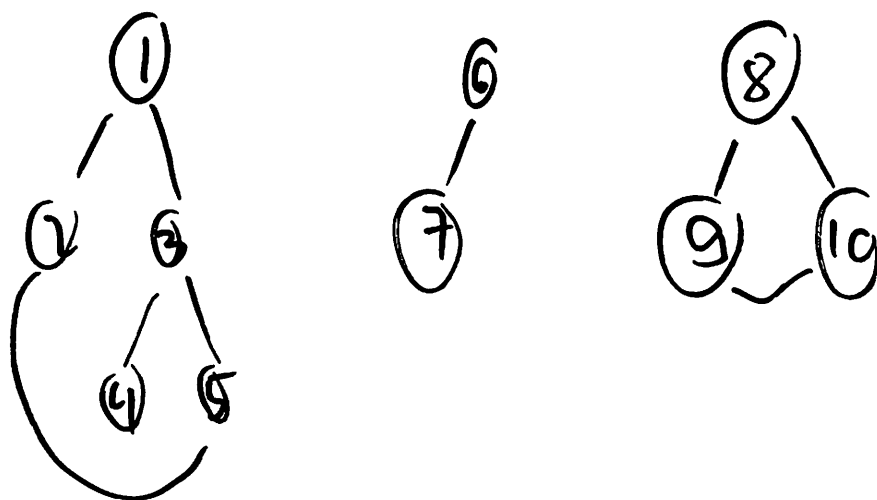
So, $O(|V| + |E|)$

Note: the time per vertex differs in loop

Can we do better? No

$O(|V| + |E|)$ is optimal

Connected Components in undirected graphs



3 Connected Components
 $\{1, 2, 3, 4, 5\}$, $\{6, 7\}$, $\{8, 9, 10\}$

Try to define yourself the
Connected Components
of a graph!

Observation:

The DFS procedure finds all connected components of a graph.

Each time the algorithm starts to examine a new ~~new~~ vertex in the loop-body (for all $v \in V$) that has not been visited before. I.e..

Explore(v) is called, a new connected component is built.

Procedure previsit (v)

$ccnum[v] = cc$

with cc initialized to 0
and then incremented each
time the loop calls explore

previsit will give each vertex
the number of the connected
component it belongs to

Previsit and postvisit Orderings

Procedure previsit (v)

$pre[v] = clock$

$clock = clock + 1$

Procedure postvisit (v)

$post[v] = clock$

$clock = clock + 1$

previsit captures the "time"
of first discovery of a vertex
postvisit captures the "time"
of final departure.

Can you link this to our
discussion of tree traversals?

Master theorem: (general form)

$$T(n) = a T\left(\frac{n}{b}\right) + f(n) \quad \text{where } a \geq 1, b > 1$$

Case 1: $f(n) = O(n^{\log_b a - \epsilon})$ for $\epsilon > 0$

$$T(n) = \Theta(n^{\log_b a})$$

Case 2: $f(n) = \Theta(n^{\log_b a} \log^k n)$

$$T(n) = \Theta(n^{\log_b a} \log^{k+1} n)$$

Case 3: $f(n) = \Omega(n^{\log_b a + \epsilon})$

and it's also true that $a f\left(\frac{n}{b}\right) \leq c f(n)$ for $c < 1$
and sufficiently large n

$$\Rightarrow T(n) = \Theta(f(n))$$

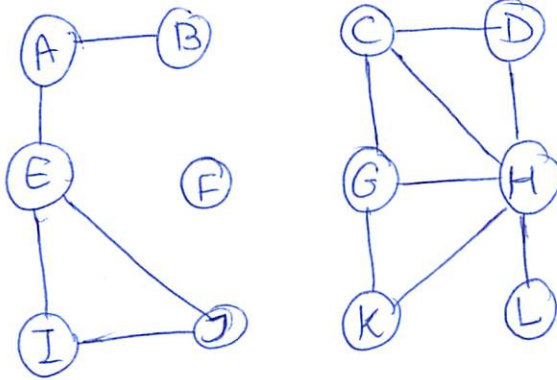
ex: $T(n) = 2T\left(\frac{n}{2}\right) + n^2$ Case 3: $T(n) = \Theta(n^2)$

Announcement: Errata for the text book:

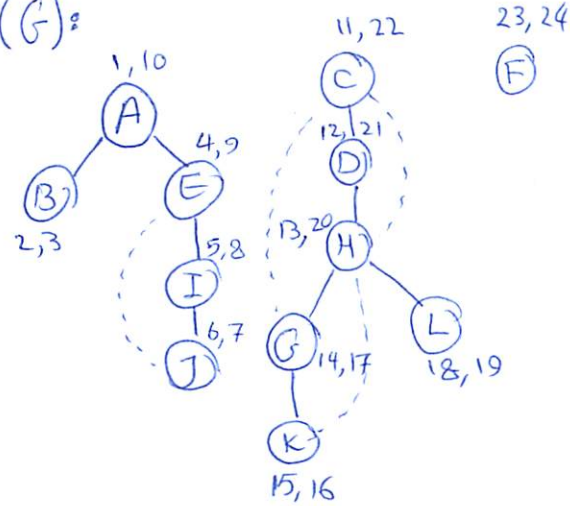
cseweb.ucsd.edu/~dasgupta/book/errata.pdf

Review DFS by an example:

G :



DFS(G):



to collect more information about the graph, we record two type of

- events:
- First discovery (Pre visit)
 - Final departure (Post visit)
- [Pre, Post]
node

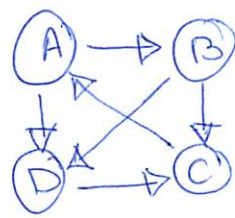
Property 1: $\forall u, v \in V$, two intervals $[Pre(u), Post(u)]$ and $[Pre(v), Post(v)]$ are either disjoint or one is contained in the other one.

why? last-in, first-out behaviour of stack

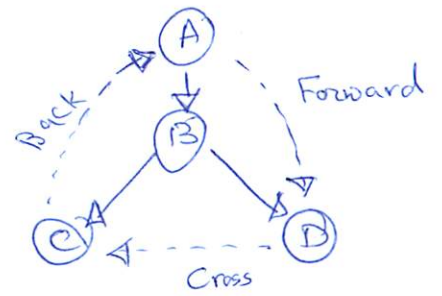
DFS for directed graphs: - What's a directed graph?

Algorithm is the same but movements by edges are in the direction of these edges.

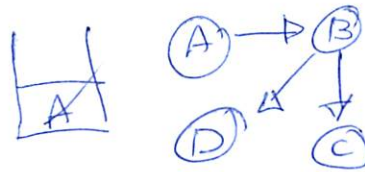
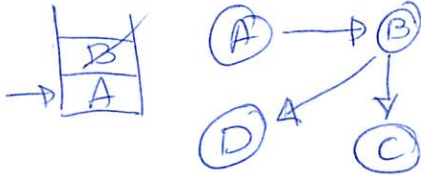
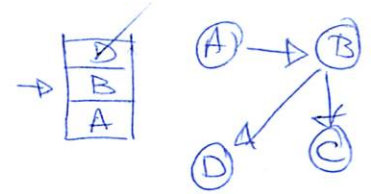
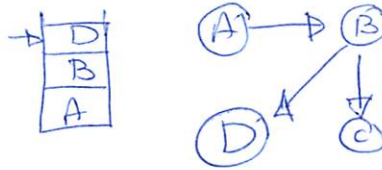
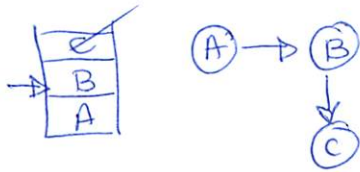
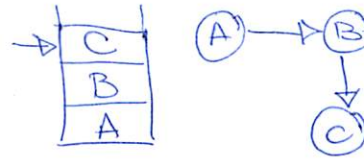
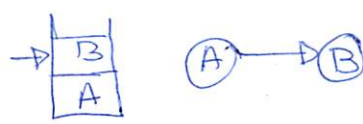
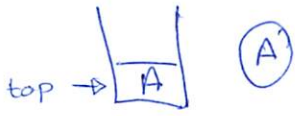
Example:



DFS



DFS steps with stack:



Terminology: - "A" is the root of the DFS tree, everything else is its descendant, and conversely, "A" is their ancestor.

- Parent: direct ancestor (immediate) } A is parent of B
 - child: ~ descendant (~) } B is a child of A

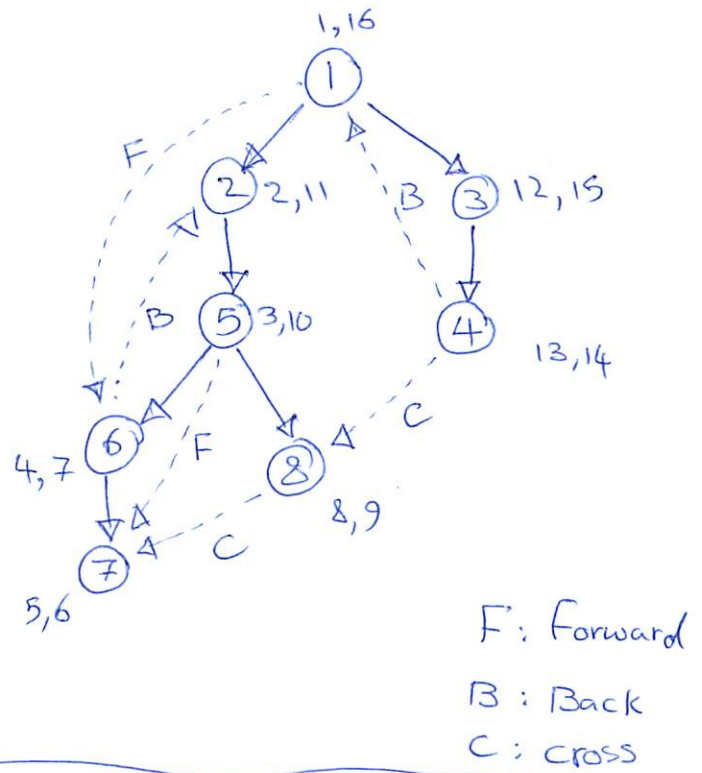
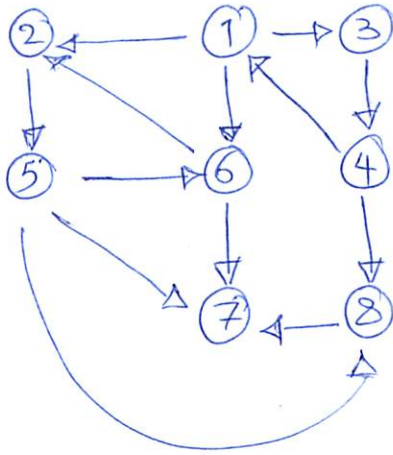
- Source: if a node has only outgoing edges is called source

- Sink: " " " " " " incoming " " " Sink

- Types of edges in DFS forest:

- Tree edges $(u, v) \rightsquigarrow \begin{bmatrix} [&] \\ u & v \end{bmatrix}$
- Forward " $\rightsquigarrow \begin{bmatrix} [&] \\ u & v \end{bmatrix}$
- Back " $\rightsquigarrow \begin{bmatrix} [&] \\ v & u \end{bmatrix}$
- Cross " $\rightsquigarrow \begin{bmatrix} [&] & [&] \\ v & v & u & u \end{bmatrix}$

Example:



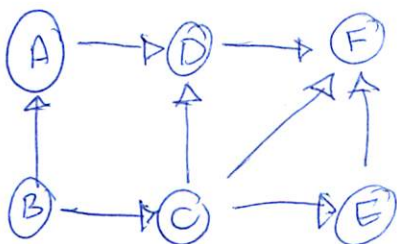
Q: why this pattern $\begin{bmatrix} [&] \\ u & v \\ [&] \\ u & v \end{bmatrix}$ is not possible?

Q: What we can infer from $\begin{bmatrix} [&] \\ u & u \\ [&] \\ v & v \end{bmatrix}$? We infer that v is not accessible from u but the reverse is unknown.

Property 2: A directed graph has a cycle iff its DFS forest has a back edge. Proof? (both directions of proof)

Directed Acyclic Graph (DAG): good for modeling dependencies.

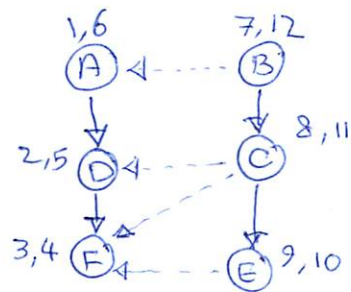
We are interested in an ordering of nodes, that dependencies constraints are satisfied \rightsquigarrow Topological Sort



One orders: B, C, A, D, E, F
 not unique
 B, A, C, D, E, F
 B, C, E, A, D, F

One simple algorithm (in linear time) is to run DFS on DAG and keep ~~nodes~~ nodes in a decreasing order of their post value.

Note: to get a linear-time complexity you should not sort them after running DFS, because we know sorting is $\Omega(n \log n)$. figure out how?



post values: 12, 11, 10, 6, 5, 4
order: B, C, E, A, D, F

Why this algorithm is correct and give us a true order?

- Because only Back edges has a pattern like $post(u) < post(v)$ for (u,v) as an edge. Since it's a DAG \Rightarrow no cycle $\xrightarrow{\text{PROPERTY 2}}$ no Back edge \square

Since we can linearize a DAG, always exist a highest value for post (that's the source) and a lowest value (that's the sink).

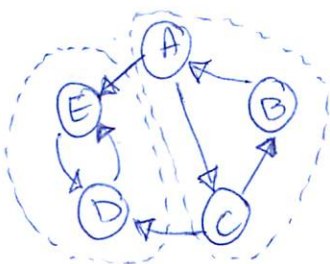
Alternative algorithm for topological sort: - Find a source, output it & delete it
- Repeat until the graph is empty

Strongly Connected Components: (SCC)

Relation: $\forall u, v \in V$, they are connected iff \exists a path from u to v and from v to u ($u R v$)

This relation partition vertices into disjoint sets, called SCC.

Ex.



meta-graph:

It's always a DAG
why?

That's a very nice property that we have two-tiered connectivity structure for directed graphs. It's very helpful to analyze the graph, for example, when you are analyzing web pages graph.

An efficient algorithm to find SCCs:

Observation: If we call `explore()` on a node that lies in a sink SCC, then that component will be retrieved

Q1: How to find a node that is inside a sink SCC? Note that that node is not a sink itself, the containing SCC in meta-graph is a sink.

Q2: How to continue when first SCC is discovered?

To answer Q1: there is no easy way, but we have the following property:

Property 3: If C and C' are two SCCs and there is an edge from C to C' (in meta-graph), then highest post value in C is bigger than highest post value in C' .

Proof:



If C is visited first by DFS, then first node in C will have highest value among all nodes in C and C' .

If C' is visited first, it's clear.

Note: - A source in G , is a sink in G^R (reverse graph)

- G^R has exactly the same SCCs as G

From Property 3, it follows that the node with highest post value is inside a source SCC.

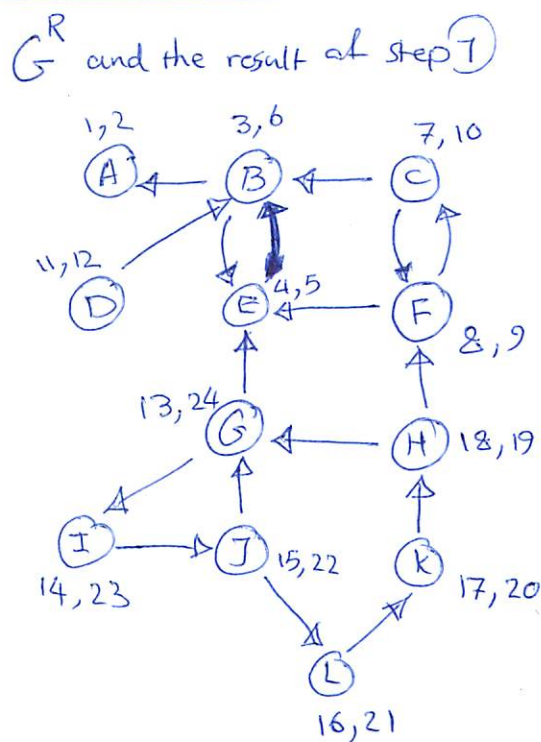
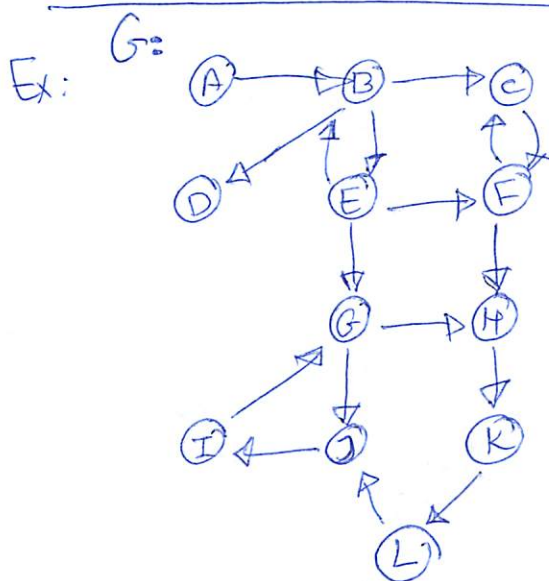
Algorithm:

① Run depth-first search on G^R , get a decreasing order of Post values (highest is inside a source SCC in G^R , that is a sink in G).

② Run DFS on G based on this order:

- you have a counter that shows component's ~~xxx~~ numbers, initial value is zero.

- plus one this counter on each explore() call



Order Based on Post: G, I, J, L, K, H, D, C, F, B, E, A
 SCC-num(\cdot), from Step 2: {1 1 1 1 1 1} {2} {3, 3} {4, 4} {5}

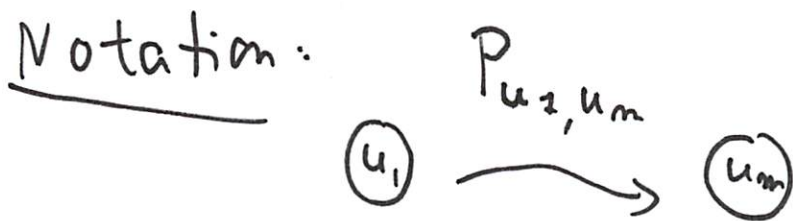
Chapter 4 [textbook]

Paths in Graphs

Let u_1, \dots, u_m be vertices (all distinct) of a graph $G = (V, E)$ so that $(u_i, u_{i+1}) \in E$ $i = 1, \dots, m-1$

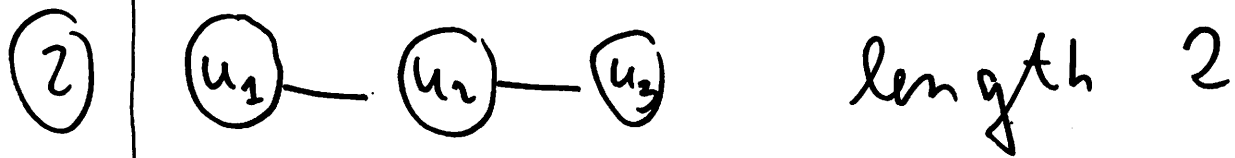
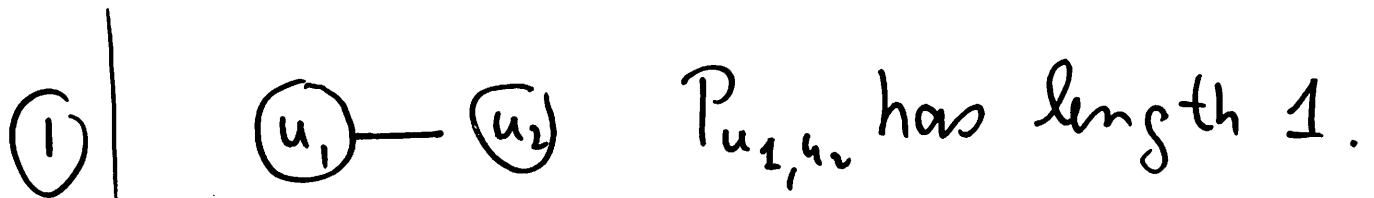


we say that there is a path P_{u_1, u_m} (or $P(u_1, u_m)$) from u_1 to u_m in G .

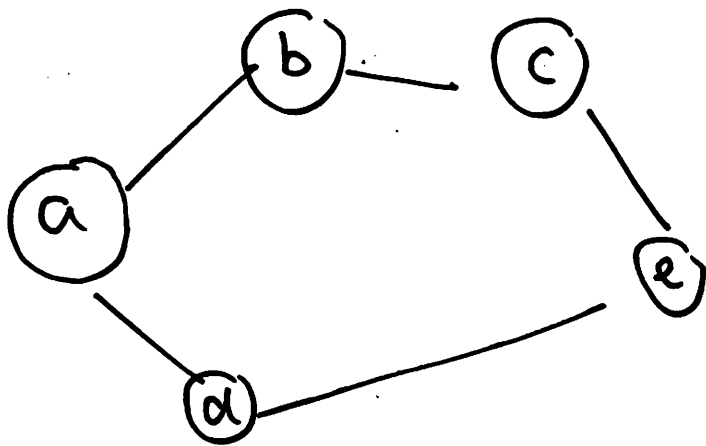


Def The Length of a Path is the number of its edges.

Example:



In general there are many paths between a pair of vertices in a graph.



One path is $a - b - c - e$
another path $a - d - e$

(a)-(b)-(c)-(e) has length: 3

(a)-(d)-(e) " " : 2

Df. The distance between two vertices in a graph is the length of the shortest path connecting them.

Notation. $|P_{a,b}| = \text{length of Path } a \xrightarrow{P_{a,b}} b$
 $d(a,b) = \text{distance between } a \text{ and } b$
 $(= \min\{|P_{a,b}|\} \mid \text{all paths } P_{a,b} \text{ in } G \}$

For the above: $|P_{a,b}| = 3$

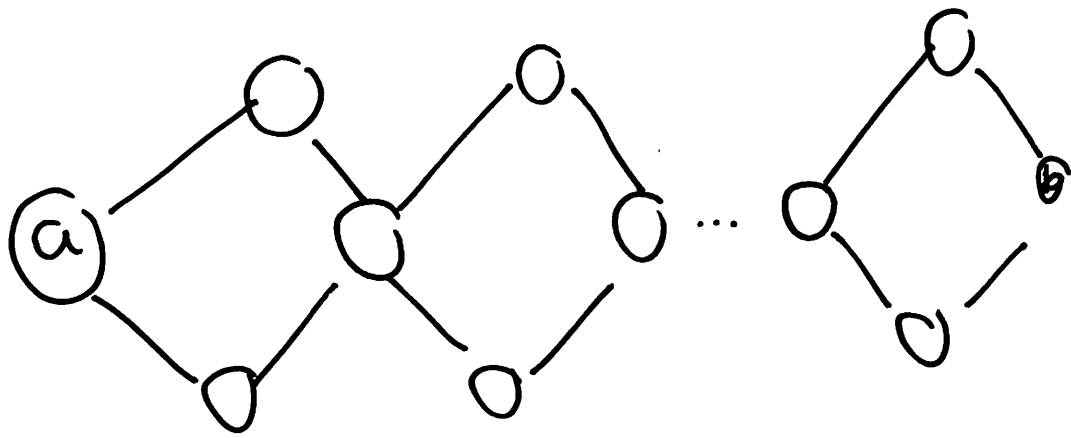
where $P_{a,e} = (a)-(b)-(c)-(e)$

but distance $d(a,e) = 2$

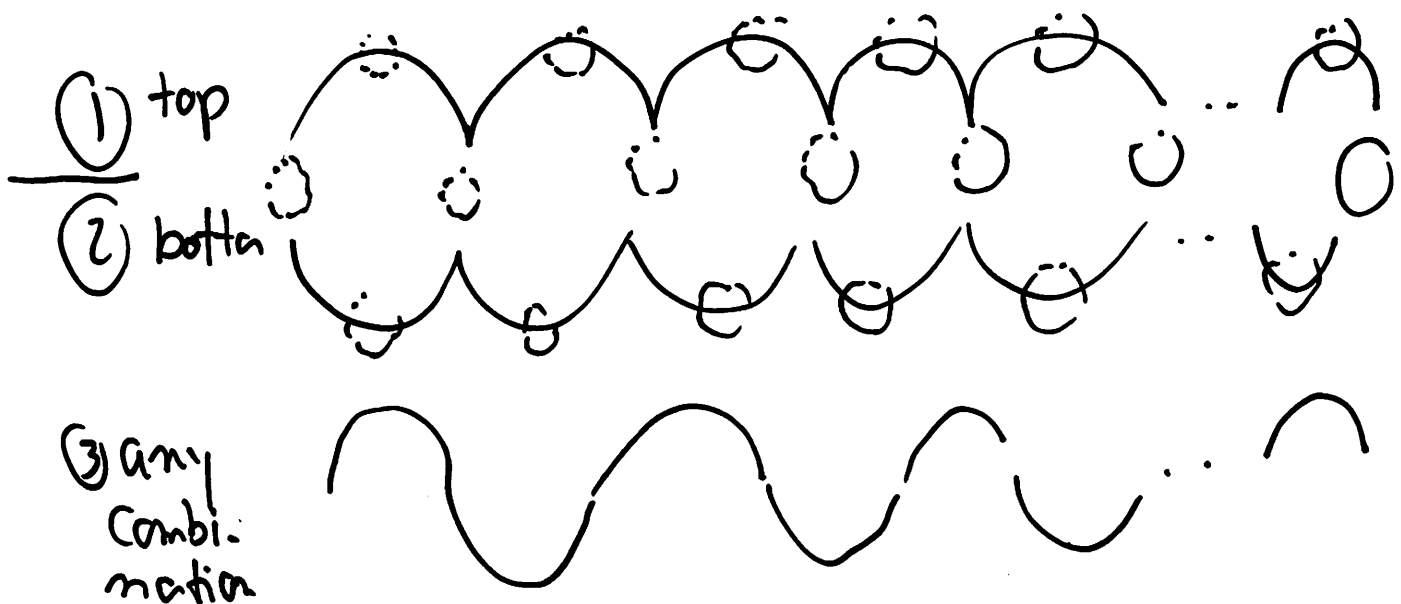
because the shortest path $P_{a,e}$ has 4.3

length = 2, (a)-(d)-(e)

How many paths could there be, in a graph G , connect two vertices a and b ?



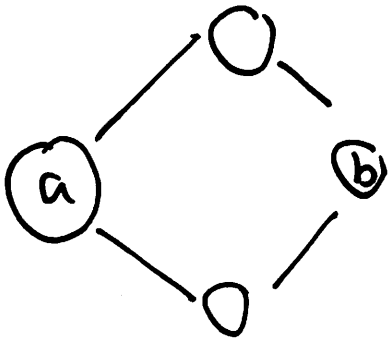
Exponentially many!



How many exactly!

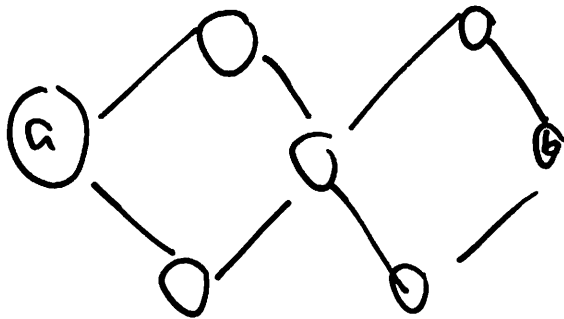
assume # of vertices $n = 1 + k \cdot 3$

Let $k = 1$ then $n = 4$



2 paths
(top, bottom)

(for $k = 2$ $n = 7$



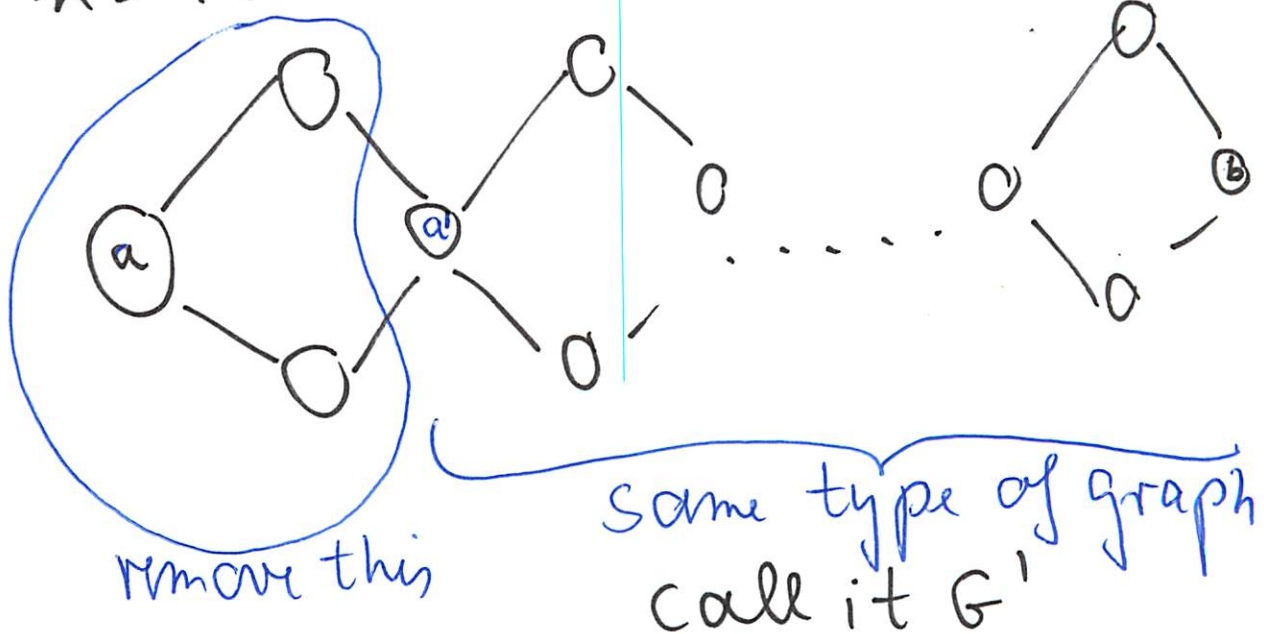
4 paths
top top ∞
top bottom ∞
bottom top ∞
bottom bottom ∞

Claim: Let $n = 1 + k \cdot 3$ be the number of vertices in this type of graph. Then there are 2^k different paths from a to b .

Proof: $k = 1$ we got $2^1 = 2$ paths

Assume that for $k-1$ we get 2^{k-1} different paths.

Consider now the graph with $n = 1 + k \cdot 3$ vertices.



Any path from a to b in G must go through a' .

Now by induction, there are 2^{k-1} paths in G' from a' to b .

There are 2 paths from a to a' in G . Any of these 2 paths can be concatenated with a path from $a' \rightsquigarrow b$ in G' but also in G .

these are paths from $a \rightsquigarrow b$ in G .

So, the number of paths in G from a to $b = 2 \cdot 2^{k-1} = 2^k$

□

So, there are graphs with an exponential number of paths between pairs of vertices.

So how to find the distance between 2 vertices?

Algorithm "STOPID" or
"BRUTE FORCE"

1. Generate all paths
2. Determine their lengths and compute the minimum

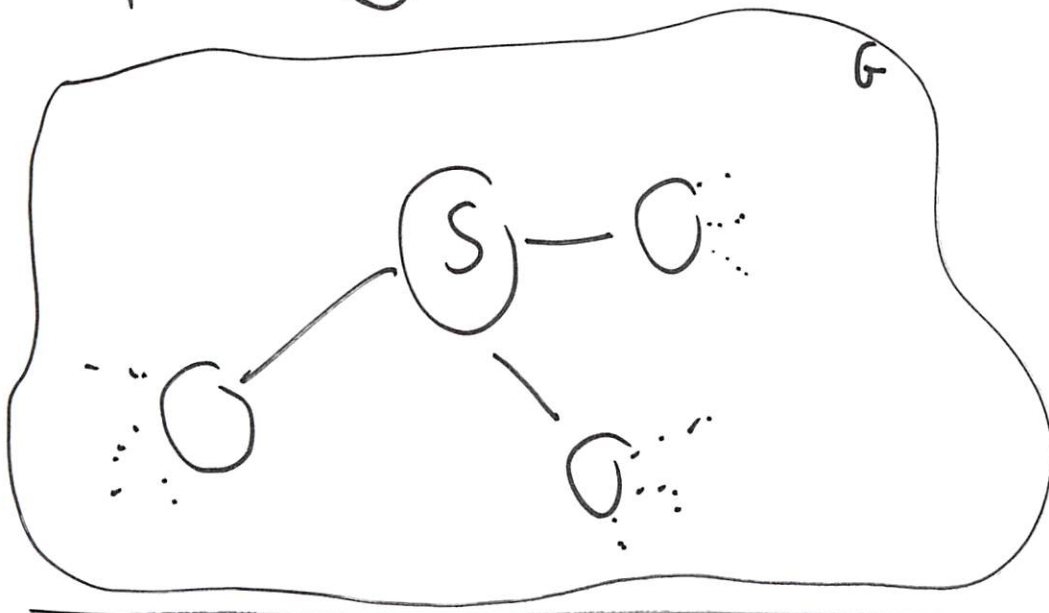
Time Complexity: exponential

can we do better?

Consider a vertex, s , called **SOURCE**.



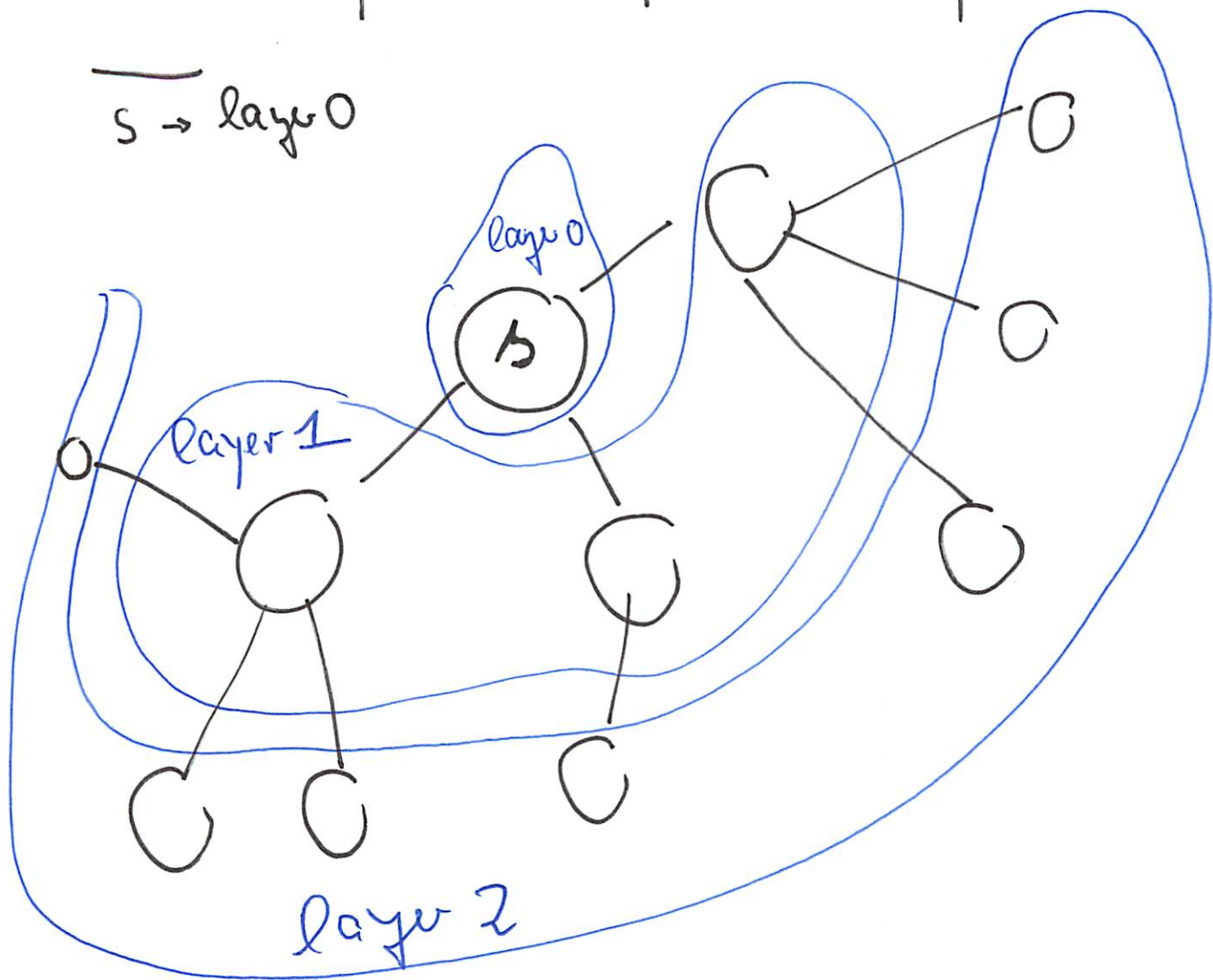
Some vertices are at distance 1 from s .



let us call these vertices layer 1 vertices.

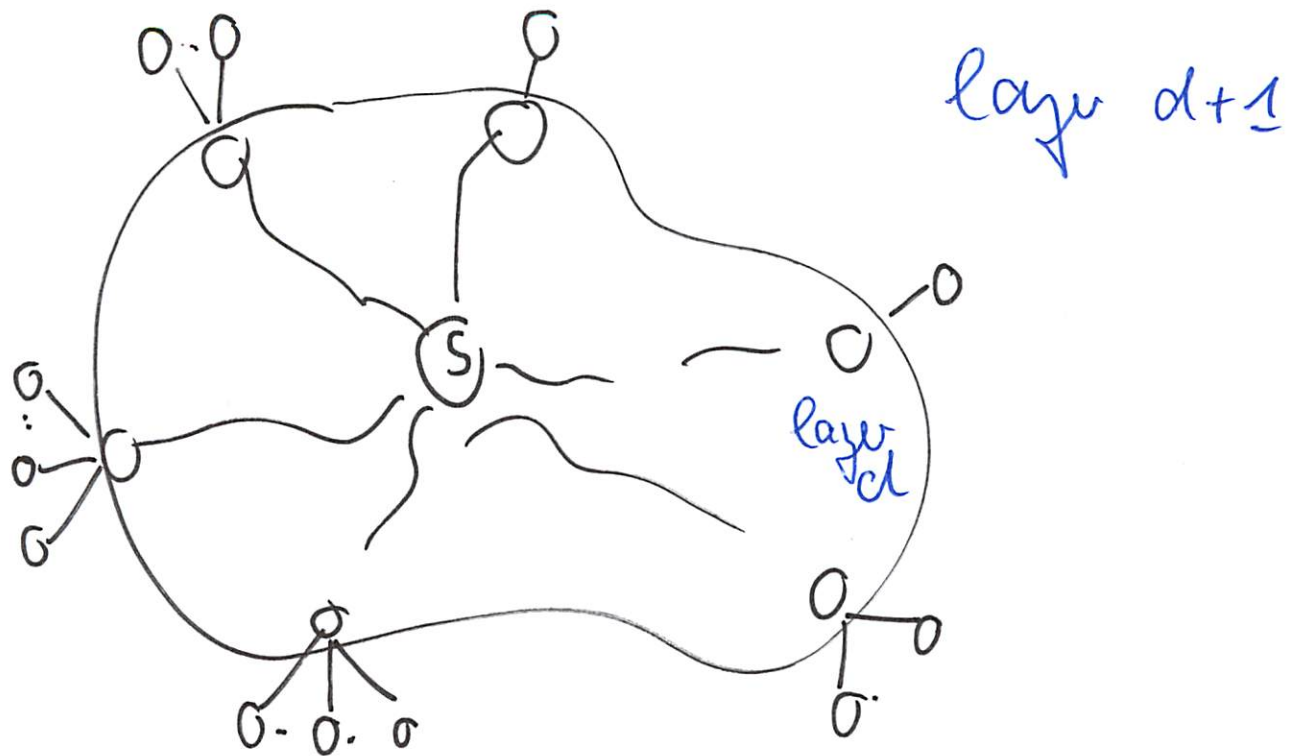
What are the layer 2 vertices?

All those "new" vertices that can be reached via an edge (just one not a path) from layer 1 vertices



We say that 2 vertices are adjacent if there is an edge connecting them.

in general, layer $d+1$ are all vertices that are adjacent to layer d vertices (but not already on layer $0, 1, \dots, d$)



Note: A vertex v is at distance d from s if and only if it is on layer d for vertex s .

So to compute the distance from s to a vertex t , we can determine the layer on which t appears.

Note: We get more information than just the distance $d(s, t)$.

Procedure bfs (G, s)

bfs : breadth first search

Input: Graph $G = (V, E)$ directed
or undirected; vertex $s \in V$

Output: For all vertices u
reachable from s , $\text{dist}(u)$
is set to the distance from s to u ,
i.e., $d(s, u)$.

for all $u \in V$
 $\text{dist}(u) = \infty$

$\text{dist}(s) = 0$

$Q = [s]$ (queue containing
just s)

while Q is not empty

$u = \text{eject}(Q)$

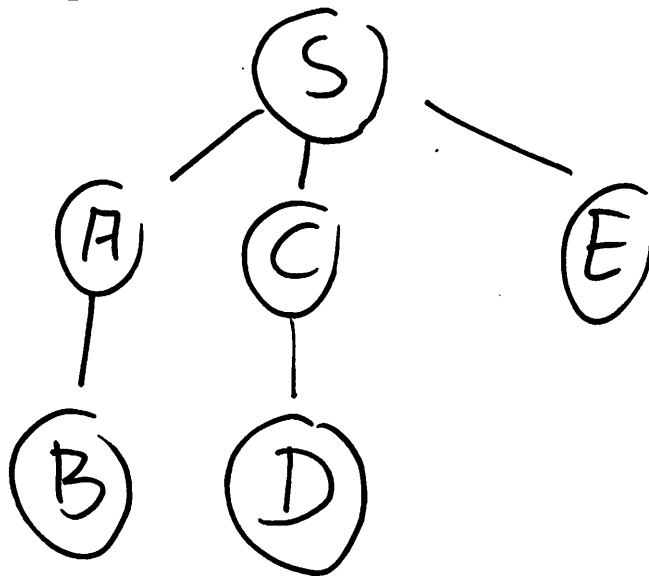
for all edges $(u, v) \in E$

if $\text{dist}(v) = \infty$

$\text{inject}(Q, v)$

$\text{dist}(v) = \text{dist}(u) + 1$ 4.13

Example



<u>Order of Visitation</u>	<u>Level dist()</u>	<u>Queue contents after processing vertex</u>
		[S]
S	0	[A C E]
A	1	[C E B]
C	1	[E B D]
E	1	[B D]
B	2	[D]
D	2	[]

Correctness

Inductive Hypothesis

for each $d=0,1,\dots$ 1) all nodes at distance $\leq d$ have correct distance value

2) all other nodes have distances ∞

3) the queue contains exactly the nodes at distance d (once all distance $d-1$ nodes have been processed)

The inductive Hypothesis is easily turned into a proof by induction.

Why does this prove the correctness?

Complexity!

The overall running time of the algorithm is $O(|E| + |V|)$ for a graph $G = (V, E)$.

Argument: Each vertex is added exactly once to the Queue (if it is reachable from s). \Rightarrow

There are $2|V|$ queue operations.

The inner loop executes for each edge a constant amount of operations whose cost is a constant.

We cannot improve in terms of "big oh".

dfs Depth-First Search and
bfs Breadth-First Search
are 2 alternate ways in
which to explore graphs.

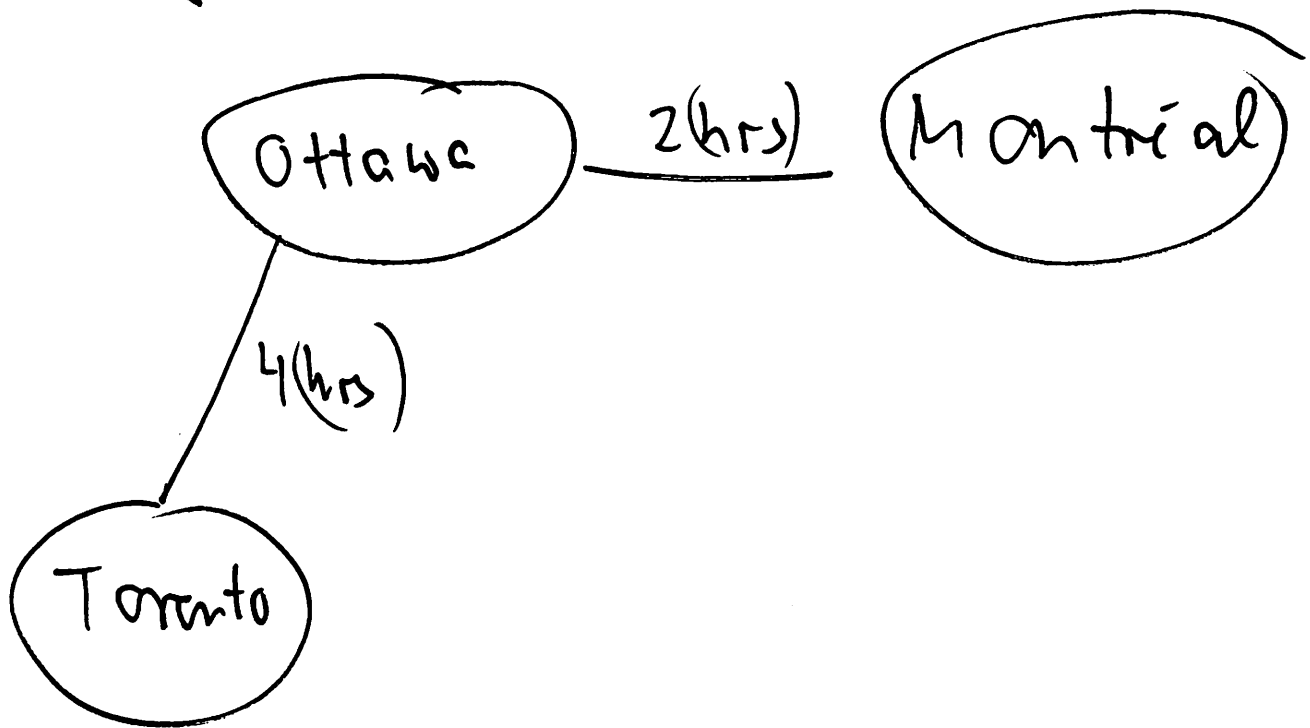
Find examples for each method.

e.g.; Chess is suited for
which strategy!

Lengths / weights on edges

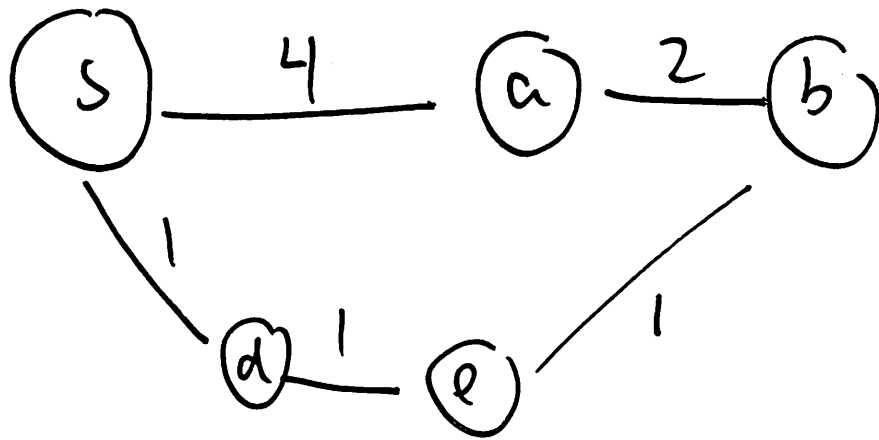
bfs considers all edges to have the same length or weight

if you drive though you have different lengths



Now the distance $d(s, u)$ is still the min cost of all path from s to u .

But the cost of a path = sum of the ^{weights} lengths of each edge on the path.



$$\text{cost}(s-a-b) = 6$$

$$\text{cost}(s-d-e-b) = 3$$

distance $d(s, b) = 3$

number of edges | $s-d-e-b$ | = 3 | $s-a-b$ | = 2 4.15

How to find the distance
in this setting!

Dijkstra's algorithm (G, l, s)

Input: Graph $G = (V, E)$; $s \in V$
positive edge length $\{l_e : e \in E\}$

Output: for all vertices u reachable from s ;
distance, $\text{dist}(u)$, is $d(s, u)$

for all $u \in V$
 $\text{dist}(u) = \infty$
 $\text{prev}(u) = \text{nil}$

$\text{dist}(s) = 0$

$H = \text{make_queue}(V)$ (using dist-values as keys)

while H is not empty

$u = \text{deletemin}(H)$

for all edges $(u, v) \in E$

if $\text{dist}(v) > \text{dist}(u) + l(u, v)$

$\text{dist}(v) = \text{dist}(u) + l(u, v)$

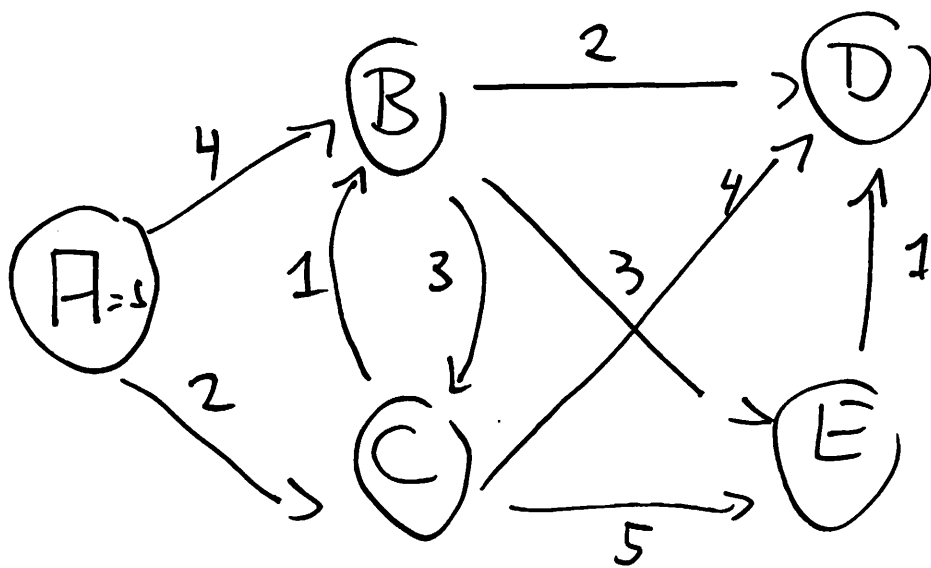
$\text{prev}(v) = u$

$\text{decreasekey}(H, v)$

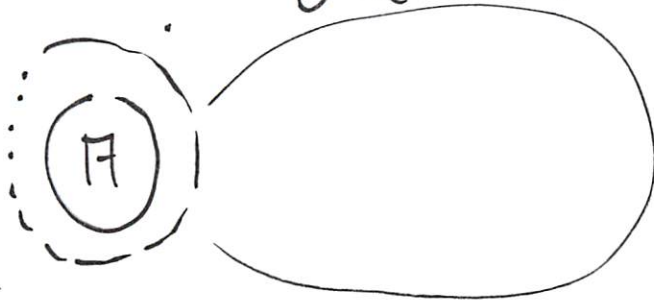
$\text{dist}(u) = \infty$ means no value found
for u yet

When u is removed from H the
value of $\text{dist}(u)$ is $d(s, u)$
and $\text{prev}(u)$ is the node on the
shortest path from s to u
just before u .

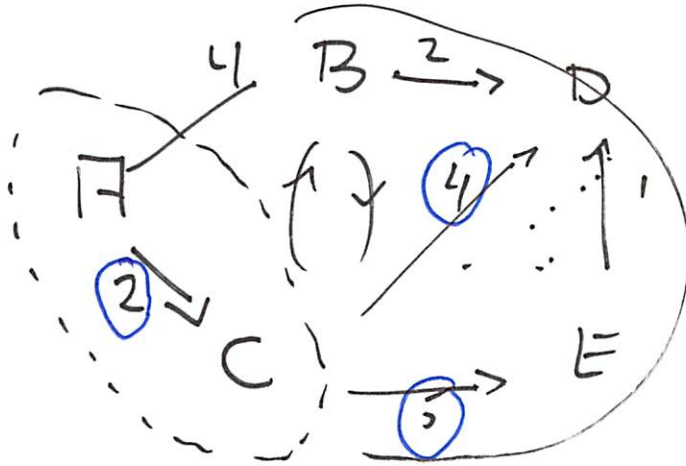
Example: (see book)



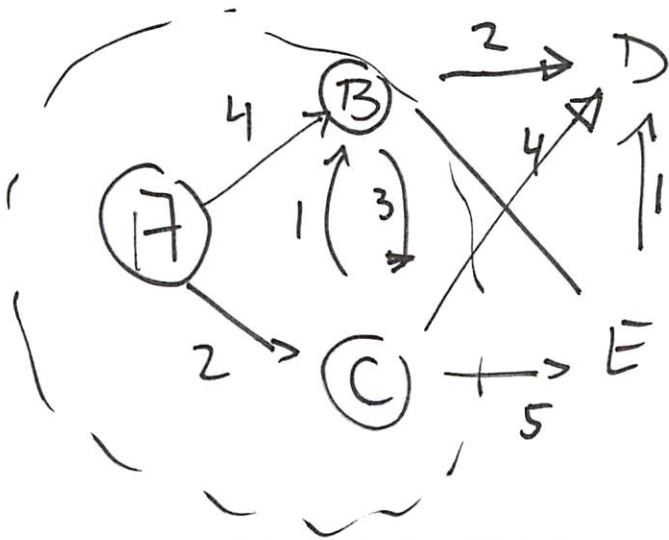
see textbook



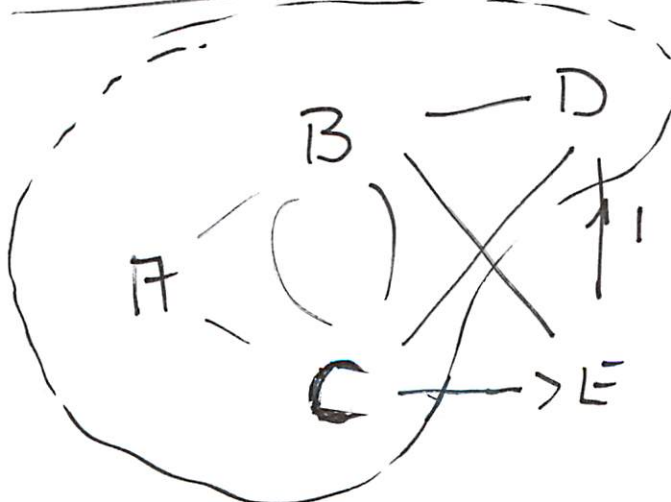
A: 0	D: ∞
B: 4	E: ∞
C: 2	



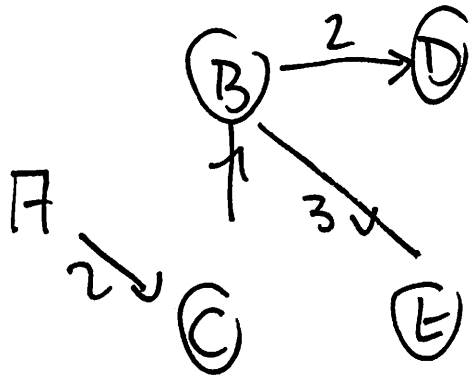
A: 0	D: 6
B: 3	E: 7
C: 2	



A: 0	D: 5
B: 3	E: 6
C: 2	



A: 0	D: 5
B: 3	E: 6
C: 2	



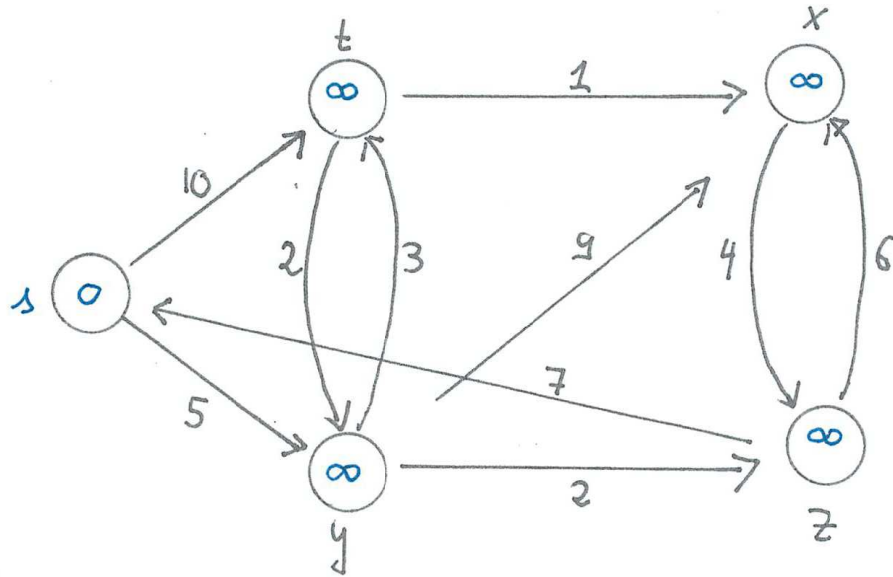
using the pointers



① We have seen an example (pp. 4.2-4.7) of a graph with an exponential, 2^k , $k = \theta(n)$ paths. This example was chosen so that it demonstrates that there can be an exponential number of shortest paths.

Think: how can Dijkstra's algorithm run in $O(|E| + |V| \log |V|)$ time if there can be an exponential number of shortest paths?

② If we did not want shortest paths, but (simple) paths. Is there a very simple graph with many many paths? What is the graph(s) and how many paths are there?



Initialization

all distances (except from s to s) are ∞

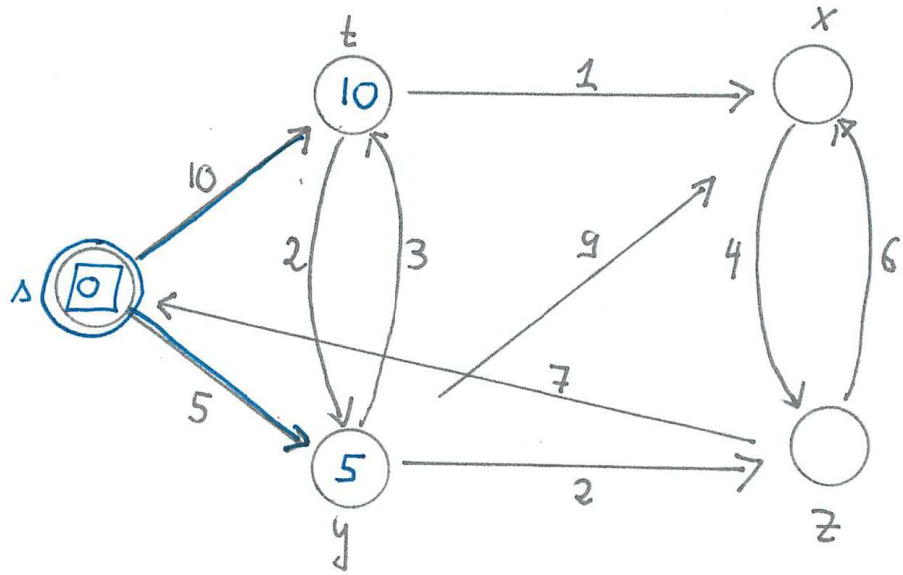
$dist(u) = \infty$ for all $u \neq s$

$dist(s) = 0$

$prev(u) = nil$ for all $u \neq s$

u = s

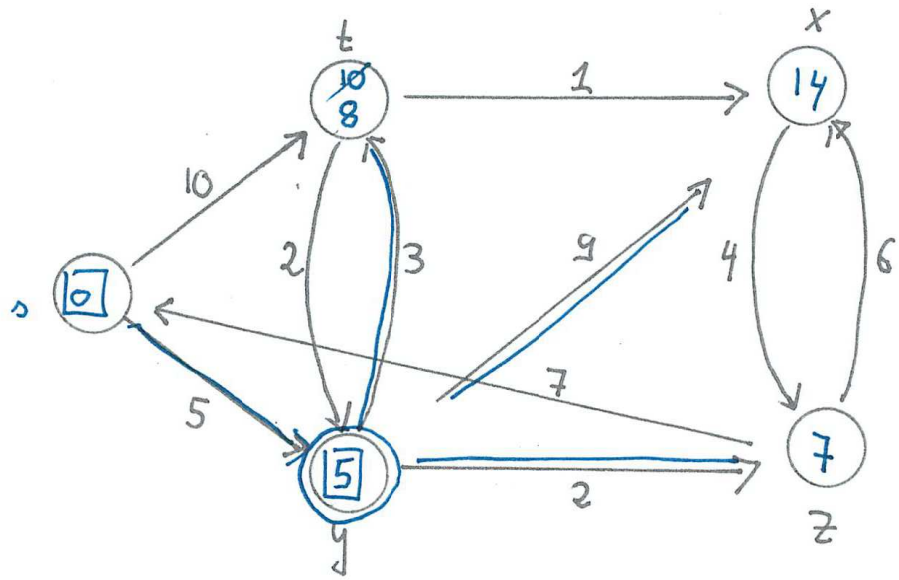
<u>dist()</u>	<u>Prev</u>
t	s
x	nil
y	s
z	nil



$d(s) = 0$ has been found

$$u = y$$

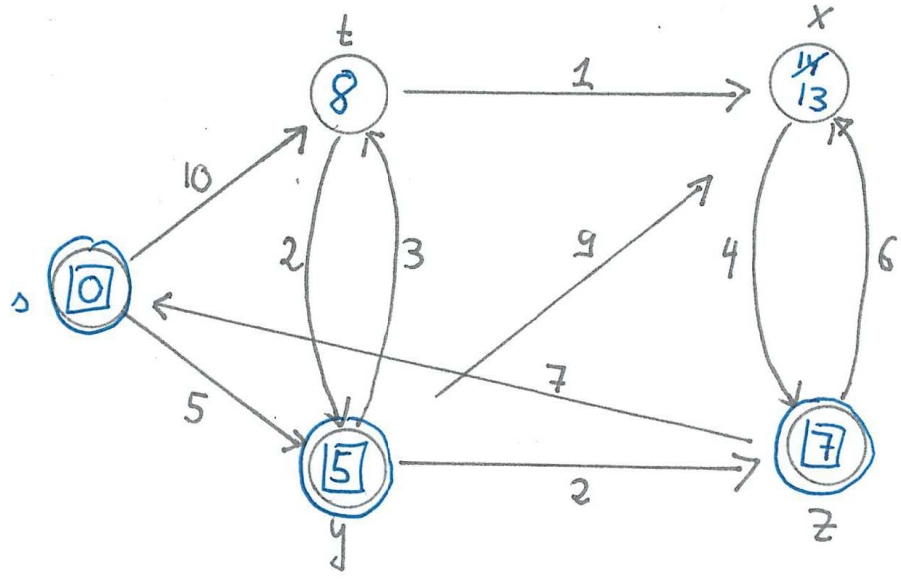
	dist()	prev
t	8	y
x	14	y
z	7	y



$d(y) = 5$ has been found

u = z

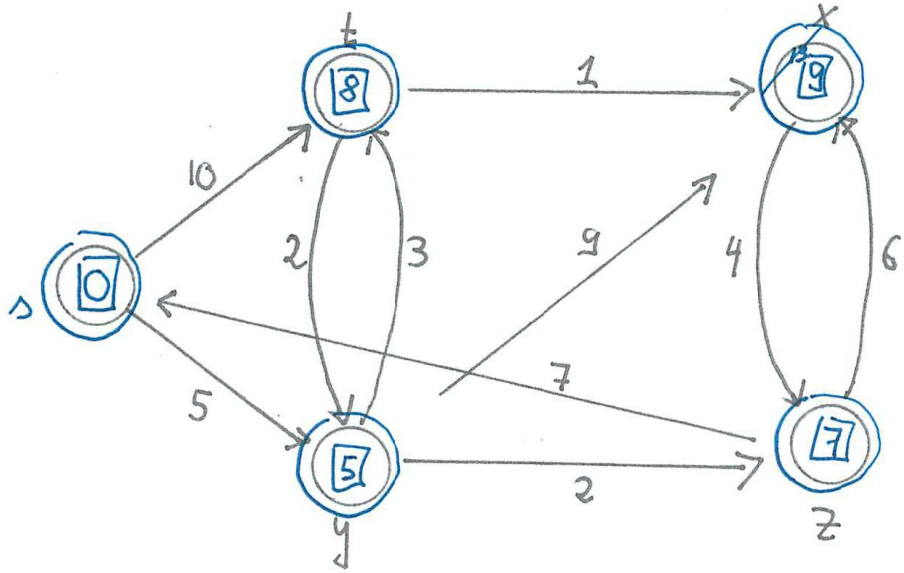
	dist()	prev
t	8	y
x	13	z



$d(z) = 7$

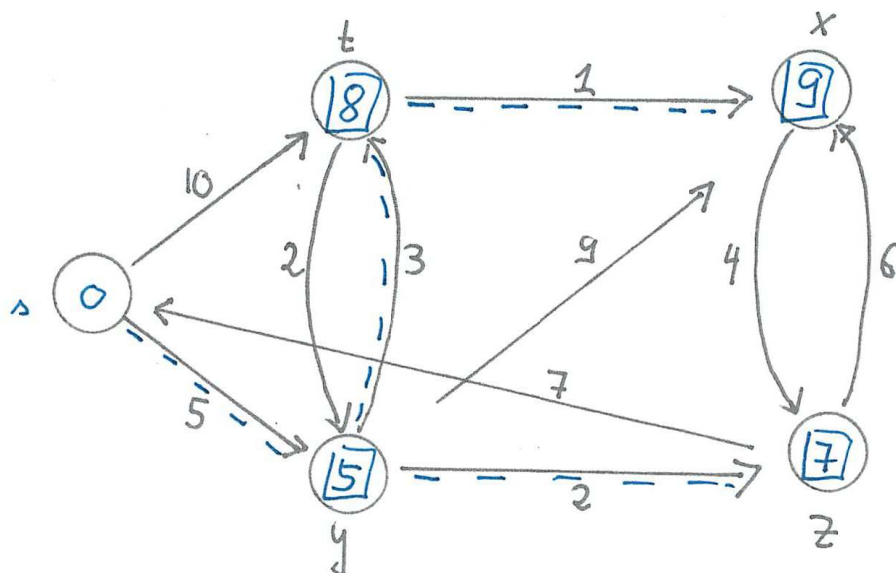
$U = t$

x	dist	prev
	9	t



$d(t) = 8$

$$U = X$$



- - - : previous relationship

$$d(x) = 9$$

Theorem Given a weighted directed Graph $G = (V, E)$ with non-negative weights on each edge of G , source s ; the algorithm, Dijkstra, terminates with $d(u) =$ distance from s to u for all $u \in V$.

Correctness

Loop invariant: at the start

of each iteration of the while loop for each vertex $v \in V - H ::= S$ (i.e., all vertices no longer in the queue) the correct distance from s to v has been found.

Initialization:

initially all vertices are in H , thus $S = V - H = \emptyset$. The statement is therefore trivially true.

Termination:

$H = \emptyset$, thus if we can show the invariant to hold. The algorithm is correct.

Maintenance:

We want to show that when u , the vertex with minimum key value is added, the invariant remains true.

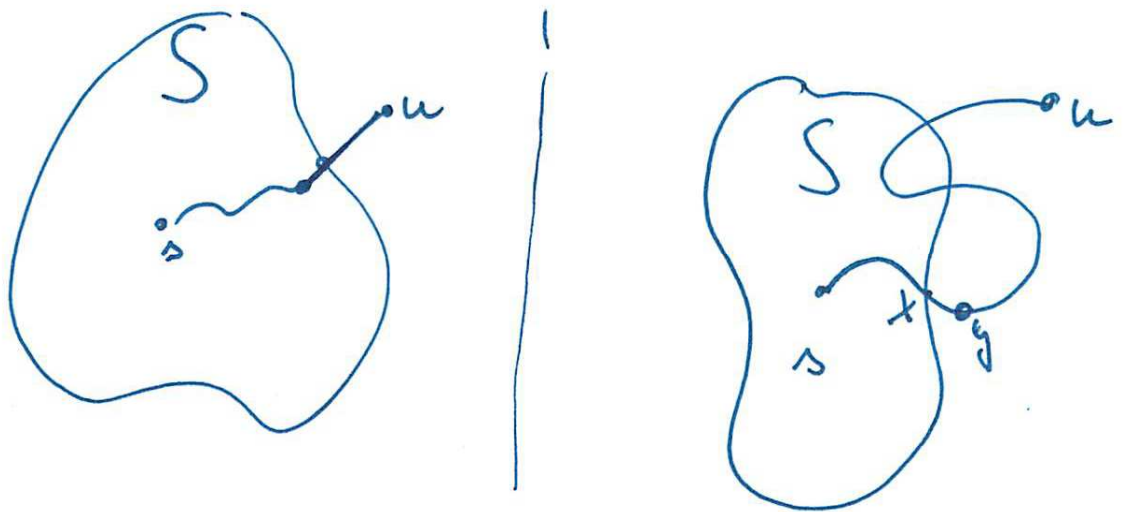
Proof by contradiction.

Assume that u is the first vertex for which $d(u)$ is incorrectly computed.

Consider the moment u is taken of the queue. $u \neq s$ as $d(s) = 0$ is correctly determined.

So, $S \neq \emptyset$, when u is added.

There is a path from s to u .
Which means there is also a shortest path from s to u .



Let y be the first vertex on that path outside S and x the predecessor of y .

(Note the part outside S before u could be empty.)
Also $x = s$ is possible

By loop invariant, $d(x)$ is correct. The edge (x, y) was used to give an upper bound on $d(y)$, i.e. $\text{dist}(y) = d(x) + \text{cost}(x, y)$
($\text{cost}(x, y) = l(x, y)$)

$$d(y) = d(s, y) \leq d(s, u) \leq \text{dist}(u)$$

But, both u and y were in H when u was chosen

$$\text{dist}(u) \leq \text{dist}(y)$$

$$\text{thus } \text{dist}(y) = \text{dist}(u)$$

$$\text{and } \text{dist}(y) = d(s, y) = d(s, u) = \text{dist}(u)$$

This is a contradiction to our assumption.

\Rightarrow the loop invariant is correct.

We can also show the correctness of the predecessor trivially.

Corollary: The algorithm also computes the shortest path tree with roots by using the predecessor relation.

Analysis:

The important factor determining the run-time, is the time for executing the priority queue operations.

- o insert well we create the entire queue
- o ~~extract~~ delete min
- o decrease key

o There are $|V|$ delete min operations

o we create the entire queue

o decrease key is often done by removing the element and then inserting it.

There are $|V| + |E|$ such operations

If we use a binary (min) heap for n elements

create takes $O(n)$

insert " $O(\log n)$

delete min " $O(\log n)$

Thus, Dijkstra's algorithm runs in $O((|V| + |E|) \log |V|)$ time using a binary heap.

Using a more complex priority queue organization known as Fibonacci heap, the time is reduced to

$$O(|E| + |V| \log |V|)$$

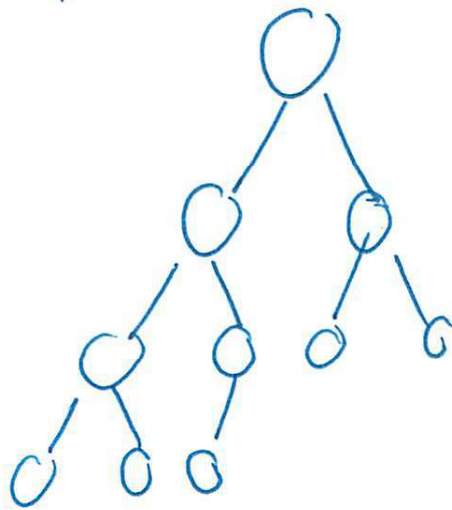
[insert + decrease key are $O(1)$ amortised]

What is the time complexity of Dijkstra's algorithm using an array implementation for the PQ.

Heaps (binary)

Structure

- ① Elements are stored in a complete binary tree, i.e. each level is filled from left to right and must be full before the next level is started. Except for the last level all levels are thus full. The last level may not be



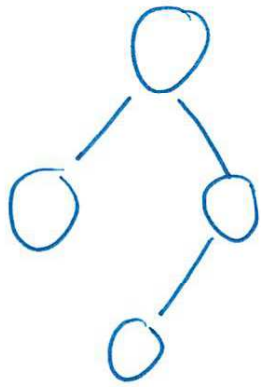
$n=10$

②

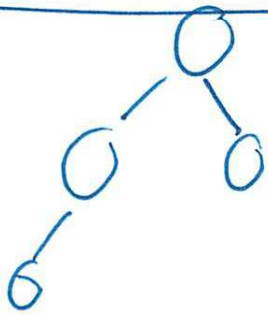
Order

The key value of a parent is \leq that of its children.

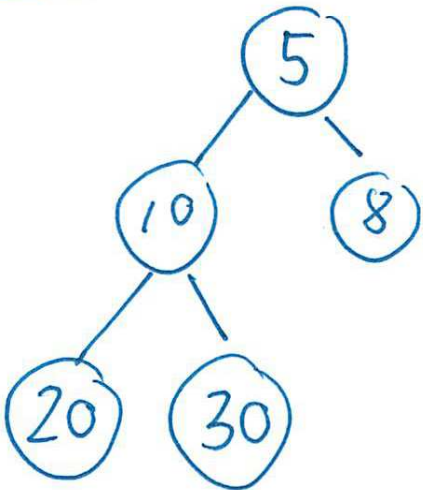
Ex:



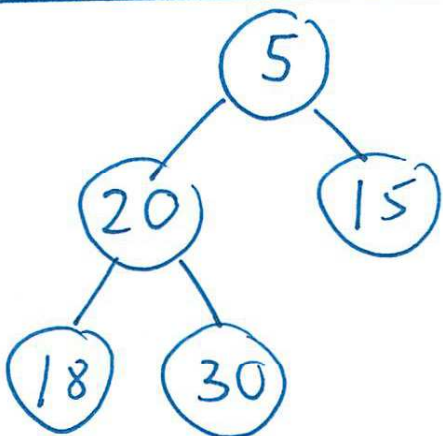
- not a heap
violate the
structure



✓ correct structure



✓ correct structure
✓ correct order



✓ correct structure
- in correct order
 $20 > 18$

Fact. The global minimum is at the root.

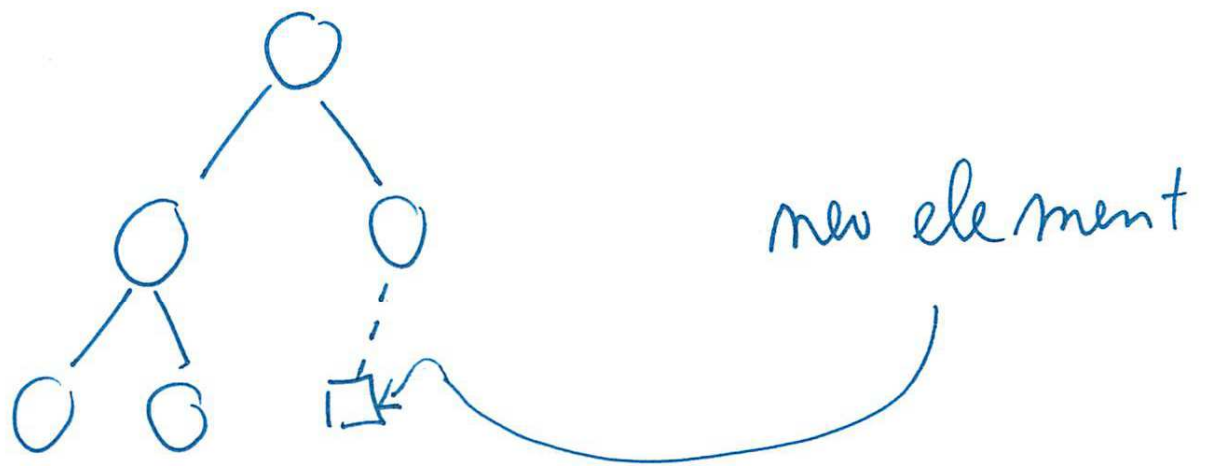
Question. Where is the maximum?

How many comparisons are required to find the maximum in a min-ordered heap?

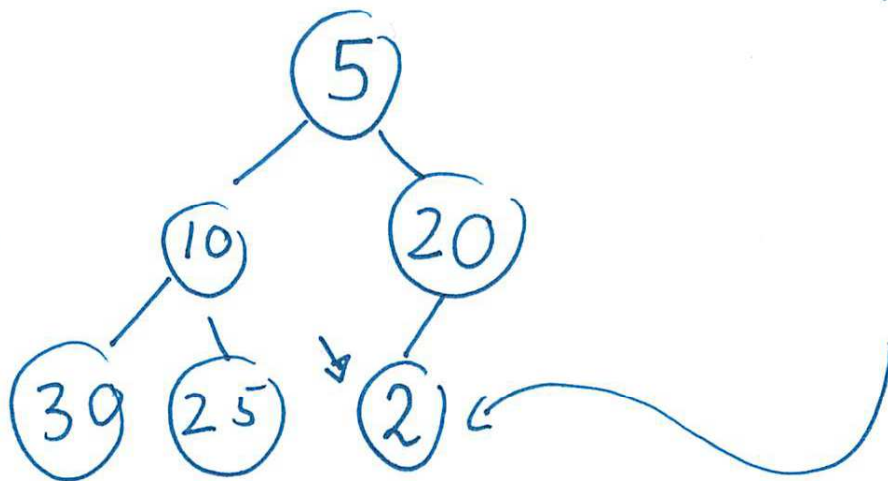
see
Class

Insertion :

a new element is inserted into a heap by first getting the structure right then the order.



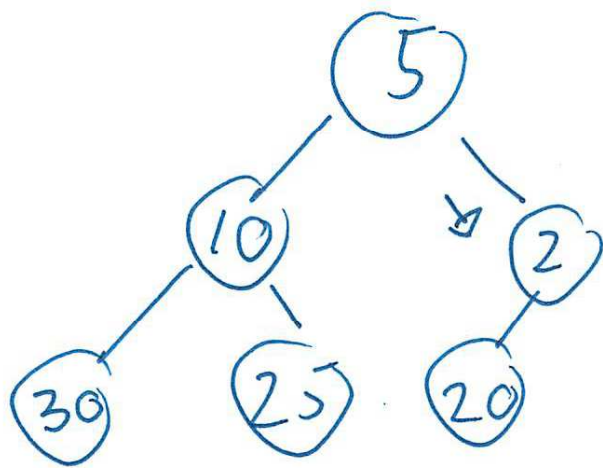
Then, to get the order right we bubble the element up.



heap order is violated.

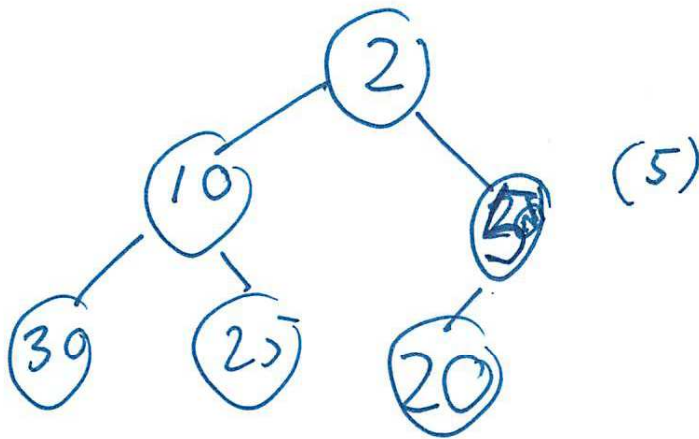
⇒ compare element to parent
if smaller swap
and move one level up

repeat



after swap

again, parent (5) > (2) \Rightarrow swap



Why is it correct?

For the heap-order to hold every
parent node key \leq child node key

all nodes not on the path
from the new leaf to the root
remain unchanged.

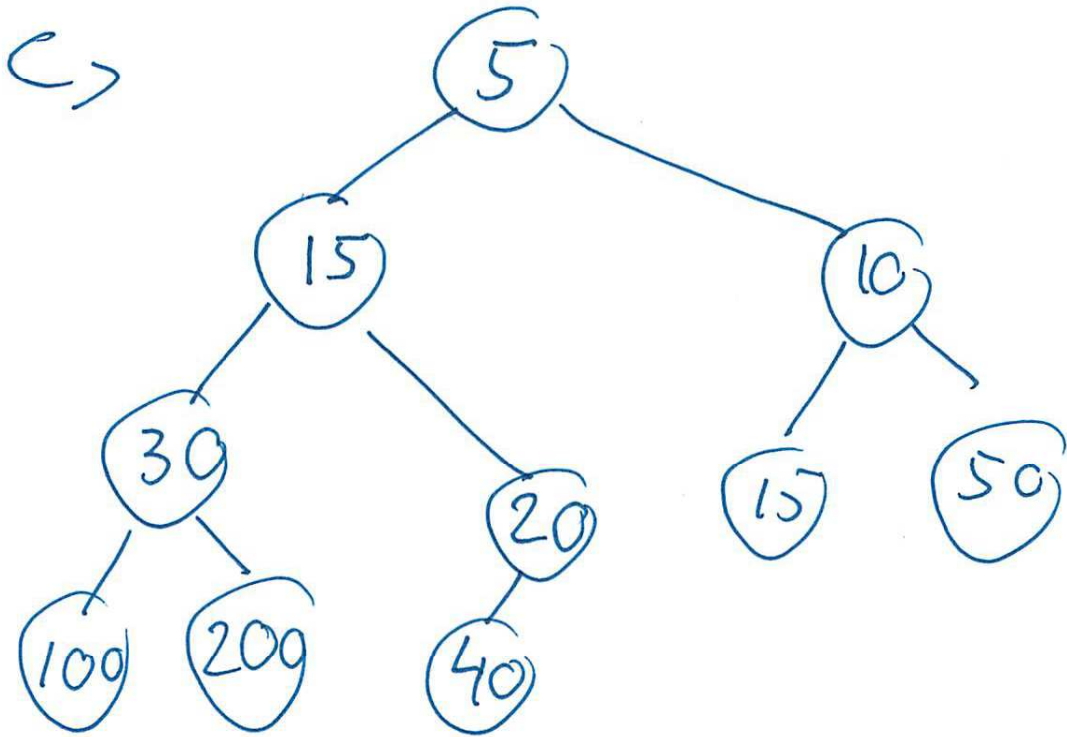
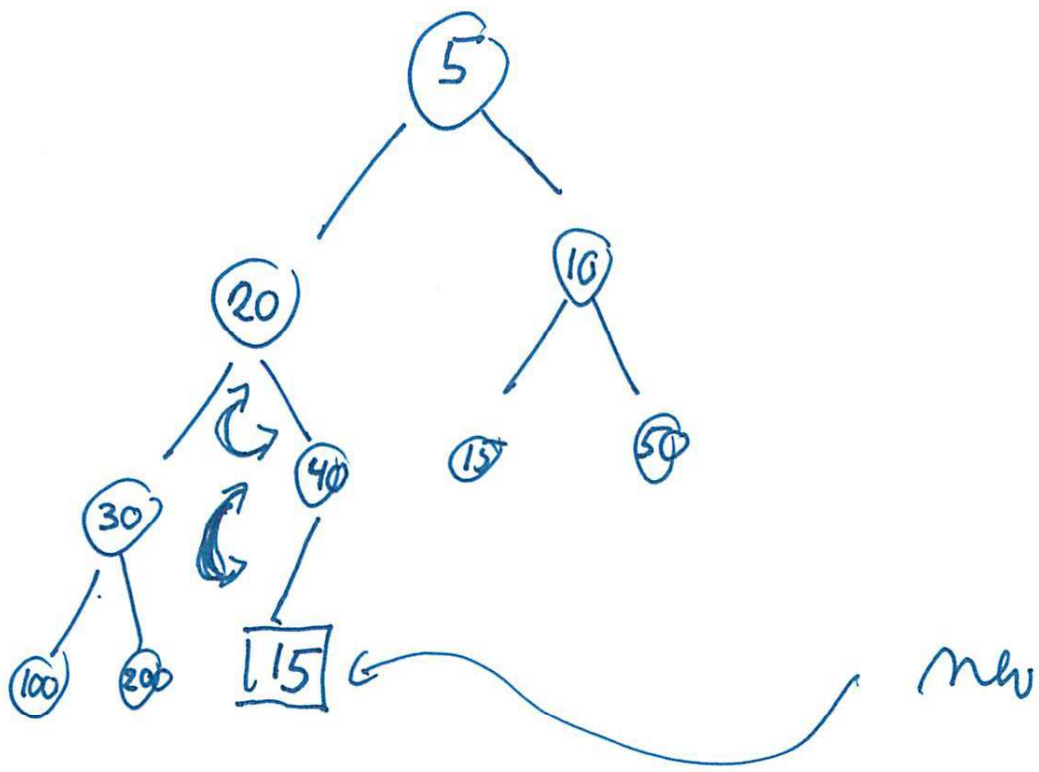
The new leaf is after possible
swap \geq parent key.

—
Every time an element pair
is swapped it results in a
parent key that is $<$ the original
so the parent is replaced by a
smaller key. The heap-order
to the other child is thus correct.

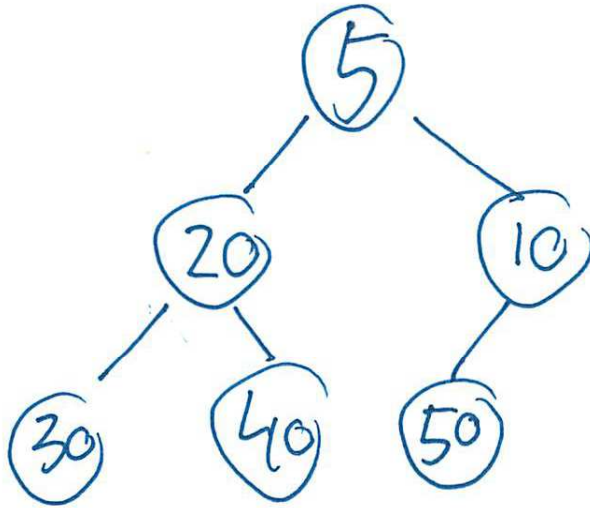
If a parent, child relation is
correct on the path no further
swaps are required

Time. $O(\log n)$: path
length

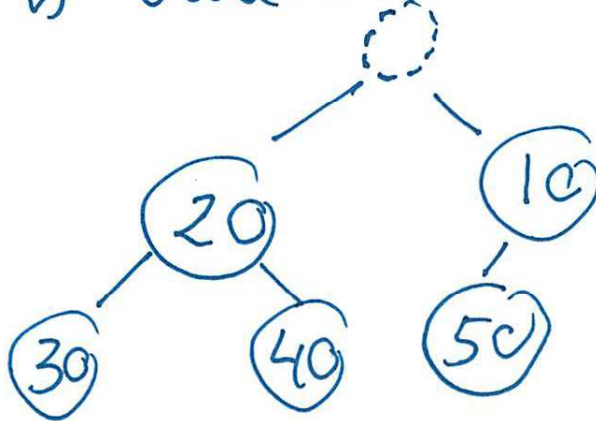
D.19



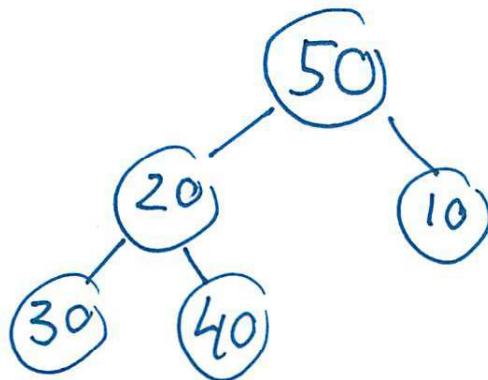
Delete Min



5 is deleted



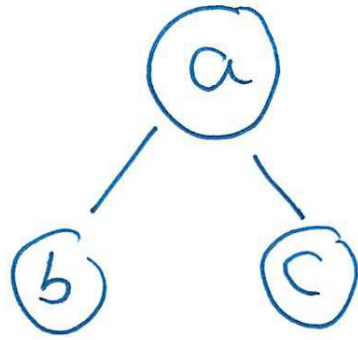
To get the structure right move 50 into root position.



now get
the heap
order fixed

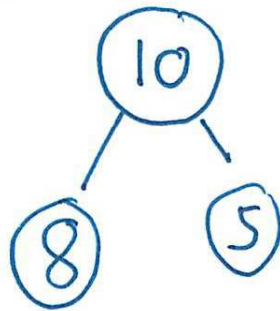
$$50 > 20 \\ > 10$$

if $a > \min(b, c)$

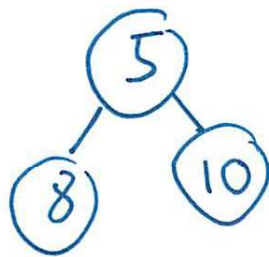


replace
a by
 $\min(b, c)$

Example :



↪

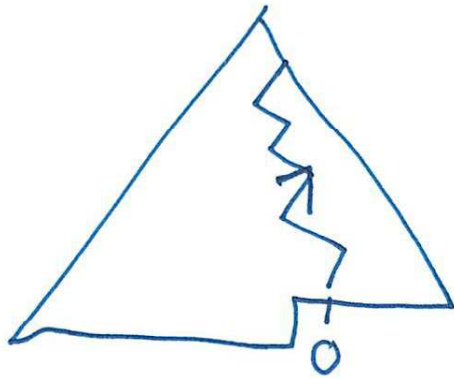


recurse to the side of the swapped element. Here to the right child.

constant work per level on a path to a leaf.

Notes: For a given n the structure of the heap is unique.

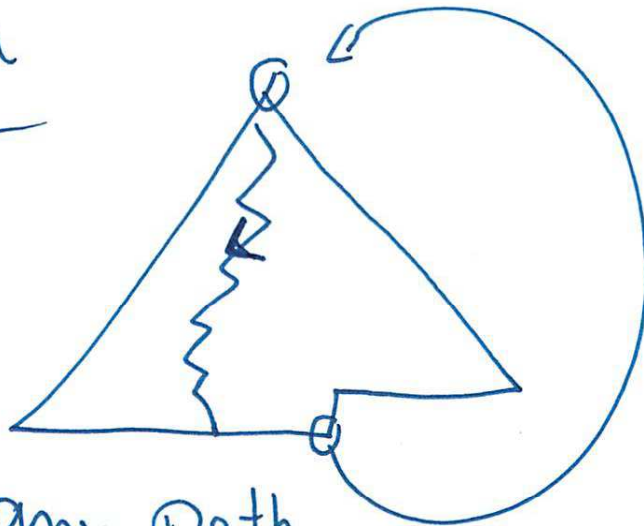
Insertion



The path is unique

"bubble-up"

Delete



"trickle down"

any path is possible

How to create a heap on n elements!

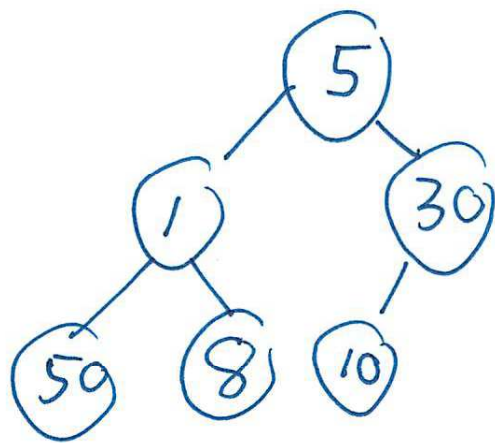
I. $n \times$ insert element

$$\Rightarrow O(n \log n)$$

II. an $O(n)$ method

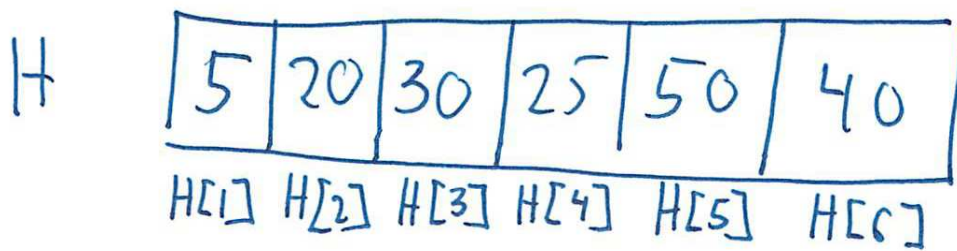
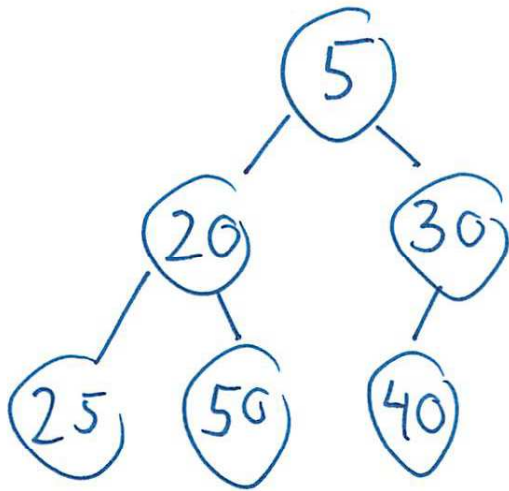
Take the n elements and put them into a heap-structured tree (possibly violating the order)

5, 1, 30, 50, 8, 10

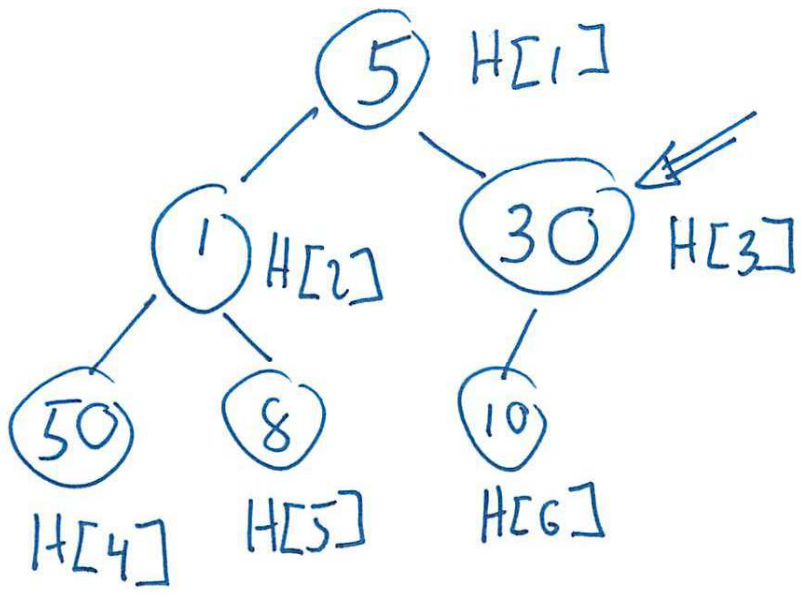


Now 10, 8, and 50 have no children so their heap-order is trivially correct.

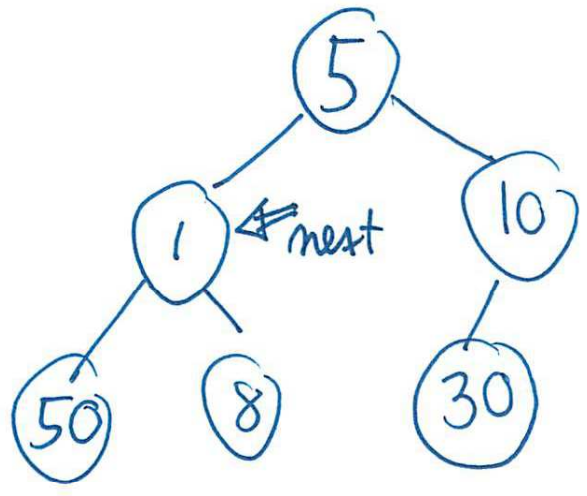
heaps are most efficiently stored in an array.



parent $H[i]$ has children
 $H[2i]$, $H[2i+1]$
(if it exists)

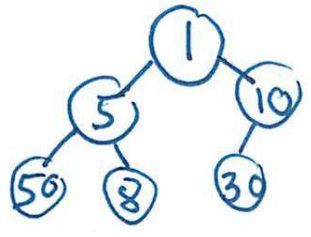


$H[3]$ is rightmost non leaf in array.
 Check if heap order is correct for $H[3]$
 If not fix (like in delete min operation)



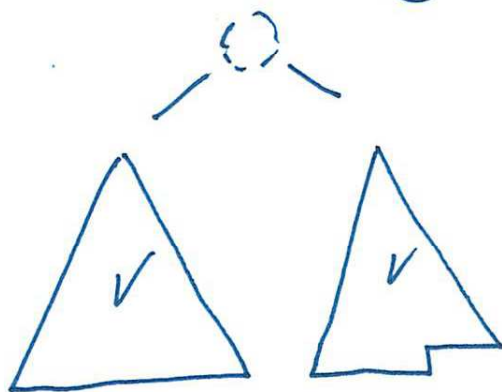
$H[2]$ is correct

 The $H[1]$ fix it.



Correctness

recursive argument



possibly incorrect

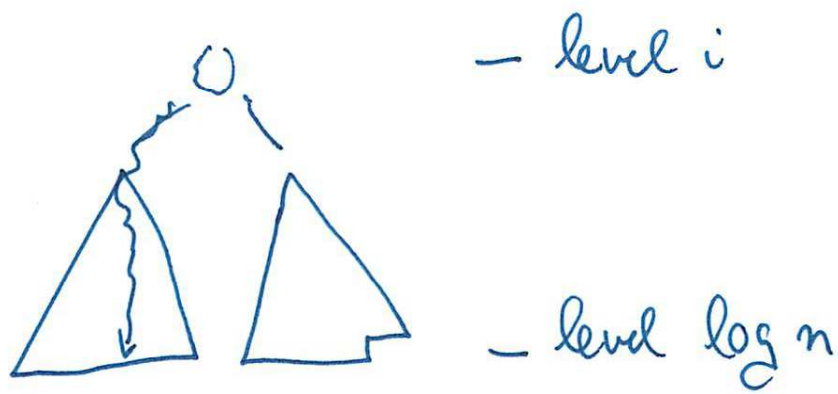
↳ gets heap-ordered.

Analysis

intuition: at the low levels (bottom)

we have many elements, but little work per element (height is small)

at the top levels (near root) we have few elements with a lot of work.



there are 2^i elements on level i
 (except for possibly the last level)

each element may need to go
 to the bottom (trickle-down)

work $O(\log n - i)$

$$\sum_{i=0}^{\log n} 2^i \cdot (\log n - i) = \sum_{i=0}^{\log n} 2^{\log n - i} \cdot i$$

$$= \sum_{i=0}^{\log n} \frac{2^{\log n}}{2^i} \cdot i = \sum_{i=0}^{\log n} \frac{n}{2^i} \cdot i$$

$$= n \sum_{i=0}^{\log n} \frac{i}{2^i} \leq 2n$$

\Rightarrow Heap construction takes
 $O(n)$ time.

Bellman-Ford Alg.

Bellman-Ford (G, w, s) better description than in textbook

Input: Graph G , weight function w , source s
 $G = (V, E)$, \uparrow also known as "E"

output: If there is a negative cycle reachable from s the algorithm returns "false" i.e. no solution exist
Otherwise, the algorithm produces the shortest path distances from s to all reachable vertices

```
1 Initialize-Single-Source ( $G, s$ )
2 for  $i = 1$  to  $|V| - 1$  do
3   for each edge  $(u, v) \in E$ 
4     do Relax ( $u, v, w$ )
5   for each edge  $(u, v) \in E$ 
6     do if  $d[v] > d[u] + w(u, v)$ 
7       then return "false"
8   return "true"
```

with

Ⓐ Initialize-Single-Source (G, s)

for each vertex $v \in V$

do $d[v] = \infty$

$\pi[v] = nil$

$d[s] = 0$

Ⓑ Relax (u, v, w)

if $d[v] > d[u] + w(u, v)$

then $d[v] = d[u] + w(u, v)$

$\pi[v] = u$

In many textbooks the Relax procedure is used.

The complexity of Bellman-Ford's algorithm is $O(|V| \cdot |E|)$.

dominates $\left[\begin{array}{l} \text{for each vertex do} \\ \quad \text{for each edge do} \\ \quad \quad [\text{constant work}] \end{array} \right]$

Correctness not covered here
but easy.

A path has at most $|V| - 1$ vertices
unless it has cycles.

So, the lines 5-7 test if
after executing the loop (2)
one can still gain \Rightarrow cycle.

If Bellman-Ford's algorithm "Sees" no changes between 2 iterations of the main loop (2) then it can terminate.

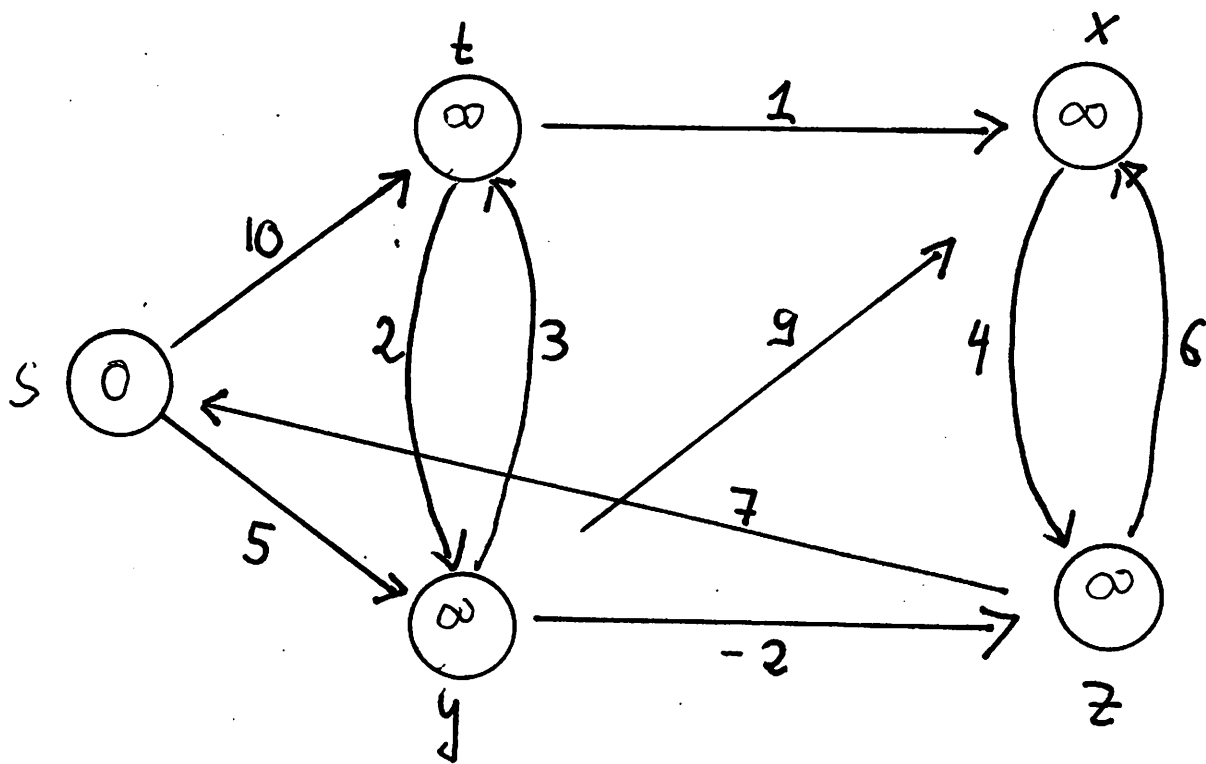
This is frequently, as easily implementable, improvement in actual run-time.

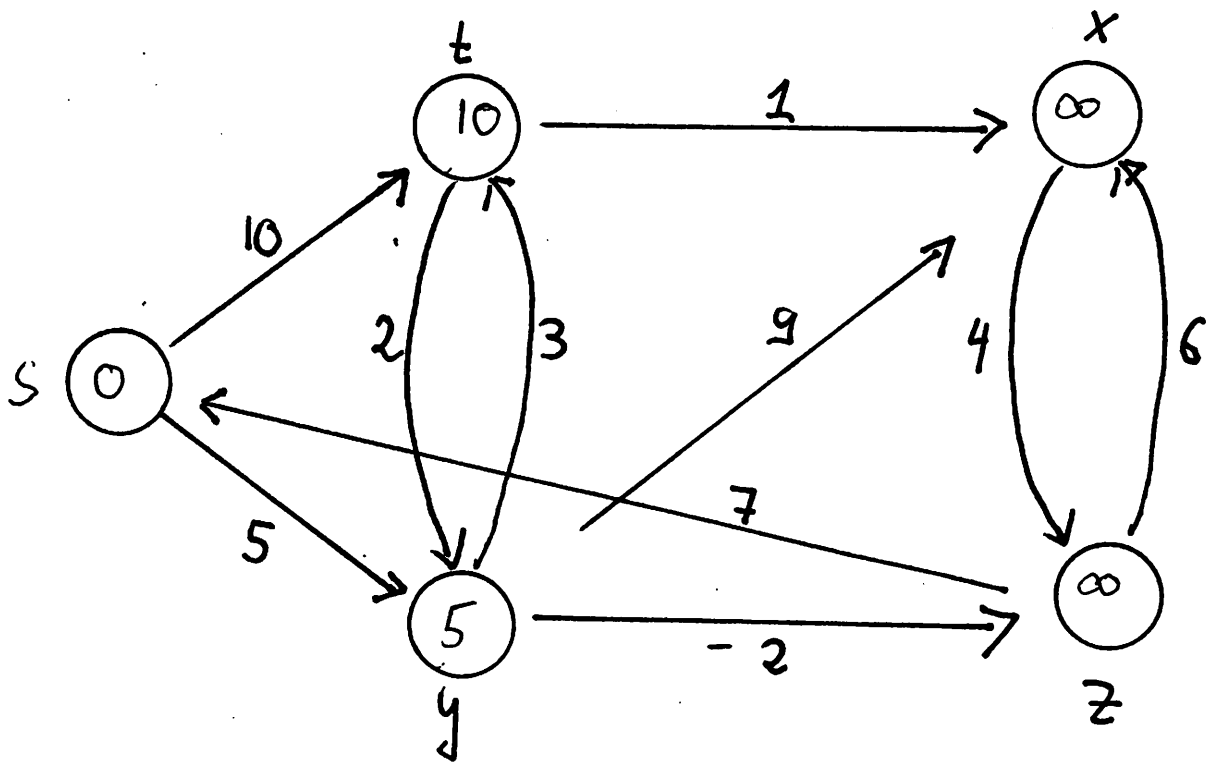
Construct for yourself 2 examples when this is a

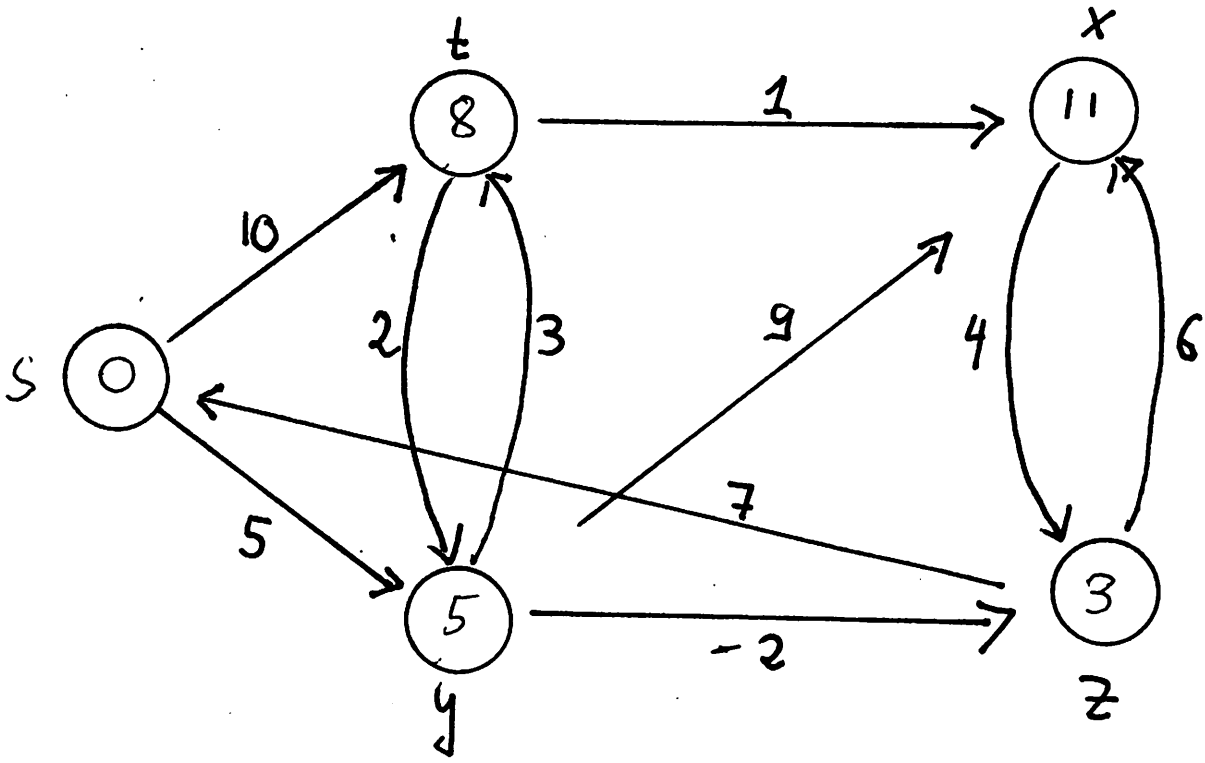
(a) major improvement and

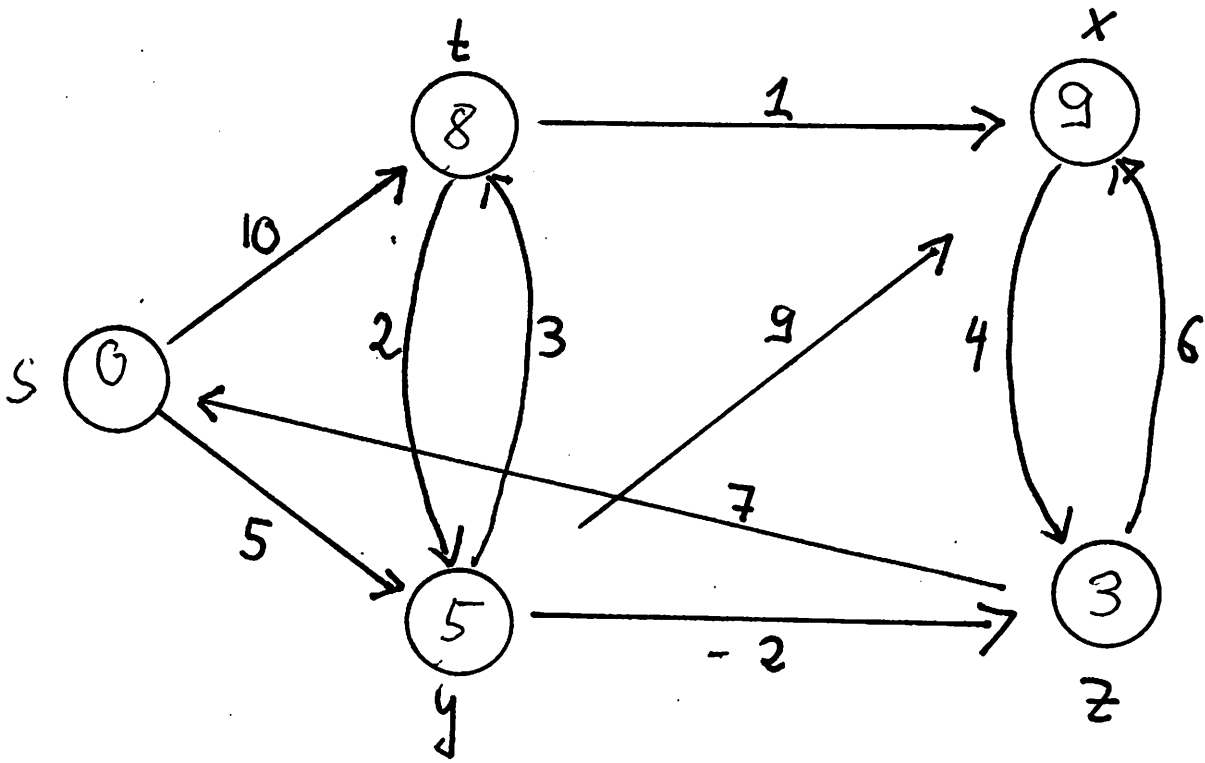
(b) no improvement.

Example









Chapter 5

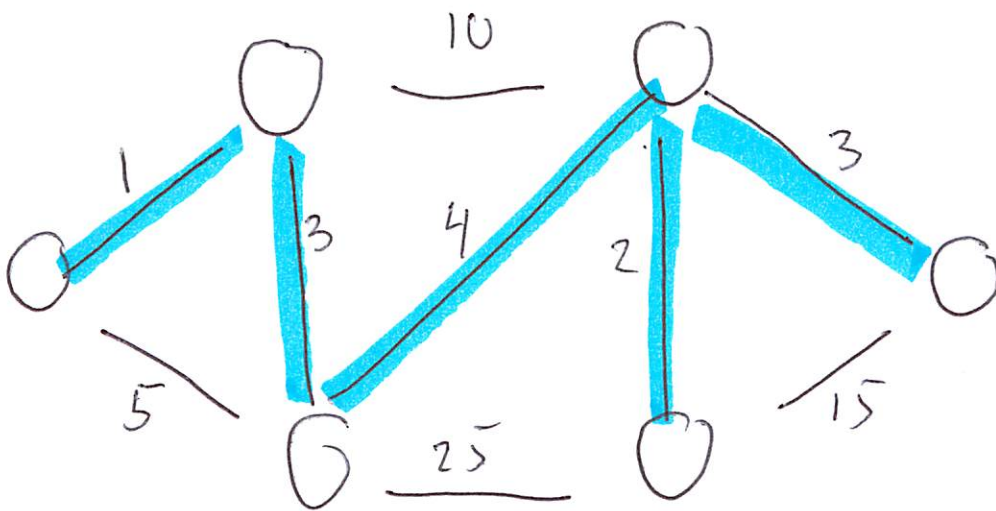
Greedy Algorithms

Informally, an algorithm is greedy if it builds a solution piece by piece, always choosing the next piece that offers the most obvious and immediate benefit.

Think about real-life examples when this greedy approach succeeds and where not.

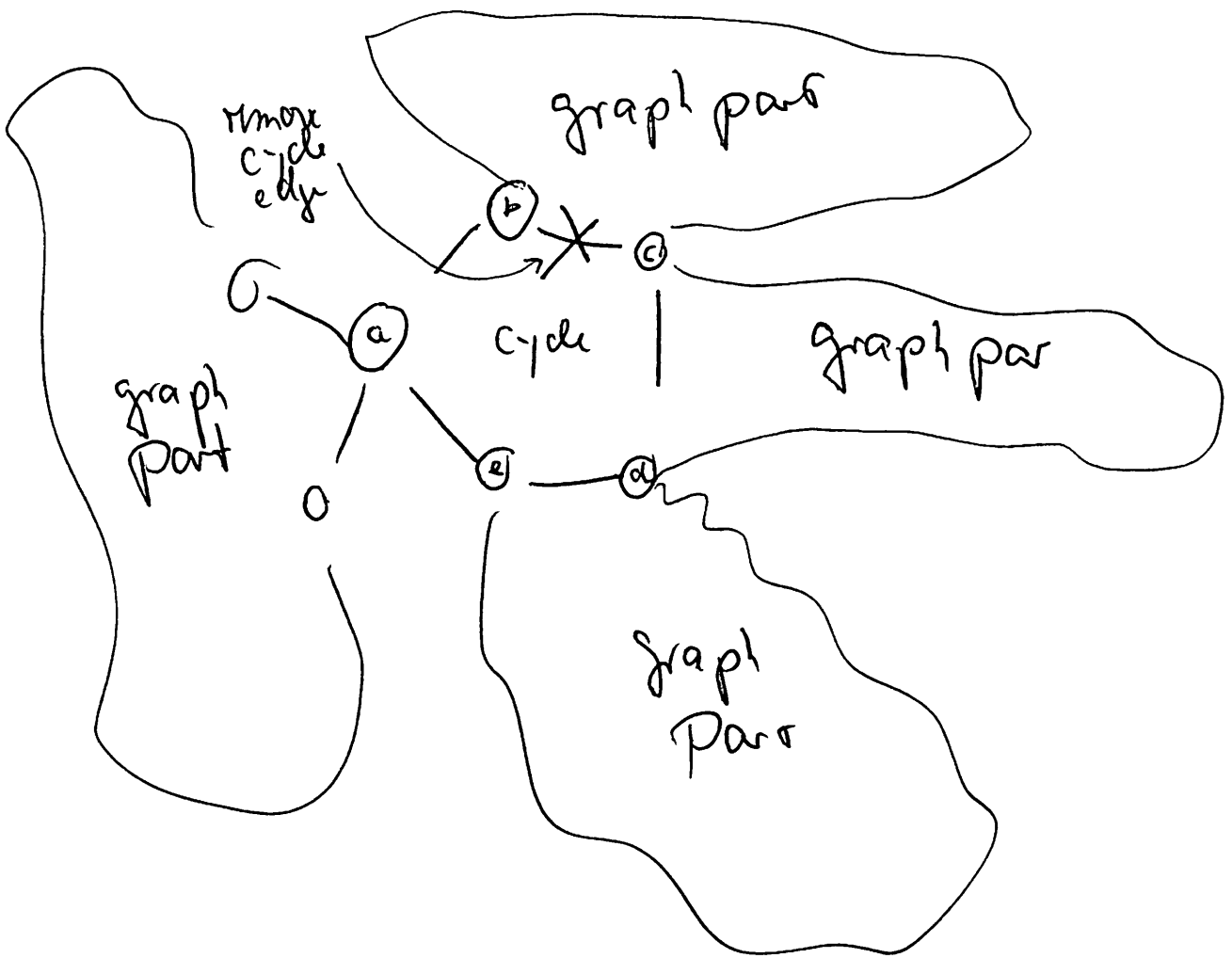
Minimum Spanning Trees

Example: Computer Network
with min amount of wire
between them.



First some facts / Properties

- ① Removing a cycle edge cannot disconnect a graph.



$a - b - c - d - e$ is a cycle.

if we remove say, $b - c$, and

there is a path

x, \dots, b, c, \dots, y

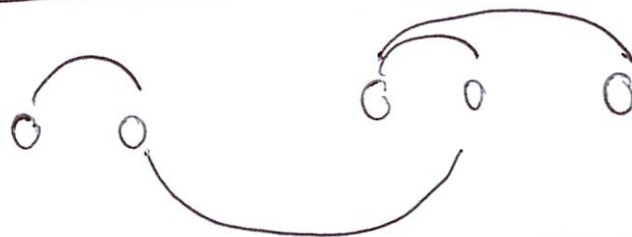
we can always replace this path by

$x, \dots, b, a, e, d, e, \dots, y$

when the parts \dots stay.

Properties cont'd

② A tree on n nodes has $n-1$ edges.
Build the tree adding one edge at a time.



each time, we add an edge 2 components that were disconnected are connected.

after $n-1$ edges all initially n isolated nodes are in one connected component

(Adding an edge inside a component creates a cycle.)



3 Also, the converse is true:

Any connected, undirected graph $G = (V, E)$ with $|E| = |V| - 1$ is a tree

Assume that G has a cycle.

Then remove one of the cycle edges.

After all cycles have been removed a graph $G' = (V, E')$, $E' \subseteq E$ is created that is acyclic.

Using Property 1 (repeatedly)

G' is connected $\Rightarrow |E'| = |V| - 1$
by Property 2. (if $E' = E$, G was

already acyclic.)

Properties

④ An undirected graph is a tree, if and only if there is a unique path between any pair of nodes.

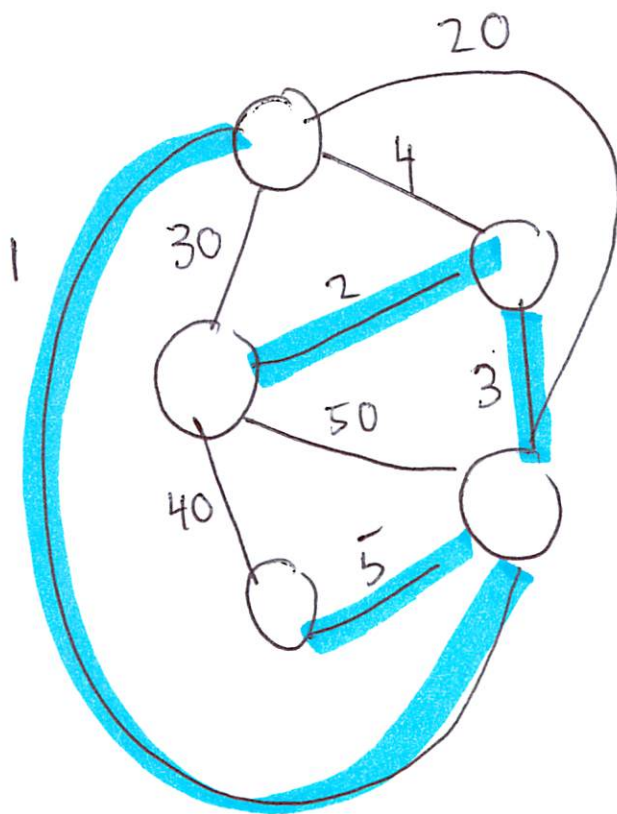
"
" \Rightarrow " If G is a tree there is a unique path as G is connected and if there were 2 paths a cycle would result.

"
" \Leftarrow " If there is a unique path between any pair of vertices, then G is connected and acyclic $\Rightarrow G$ is a tree.

Definition A minimum spanning tree for a graph $G = (V, E, w)$ is a tree $T = (V, E')$ $E' \subseteq E$ which minimizes

$$\text{Weight}(T) = \sum_{e \in E'} w_e$$

We call the minimum spanning tree MST



Minimum Spanning Tree

Kruskal's Algorithm

(Greedy approach)

Procedure Kruskal (G, w)

Input: A connected, undirected graph
 $G = (V, E)$ with edge weights w_e

Output: A minimum spanning tree for G

for all $u \in V$ do
 makeset (u)

$X = \{\}$

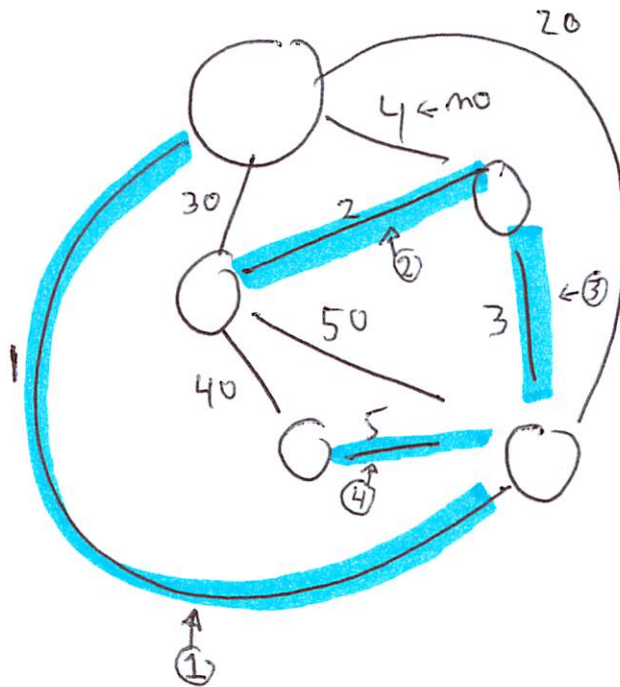
Sort the edges E by weight

for all edges $\{u, v\} \in E$, in increasing order (weight)
 do
 if find(u) \neq find(v)

 add edge $\{u, v\}$ to X
 union (u, v)

"Algorithm" for MST

repeatedly add the next lightest edge that does not produce a cycle.



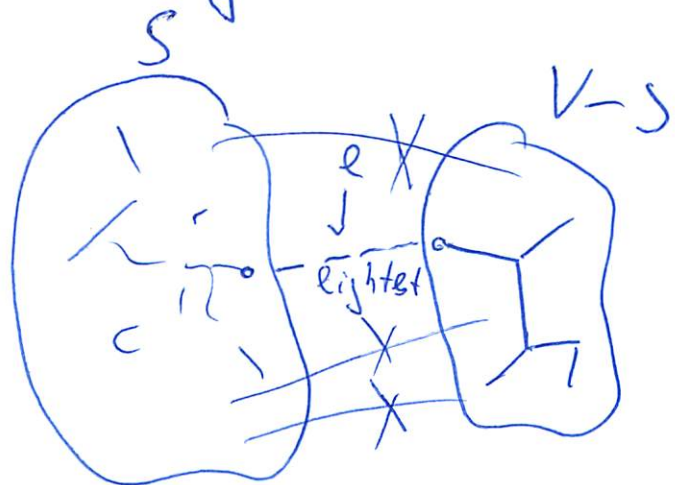
- (A) Does it work?
- (B) What is the time complexity?
- (C) Can we do better?

(B) depends on how fast we find the lightest edge
 (how fast we can test if we create a cycle which component does a vertex belong to) — easy!
 5.8

CUT Property (MST)

Minimum spanning tree

Suppose edges X are part of a minimum spanning tree of $G = (V, E)$. Pick any subset of nodes S for which X does not cross between S and $V-S$, and let e be the lightest edge across this partition. Then $X \cup \{e\}$ is part of some MST.



X : not in MST

lightest = minimum weight edge

So, we have 3 operations

make set (x): create singleton set for x

find (x): to which set does x belong?

union (x, y): merge the sets containing x and y

For Kruskal's algorithm:

$|V|$ make set operations

$2|E|$ find

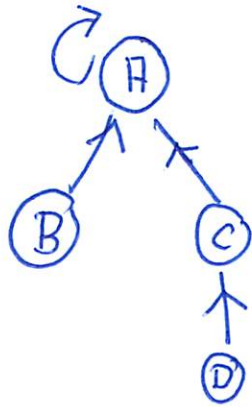
$|V| - 1$ union

so, how to implement these?

The operations are also used in other algorithms.

Disjoint Sets Problem

Store a set x in a tree,



↔ the root is the representative of the set
i.e. the name of the set

procedure makeset (x)

$$\pi(x) = x$$

$$\text{rank}(x) = 0$$

[height of subtree]

function find(x)

while $x \neq \pi(x)$ do $x = \pi(x)$

return x

(follows the up pointers to the root and return that root value)

Procedure union(x, y)

$$r_x = \text{find}(x)$$

$$r_y = \text{find}(y)$$

representative
finding

if $r_x = r_y$: return

are the equal!
sets are equal

if $\text{rank}(r_x) > \text{rank}(r_y)$
then $\pi(r_y) = r_x$

link the
smaller tree
as child of
larger

else

$$\pi(r_x) = r_y$$

if $\text{rank}(r_x) = \text{rank}(r_y)$ then

$$\text{rank}(r_y) = \text{rank}(r_y) + 1$$

rank (height)
goes up when
the trees have
equal heights

Example of sequence of
operations in class

also question:

What would happen
if we were to hang the
larger tree under the
smaller i.e.

reverse the inequality?
in procedure union?

Property 1 :

For any x , $\text{rank}(x) < \text{rank}(\pi(x))$

Proof obvious

Property 2: Any root node

of rank k has at least 2^k
nodes in its tree.

Corollary: analogously to

Property 2, the result holds
for any node not only the root.

Property 3 \Downarrow there are n elements
overall, there are at most
 $n/2^k$ nodes at rank k .

What does this imply for our
analysis?

Property 3 \Rightarrow maximum rank is $\log n$

\Rightarrow height of each tree $\leq \log n$

\Rightarrow find(x) is $O(\log n)$

Since union(x,y) calls
find(x) and find(y)
and otherwise is $O(1)$

Union(x,y) is $O(\log n)$

Note: $O(\log |E|) = O(\log |V|)$
because $|E| \leq |V|^2$ and
 $\log(|E|) \leq \log(|V|^2) = 2 \log |V|$

Kruskal's algorithm

sort the edges by weight $O(|E| \lg |V|)$

then $|V|$ makeset ops $O(|V|)$

$|E|$ union/find ops $O(|E| \log |V|)$

$O(|E| \log |V|)$

Can this be improved?

assume that the weights are sorted or are sorted faster than $O(|E| \lg |V|)$ e.g., if weights are bounded.

Then, we need to improve the complexity of $\text{find}(x)$.

This can only be done if the height is better than $\log n$.

This may not be possible w.c. but is possible over a sequence of operations.

Some operations will take longer, some will be very fast.

Let us sum the costs up for all operations.

Amortized cost per operation is this sum divided by the number of operations.

With a nice "trick" the amortized cost per find operation can be shown to be almost $O(1)$, it is $O(\log^* n)$.

$\log^* n$ is the number of times the log can be taken before 1 or < 1 is reached.

Example: $\log^* 1000 = 4$

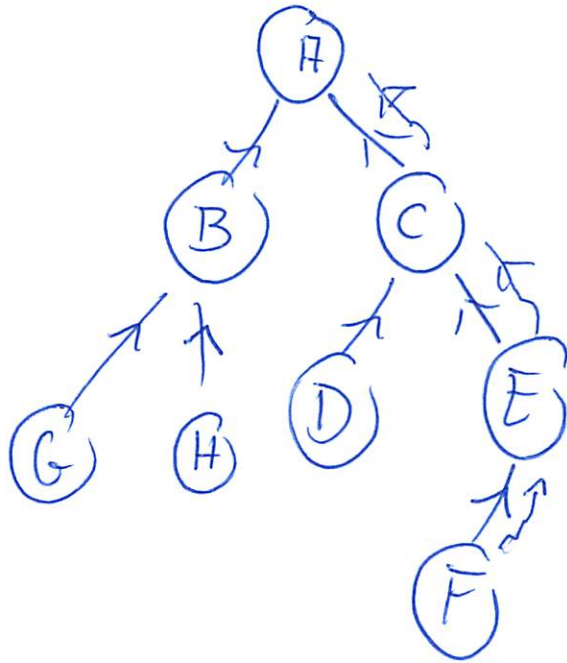
$$\log \log \log \log 1000 \leq 1$$

$$\log^* n \leq 5 \quad \text{if } n \leq 2^{65536}$$

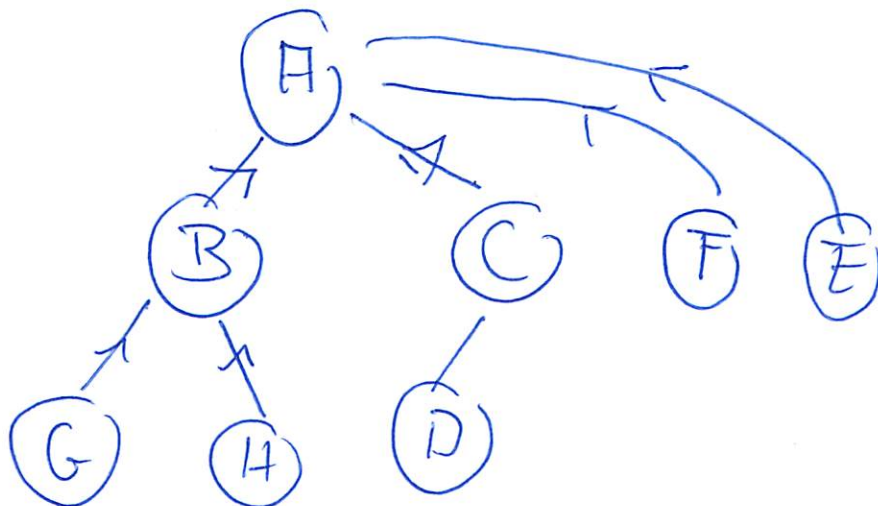
for all practical problems this will hold.

What is the "trick",

When a find(x) operation is done, we go up the tree to the root.



Why not reduce this for next time?



hang the node on the path directly under the root.

function find(x)

i) $x \neq \pi(x)$ then

$$\pi(x) = \text{find}(\pi(x))$$

return $\pi(x)$

This path compression reduces the amortized cost per find/union operation to $O(\log^* n)$

$\Rightarrow O(|E|)$ union/find operation can be carried out in

$O(|E| \lg^* n)$ time.

The idea of the analysis in the "amortized" way is to give operations some money. The cheap operations are not paying. The other operations are then shown to "pay" no more than made available - i.e., the amortized cost * # of operations.

Details not covered here but are in text book.

PRIM's Algorithm

$X = \{\}$ (edges picked so far)

repeat until $|X| = |V| - 1$

pick a set $S \subset V$ for
which X has no edges between
 S and $V - S$

let $e \in E$ be the minimum
weight edge between S and $V - S$

$X = X \cup \{e\}$.

Some what similar in flavour
to Dijkstra's algorithm.

The subtree defined by X
always grows by one edge,
which is the lightest edge between
 S and $V-S$.

it is the smallest cost vertex
 $\text{Cost}(v) := \min_{u \in S} w(u, v)$

The correctness follows from
the Cut property.

Detailed Prim's Algorithm

Procedure Prim (G, w)

input: A connected undirected graph $G=(V, E)$
with edge weights w_e

Output: A minimum spanning tree for G

for all $u \in V$

cost(u) = ∞

prw(u) = nil

pick any initial node u_0

cost(u_0) = 0

$H = \text{makequeue}(V)$ [PQ : keys = costs]

while H is not empty do

v = delete min (H)

for each $(v, z) \in E$

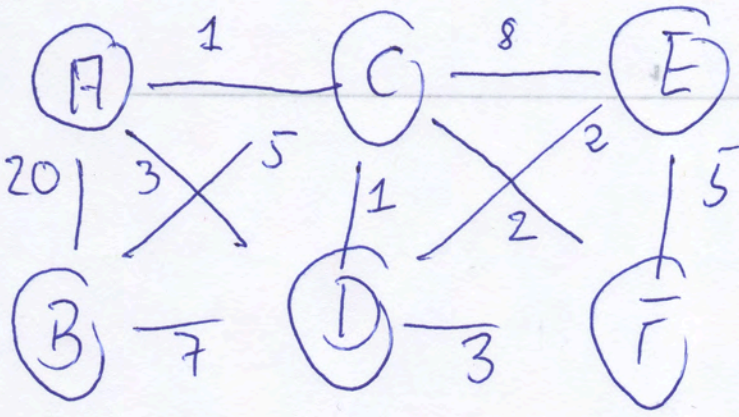
if cost(z) > $w(v, z)$

then cost(z) = $w(v, z)$

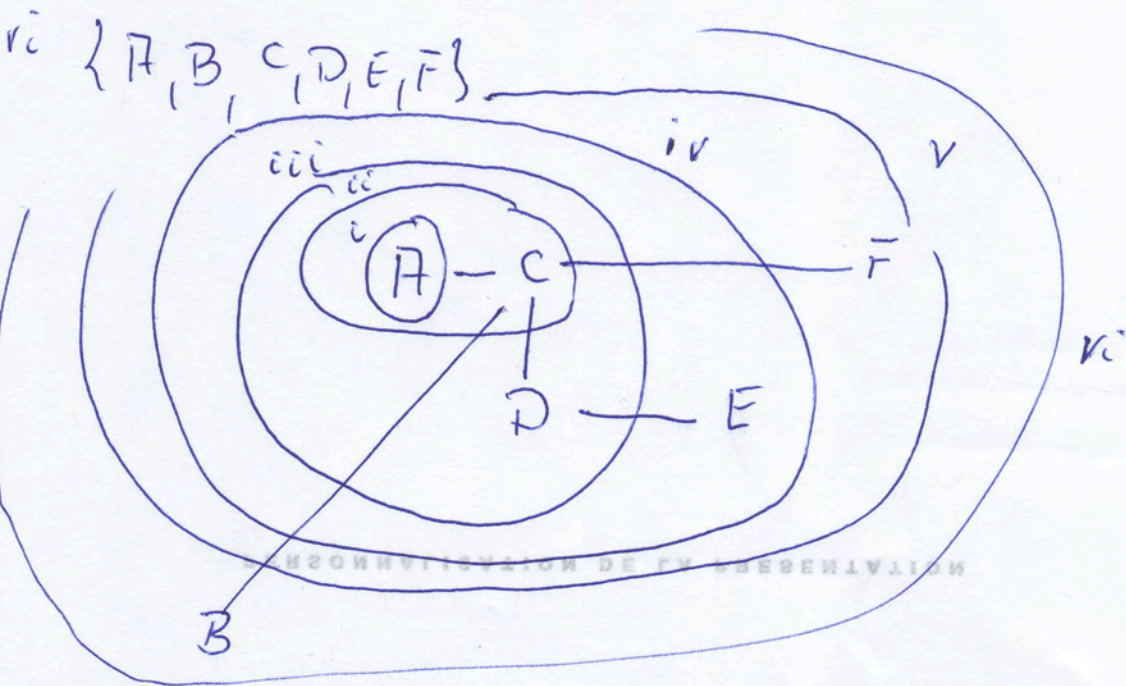
prw(z) = v

decreasekey(H, z)

} note
errors
in
textbook



Set S	A	B	C	D	E	F
$\{\}$	0/ml	∞ /ml	∞ /ml	∞ /ml	∞ /ml	∞ /ml
i $\{A\}$		20/A	1/A	3/A	∞ /ml	∞ /ml
ii $\{A, C\}$			5/C		1/C	8/C
iii $\{A, C, D\}$			5/C		2/D	3/C
iv $\{A, C, D, E\}$			5/C			2/C
v $\{A, C, D, E, F\}$			5/C			



Dynamic Programming

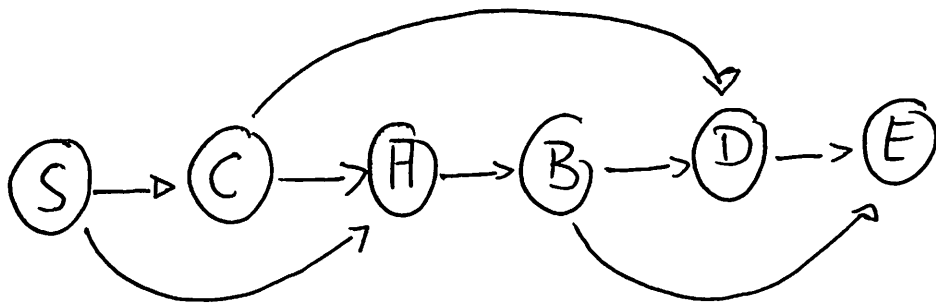
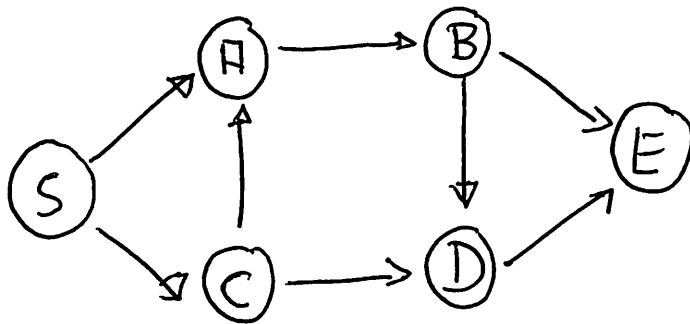
chapter 6 textbook

A useful, albeit sometimes costly, tool.

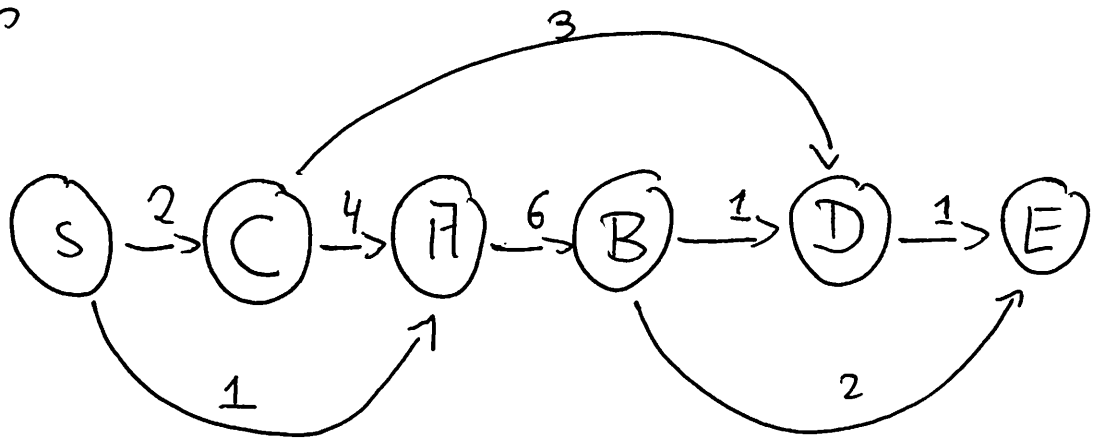
An example Shortest Paths in DAGs
(DAG = directed acyclic graph)

Property DAGs can be linearized, i.e., arranged on a line and edges are oriented left to right.

Reason: topological sort



Now consider the shortest path problem on DAGs. So, introduce costs on the edges



Now realize that, in order to compute the distance to a node further right we need to know the distance to its left vertices. We start from S, as always.

$$\text{ex.: 1) } \text{dist}(D) = \min(\text{dist}(C) + 3, \text{dist}(B) + 1)$$

$$2) \text{dist}(E) = \min(\text{dist}(B) + 2, \text{dist}(D) + 1)$$

This holds for all vertices analogously.

A single pass over the linearized graph gives all distances from S .
 $D(v) = \text{dist}(v)$

Procedure SP(DAG):

input: linearized DAG $G=(V, E)$, weights l

output: all distances $D(v), v \in V$
 $D(v) = \text{distance } S \text{ to } v \text{ in } G$

initialize $D(\cdot)$ to ∞

$D(s) = 0$

for each $v \in V \setminus \{s\}$ do
in linearized order

$$D(v) = \min_{(u,v) \in E} \{D(u) + l(u,v)\}$$

Analogously, we can compute the longest path. Note that longest path is not always as easy as shortest path.

Dynamic programming computes the value at a node using a function (min or max) of values of predecessor node.

A problem is solved by solving (many) subproblems & combine their solutions via minimization (maximization) to get a solution to the original problem.

Another example, LIS

(longest increasing subsequence)

Input: A sequence of numbers a_1, \dots, a_n

Output: A subsequence $a_{i_1}, a_{i_2}, \dots, a_{i_k}$
 $i_1 \leq i_2 \leq \dots \leq i_k \leq n$

s.t. $a_{i_1} \leq a_{i_2} \leq \dots \leq a_{i_k}$ and
 k is the largest integer for which this holds.

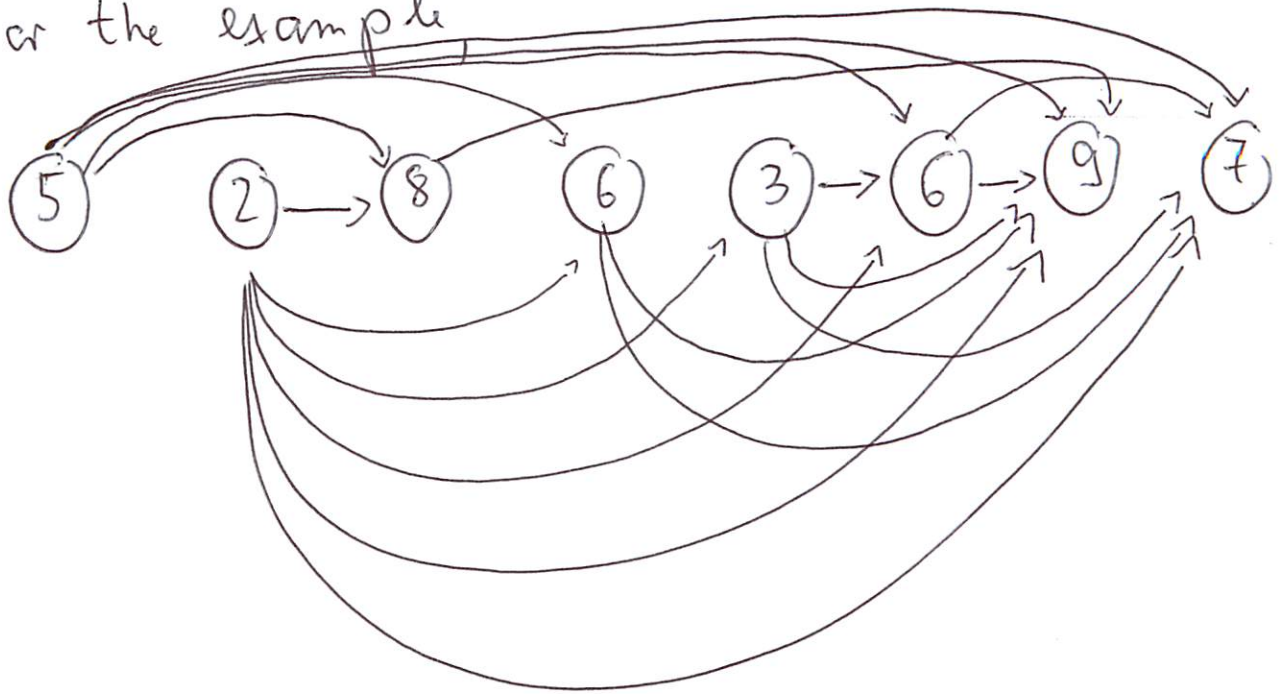
Example. 5 2 8 6 3 6 9 7
 a_1 a_2 a_3 a_4 a_5

5 8 9 is an increasing sequence $k=3$

but
2 3 6 9 is both increasing and longer $k=4$

Let us rotate this in terms of a graph
 The vertices are linearized by the input
 order, the edges $(a_i) \rightarrow (a_j)$ are $(a_i, a_j) \in E \Leftrightarrow i < j$
 and $a_i < a_j$

For the example



Observation : This is a DAG. G

LIS problem $\hat{=}$ longest path in G

for $j = 1, \dots, n$ do

$$L(j) = 1 + \max \{ L(i) : (i, j) \in E \}$$

return $\max_j L(j)$

Why is dynamic programming sometimes good?

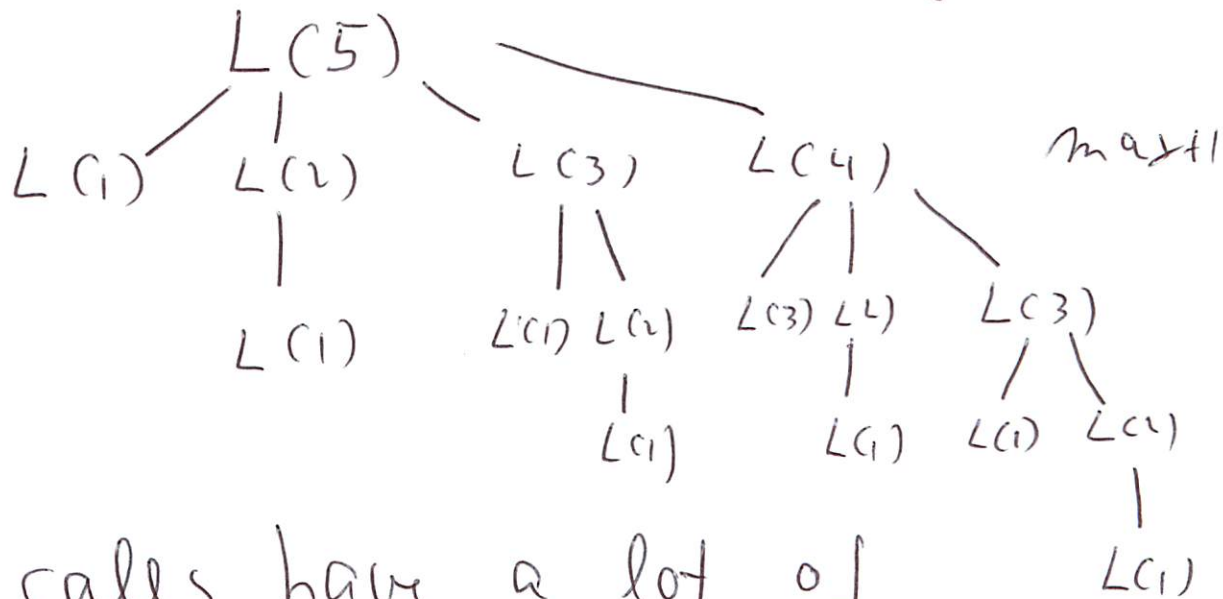
Consider the LIS problem.

Now, solve it via a recursive algorithm. [bad here but we want to understand why]

Say that the a_i 's happen to be sorted. Then

$$L(j) = 1 + \max\{L(1), L(2), \dots, L(j-1)\}$$

Visualize the recursion, say $j=5$



The calls have a lot of redundancies. Dynamic programming stores the results so that they are available.

Question how many calls are
made to $L(1), L(2), \dots$?

Divide & Conquer (recursion)
does well when subproblem sizes
are dramatically reduced.

Dynamic programming builds
up solutions from all smaller
sized problems.

A geometric problem for dynamic programming.

Min Weight Triangulation

Motivation: e.g. 1) CAD

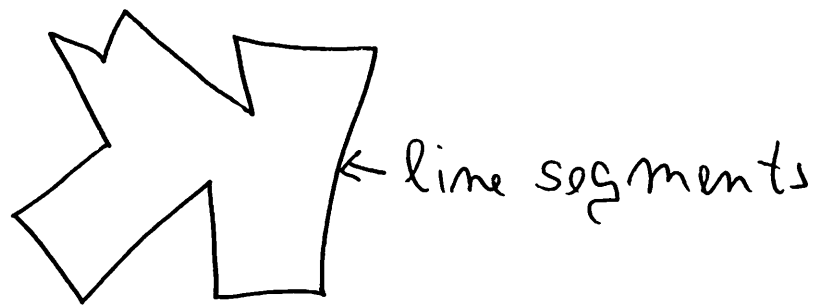
2) estimation of ice thickness

See class for details

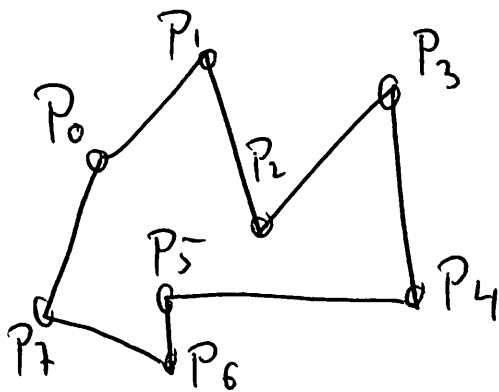
Triangulation:

Problem setting for polygons
in fact let us look at convex
polygons

A polygon is a closed plane figure bounded by line segments



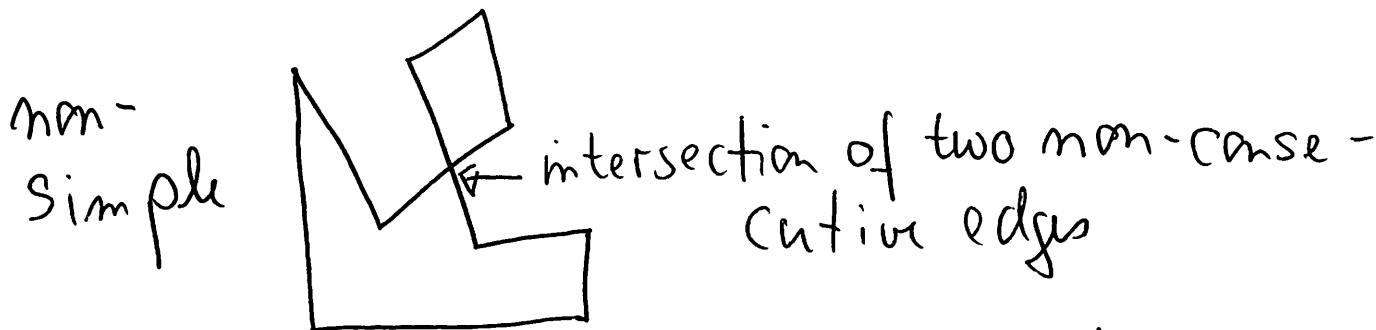
We order the vertices of the polygon clockwise around its boundary.



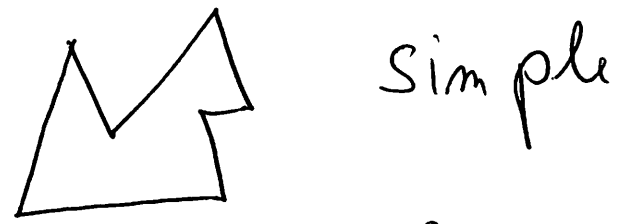
polygon
 $= (P_0, P_1, \dots, P_n)$
 P_{i-1}, P_i is a line segment
 $i = 1, \dots, n.$

Typically, one distinguishes between simple polygons and non-simple or arbitrary polygons.

A polygon is non-simple if there are two edges e_i, e_j $j \neq i+1, j \neq i-1$ which intersect.

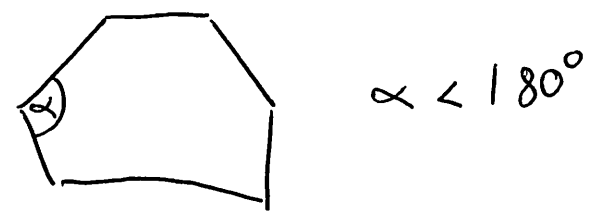


consecutive edges are only allowed to intersect at their common end point.

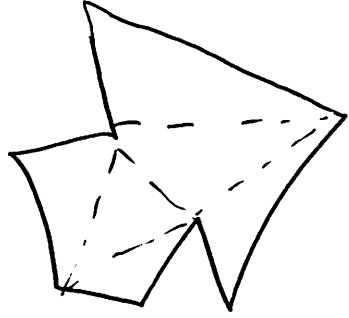


⊗ We consider from now on only simple polygons.

A polygon is convex if each internal angle $\leq 180^\circ$.



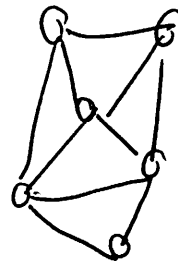
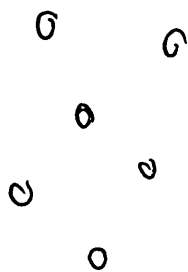
A triangulation of a polygon is a partition of the polygon into triangles. The partition edges must join vertices of the polygon.



each face is a triangle
additional edges
join vertices

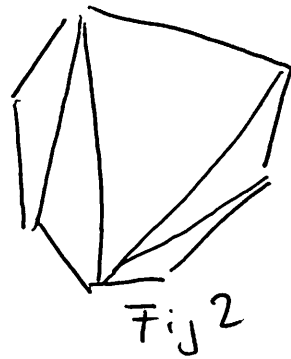
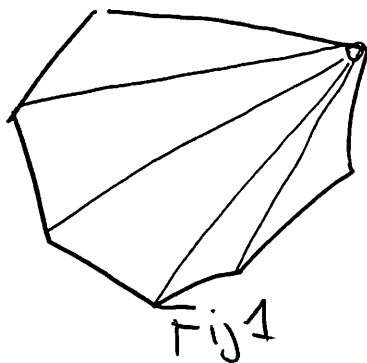
An alternate definition is that a triangulation is a planar graph with the maximal # of edges build on the vertices (+ edges of the polygon).

The latter definition extends to triangulation of point sets



Triangulations are important intermediate steps for solving many geometric problems.

There are many different triangulations of the same input polygon. They differ substantially.



Assume you had to draw these with the min ink possible.

$$\min_T \sum_{e \in T} d(e), \text{ where } T \text{ is a triangulation}$$

$d(e)$ = distance between the 2 vertices of e

We minimize the sum of all distances of the added diagonals over all possible triangulations.

We call $\sum_{e \in T} d(e)$ the weight of triangulation T .

Min Weight triangulation

Find a triangulation of a given polygon with min weight over all possible triangulations.

Solution Dynamic Programming

Key insight

there is a triangle T_i which partitions

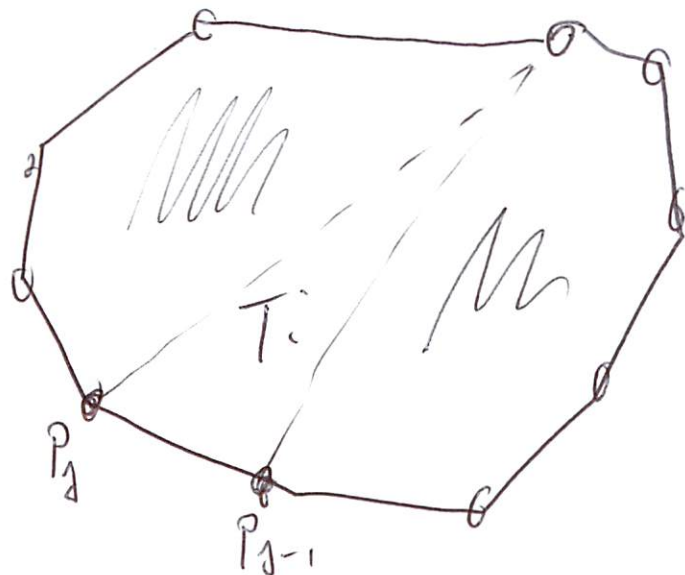
the problem

into smaller

subproblems of the same type

Then, the weight of T_i + that of the subproblems is the weight of the entire triangulation.

Can we make it 2 subproblems?



Yes! because the edge $P_{i-1}P_i$ must be in one triangle.

This is a better insight as we have less subproblems to combine and only need to iterate over all vertices with whom $P_{i-1}P_i$ can be in a triangle.

1. Let v_0, \dots, v_n be the $n+1$ vertices of an input polygon.
2. Call $C_{0,n}$ the cost (weight) of the minimum weight triangulation
 $C_{i,k}$ analogous defined for subpolygon with vertices v_i, \dots, v_k , $k \geq i+2$

Case 1: $k < i+2$ no triangulation possible

Case 2 $k \geq i+2$ then there are choices for a j
 $i < j < k$.

For each choice of j , calculate.

$\min C_{i,j} + C_{j,k} + w(i,j,k) =: C_{i,k}$
the minimum over all appropriate j

$$C_{i,k} = \begin{cases} 0 & k < i+2 \\ \min_{i < j < k} \{C_{i,j} + C_{j,k} + w(i,j,k)\} & \text{otherwise.} \end{cases}$$

Record which index j gives the minimum
call that table m ($m_{i,k} = j$)

This gives the basis for the dynamic programming.

Time complexity $O(n^3)$

Knapsack

We wish to put items in a knapsack. Each item has a weight and a value. The knapsack has a particular max. capacity. The task is to maximize the total value of the items that fit into the knapsack.

<u>Ex:</u>	Item	Weight	Value
	1	6	\$30
	2	3	\$14
	3	4	\$16
	4	2	\$9

max Capacity = 10

Two variants

- ① each item is unique
- ② every item is available in unlimited quantity

Solution to ①

$$\begin{array}{l} \text{item 1 + item 3} \\ \text{capacity reached} \quad \text{value } \$30 \\ 6 + 4 = 10 \quad \quad \quad \begin{array}{r} + \$16 \\ \hline \$46 \end{array} \end{array}$$

Solution to ②

$$\begin{array}{l} \text{item 1 + 2 * item 4} \quad \text{value } \$30 \\ \text{capacity: } 6 + 2 * 2 = 10 \quad \quad \quad \begin{array}{r} + \$18 \\ \hline \$48 \end{array} \end{array}$$

General solution to (2)

Let $k(w) = \max$ value achievable
with a knapsack of capacity w

then,

$$k(w) = \max_{i: w_i \leq w} \{ k(w - w_i) + v_i \}$$

Interpretation:

if the optimal solution with $k(w)$
contains item i , then removing
 v_i leaves an optimal solution
to $k(w - w_i)$

Algorithm

input: capacity W , weights, values

output: optimal solution for $k(w)$

$$k(0) = 0$$

for $w = 1$ to W

$$k(w) = \max \{ k(w - w_i) + v_i \mid w_i \leq w \}$$

return $k(w)$

Complexity: $O(nW)$

1-d table

General Solution to ①

So, if no repetitions are allowed

once we remove an item w_i

we cannot just take $k(w - w_i)$

as the sub solution might contain that element. Since it is unique we cannot use it twice.

Add a second parameter j , $0 \leq j \leq n$

define $k(w, j) = \text{max value achievable using a knapsack of capacity } w \text{ and items } 1 \dots j$

Final answer $k(w, n)$

Subproblems

$$k(w, j) = \max \left\{ k(w - w_j, j-1) + V_j, k(w, j-1) \right\}$$

So either V_j is in or not.
at step j

Initialize all $k(0, j) = 0$ and
all $k(w, 0) = 0$

for $j = 1$ to n do

for $w = 1$ to W do

if $w_j > w$ then $k(w, j) = k(w, j-1)$

else $k(w, j) = \max \left\{ k(w, j-1), \right.$
 ~~$k(w - w_j) + V_j$~~
 $\left. k(w - w_j, j-1) + V_j \right\}$

return $k(W, n)$

2-d table required
still $O(mW)$ complexity

Chain Matrix Multiplication

Task: 4 matrices to multiply

$$A \times B \times C \times D$$

The matrices may have different dimensions, say

$$A : 50 \times 20$$

$$B : 20 \times 1$$

$$C : 1 \times 10$$

$$D : 10 \times 100$$

Matrix multiplication is associative.

$A \times B \times C \times D$ can be bracketed in different ways

$$(A \times B) \times (C \times D) \quad ((A \times B) \times C) \times D$$

$$A \times (B \times C) \times D \quad A \times (C \times B) \times (D)$$

e.g., $A \cdot B$ has dim $50 \times \underline{1}$

which reduces subsequent computations

<u>Parenthesization</u>	<u>Cost Computation</u>	<u>total Cost</u>
$A \times ((B \times C) \times D)$	$20 \cdot 1 \cdot 10 + 20 \cdot 10 \cdot 100 + 50 \cdot 20 \cdot 100$	120,200
$(A \times (B \times C)) \times D$	$20 \cdot 1 \cdot 10 + 50 \cdot 20 \cdot 10 + 50 \cdot 10 \cdot 100$	60,200
$(A \times B)(C \times D)$	$50 \cdot 20 \cdot 1 + 1 \cdot 10 \cdot 100 + 50 \cdot 1 \cdot 100$	7,000

So, with a clever parenthesization one can save MANY computations.

Greedy fails here.

Question: Let A_1, \dots, A_m be 2-d matrices, A_i has dimension $(m_{i-1}, m_i)_{i=1, \dots, m}$.
 Find a parenthesization of $A_1 \dots A_m$ that minimizes the total cost.
 How to find this?

Fact: Valid bracket sequences are in 1-1 correspondence with full binary trees.

A bracket sequence is well-formed if

1) • $\# " (" = \# ")"$ equal number of opening and closing brackets

2) • $\# " (" \geq \# ")"$ when reading the sequence left to right (at any point)

Ex: - (()) () is well-formed

- ()) (is not

$$\begin{array}{c} \uparrow \\ \# "(" = 1 \\ \# ")" = 2 \end{array}$$

violates 2)

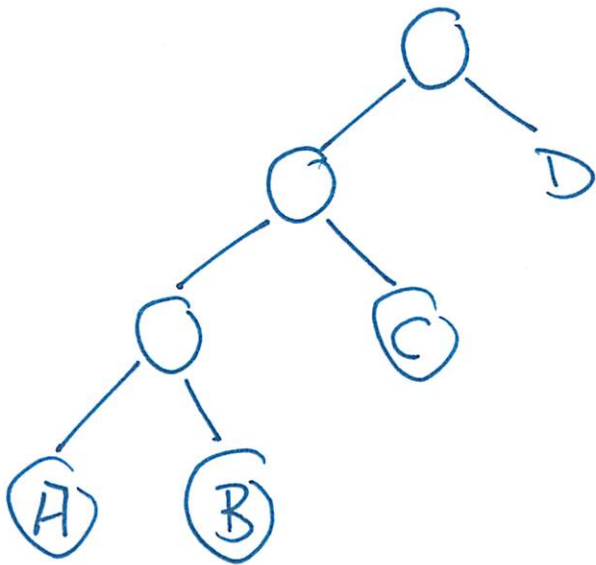
- () () (is not

$$\begin{array}{c} \# "(" = 3 \\ \# ")" = 2 \end{array}$$

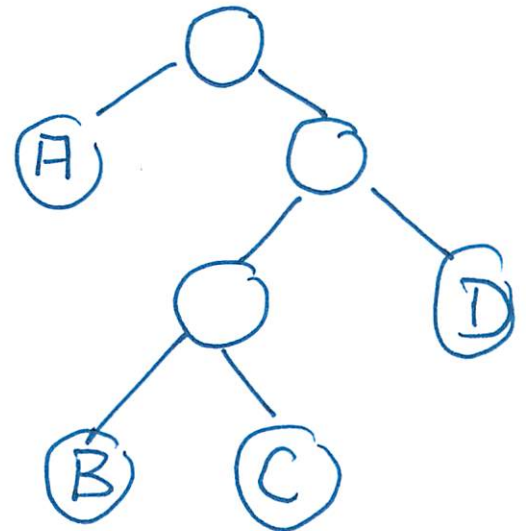
violates 1)

Here, we can see the multiplication arranged in a tree

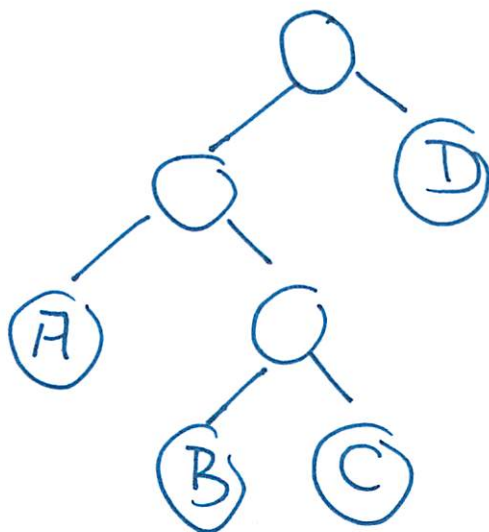
$$((A + B) + C) + D$$



$$A \times ((B \times C) \times D)$$



$$(A + (B \times C)) + D$$



Fact: Every subtree corresponds to a particular way of bracketing a subproblem

$$A_i \cdots A_j$$

Fact: For the entire tree to be optimal each subtree must be optimal.
(i.e. have optimal cost)

$$C(i, j) := \min \text{cost of multiplying } A_i \times \cdots \times A_j$$

$$C(i, i) = 0$$

$$C(i, j) = \min_{i \leq k < j} \{ C(i, k) + C(k+1, j) + m_{i-1} m_k m_j \}$$

this is

$$\underbrace{\left(\underbrace{A_i \times \cdots \times A_k}_{\min} \right)}_{\min} \cdot \underbrace{\left(\underbrace{A_{k+1} \times \cdots \times A_j}_{\min} \right)}_{\min} \cdot \underbrace{m_{i-1} m_k m_j}_{\text{cost of multiplying}}$$

Dynamic Program

for $i = 1$ to n do $C(i, i) = 0$

for $s = 1$ to $n - 1$ do

for $i = 1$ to $n - s$ do

$$j = i + s$$
$$C(i, j) = \min \left\{ C(i, k) + C(k+1, j) \right. \\ \left. + m_{i-1} m_k m_j \right. \\ \left. \text{over } i \leq k < j \right\}$$

return $C(1, n)$

2-d table each entry cost $O(n)$

to compute: $O(n^3)$

The dynamic program prevents us from having to construct all binary trees for the bracketing. This would be exponential.

MATRIX-CHAIN-ORDER(p)

```

1   $n = p.length - 1$ 
2  let  $m[1..n, 1..n]$  and  $s[1..n - 1, 2..n]$  be new tables
3  for  $i = 1$  to  $n$ 
4       $m[i, i] = 0$ 
5  for  $l = 2$  to  $n$            //  $l$  is the chain length
6      for  $i = 1$  to  $n - l + 1$ 
7           $j = i + l - 1$ 
8           $m[i, j] = \infty$ 
9          for  $k = i$  to  $j - 1$ 
10              $q = m[i, k] + m[k + 1, j] + p_{i-1}p_kp_j$ 
11             if  $q < m[i, j]$ 
12                  $m[i, j] = q$ 
13                  $s[i, j] = k$ 
14  return  $m$  and  $s$ 

```

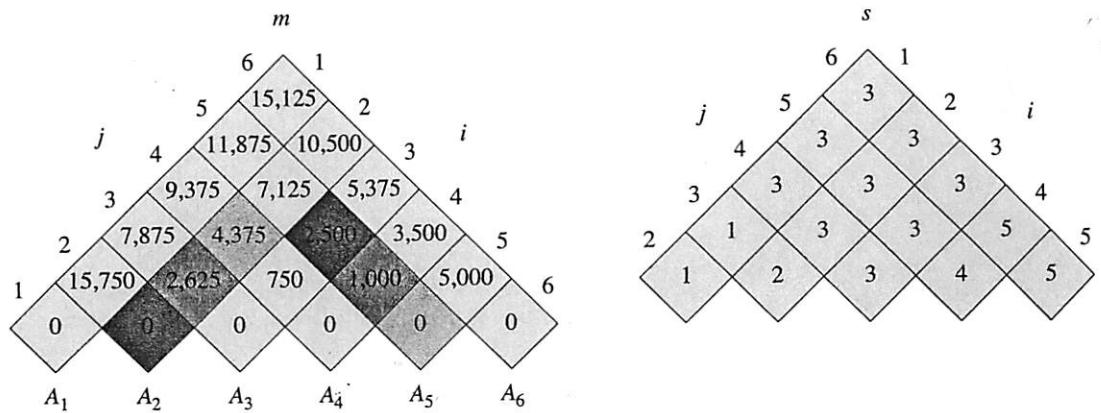


Figure 15.5 The m and s tables computed by MATRIX-CHAIN-ORDER for $n = 6$ and the following matrix dimensions:

matrix	A_1	A_2	A_3	A_4	A_5	A_6
dimension	30×35	35×15	15×5	5×10	10×20	20×25

The tables are rotated so that the main diagonal runs horizontally. The m table uses only the main diagonal and upper triangle, and the s table uses only the upper triangle. The minimum number of scalar multiplications to multiply the 6 matrices is $m[1, 6] = 15,125$. Of the darker entries, the pairs that have the same shading are taken together in line 10 when computing

$$\begin{aligned}
 m[2, 5] &= \min \begin{cases} m[2, 2] + m[3, 5] + p_1 p_2 p_5 = 0 + 2500 + 35 \cdot 15 \cdot 20 = 13,000, \\ m[2, 3] + m[4, 5] + p_1 p_3 p_5 = 2625 + 1000 + 35 \cdot 5 \cdot 20 = 7125, \\ m[2, 4] + m[5, 5] + p_1 p_4 p_5 = 4375 + 0 + 35 \cdot 10 \cdot 20 = 11,375 \end{cases} \\
 &= 7125.
 \end{aligned}$$

7. Linear Programming

Some material presented uses:
"Formulating Integer Linear Programs:
A Rogues' Gallery"
G. Brown, R. Dell

Many problems ask to minimize
or maximize an objective function.

given a linear function of certain
variables and (possibly) competing
constraints.

Linear-programming Problem:

- objective function is linear
- constraints equalities or inequalities
(linear)

Example

Profit Maximization

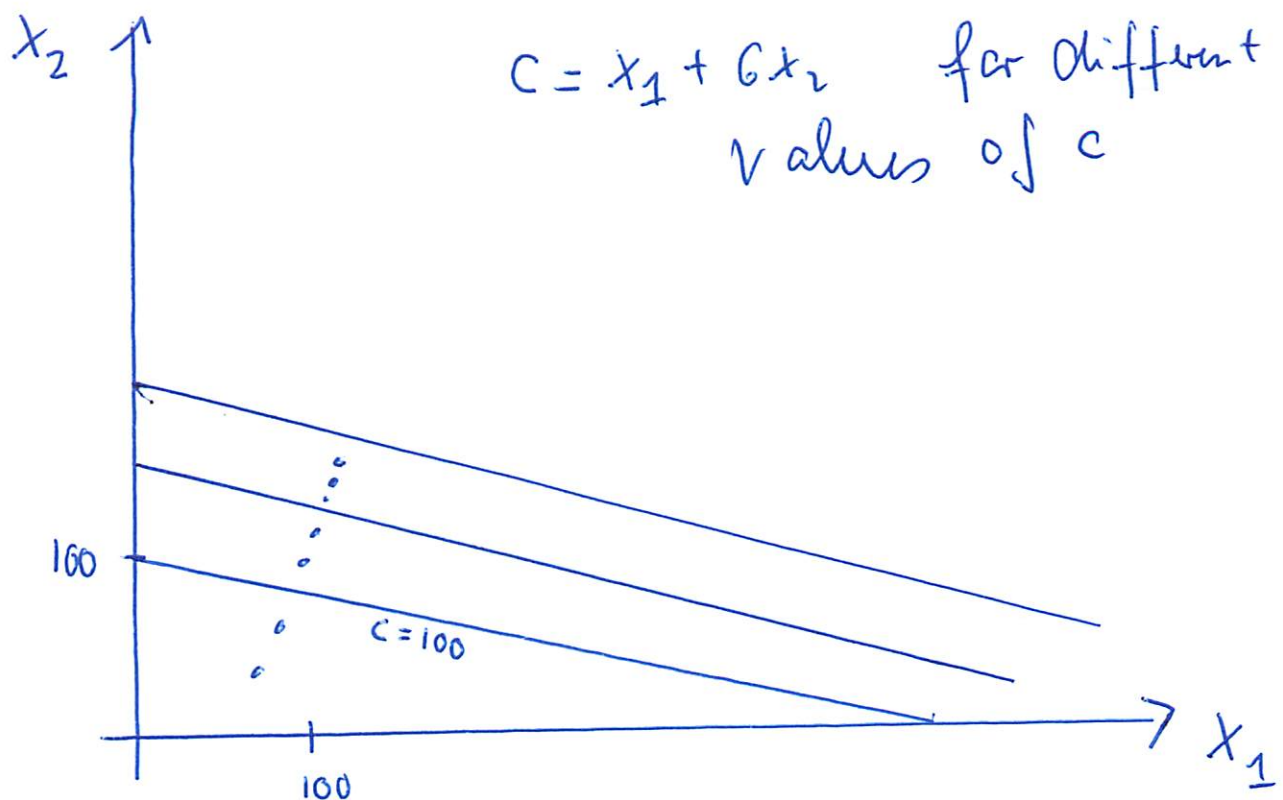
2 products

P_1 and P_2

1. x_1 boxes of P_1 can be produced per day
2. x_2 " " P_2
3. Profit from P_1 : \$1
4. Profit from P_2 : \$6
5. daily demand for P_1 : ≤ 200
6. " " " P_2 : ≤ 300
7. at most a total of 400 boxes can be produced per day

Q: How many boxes of P_1 and P_2 should we produce to maximize our profit?

Objective function, $\max 1 \cdot x_1 + 6 \cdot x_2$
it is linear in x_1, x_2 in 2-d space



for different values of c , we get different parallel lines.

The higher up the line is, the better the profit.

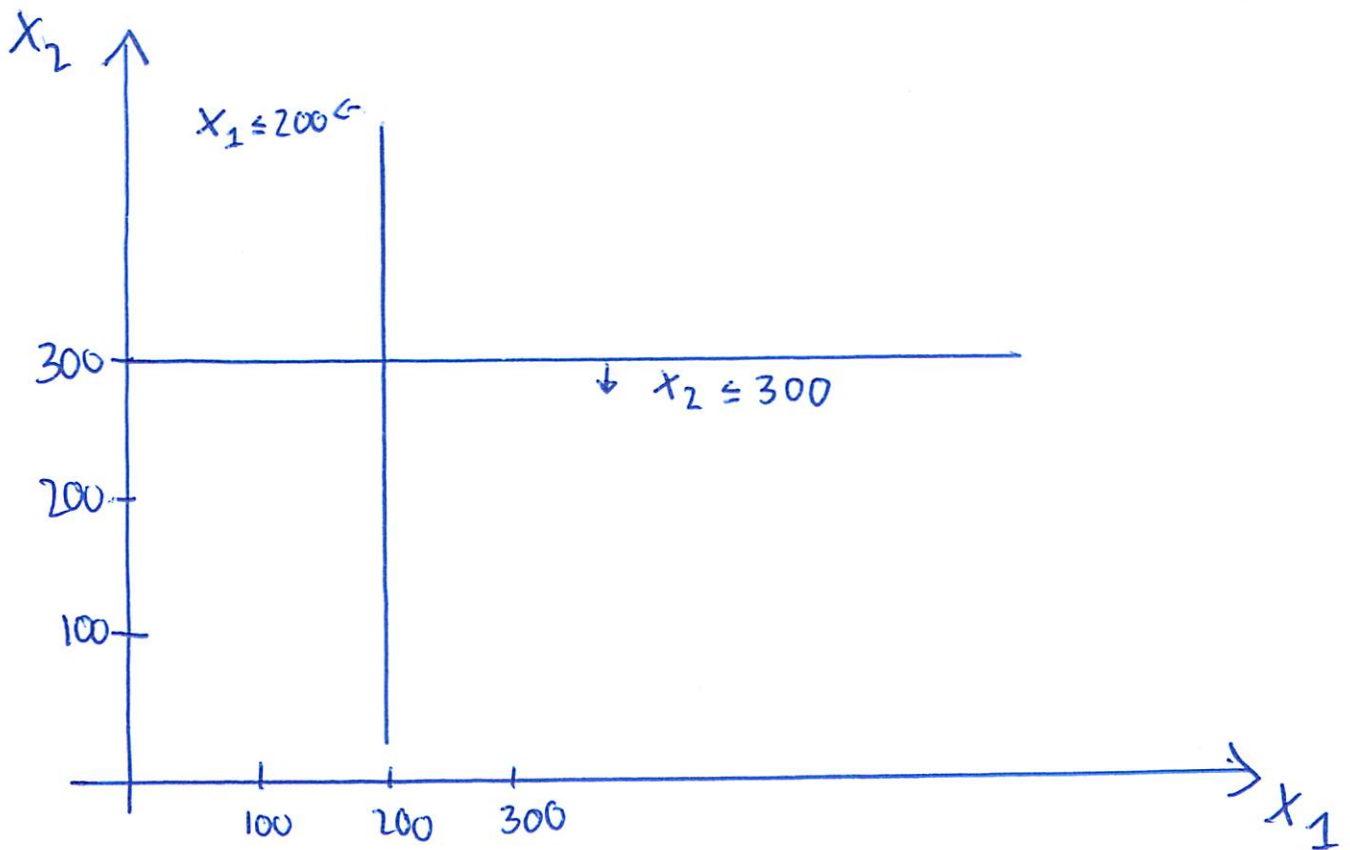
But, we have constraints preventing us to shift the line arbitrarily high up.

It is obvious $x_1 \geq 0, x_2 \geq 0$

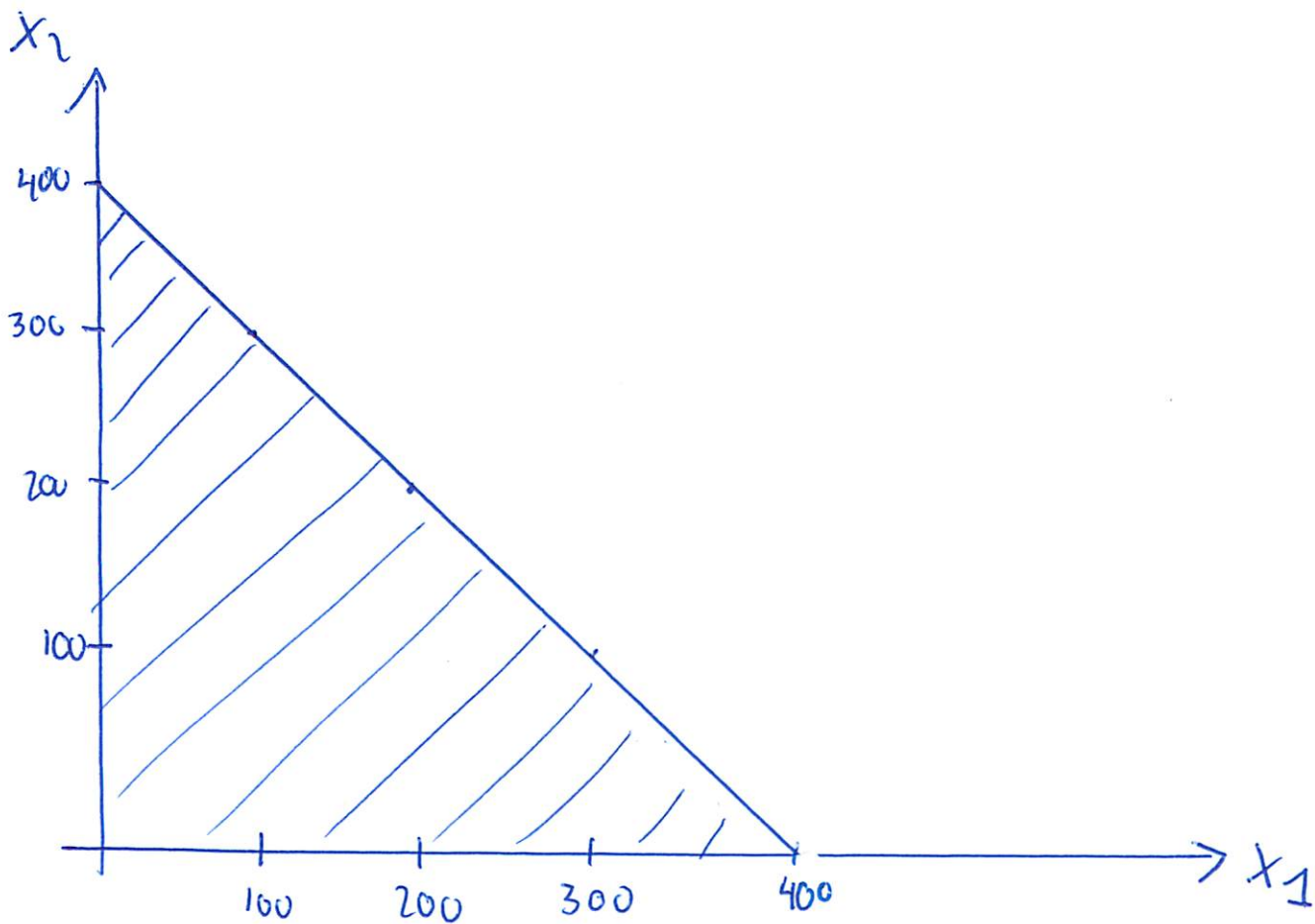
$$\begin{array}{l}
 5. \quad \Rightarrow \quad x_1 \leq 200 \\
 6. \quad \Rightarrow \quad x_2 \leq 300 \\
 7. \quad \Rightarrow \quad x_1 + x_2 \leq 400
 \end{array}
 \left. \vphantom{\begin{array}{l} 5. \\ 6. \\ 7. \end{array}} \right\} \begin{array}{l} \text{these are} \\ \text{the} \\ \text{constraints} \end{array}$$

linear constraints.

How do they look like graphically?



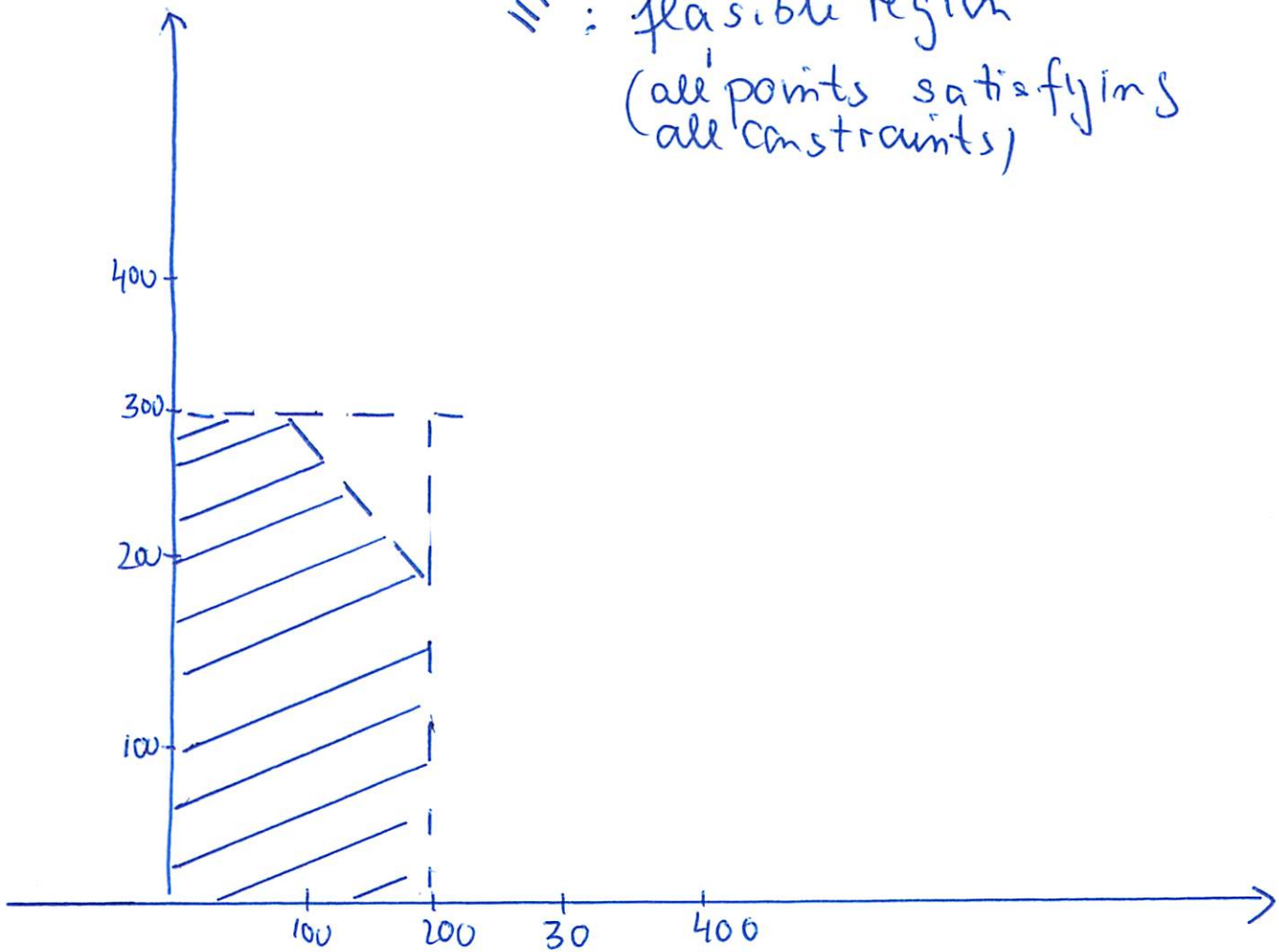
Constraints 5. + 6.



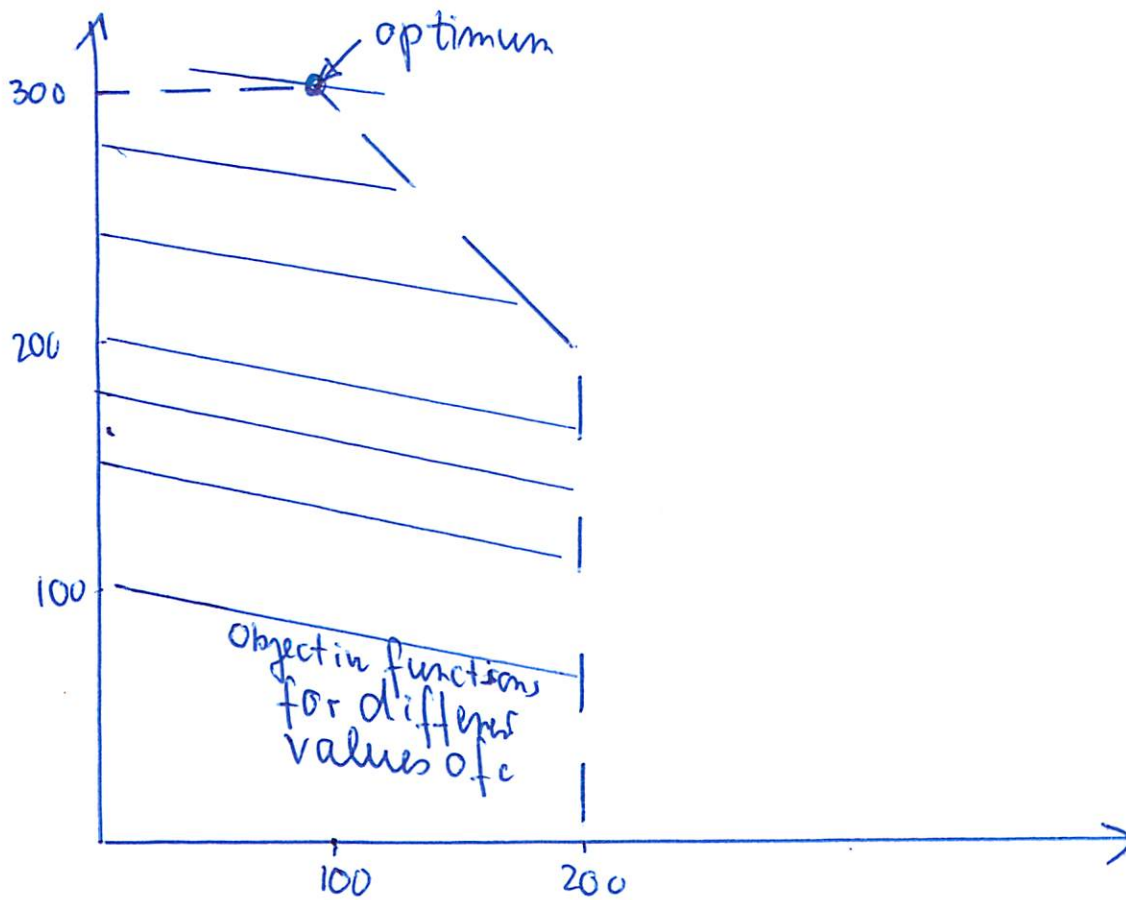
Constraint 7.

Now let us put the constraints into one graph

≡ : feasible region
(all points satisfying
all constraints)



Now we wish to find a point
which maximizes the profit.



the function maximization means going up & right until we are about to leave the feasible region.

This happens at a vertex of the feasible region.

Two cases where the optimum is not at a vertex

① the linear program is infeasible

② there is an unbounded feasible region

① $x_1 \leq 1$, and $x_1 \geq 2$
these are (obviously) conflicting / contradictory constraints.
 \Rightarrow no solution

② $\left. \begin{array}{l} \max \quad x_1 + x_2 \\ x_1 \geq 0 \quad x_2 \geq 0 \end{array} \right\} \text{unbounded}$

Solving linear Programs (LP)

Here is the good news:

there are packages!

We don't solve, but simply call the LP Solver.

These solvers are hot commercial commodities.

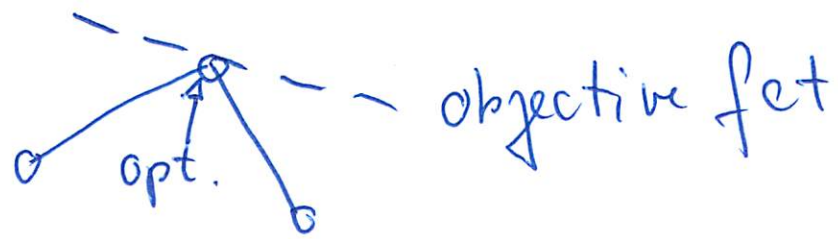
First solution by George Dantzig.

Method is based on hill climbing.

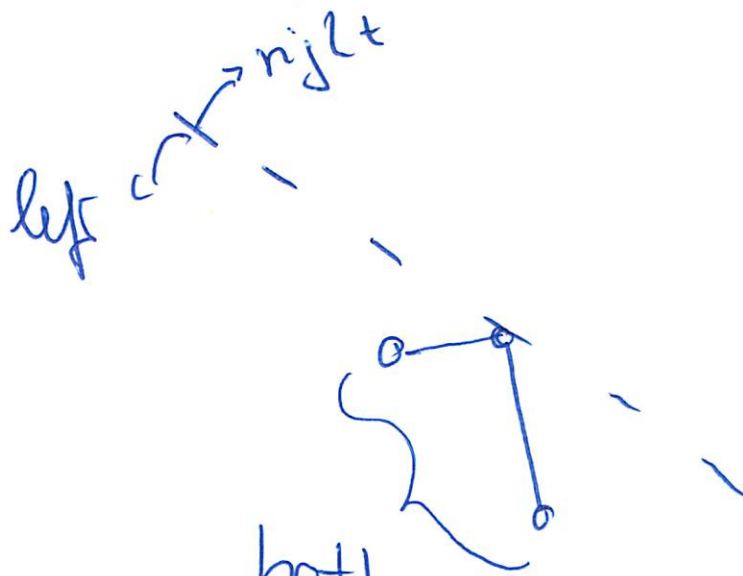
Start at a vertex of the feasible region. Calculate value of objective function.

Look for an adjacent vertex of better value \Rightarrow hill climbing.

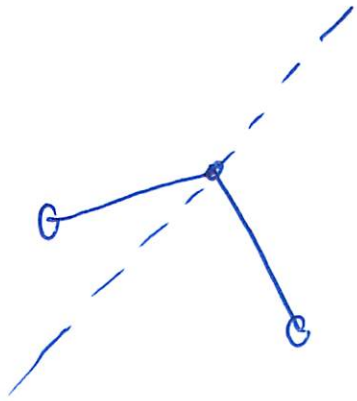
done when a vertex is reached with no better neighbor.



geometrically, both vertices lie on the same side of the objective function



both lie of the left of the line (objective function)



a better solution
is possible as
the 2 vertices
are on opposite
sides

Example 2

1. x_1 boxes of P_1 can be produced per day
 2. x_2 " " P_2 " " " " "
 3. x_3 " " P_3 " " " " "

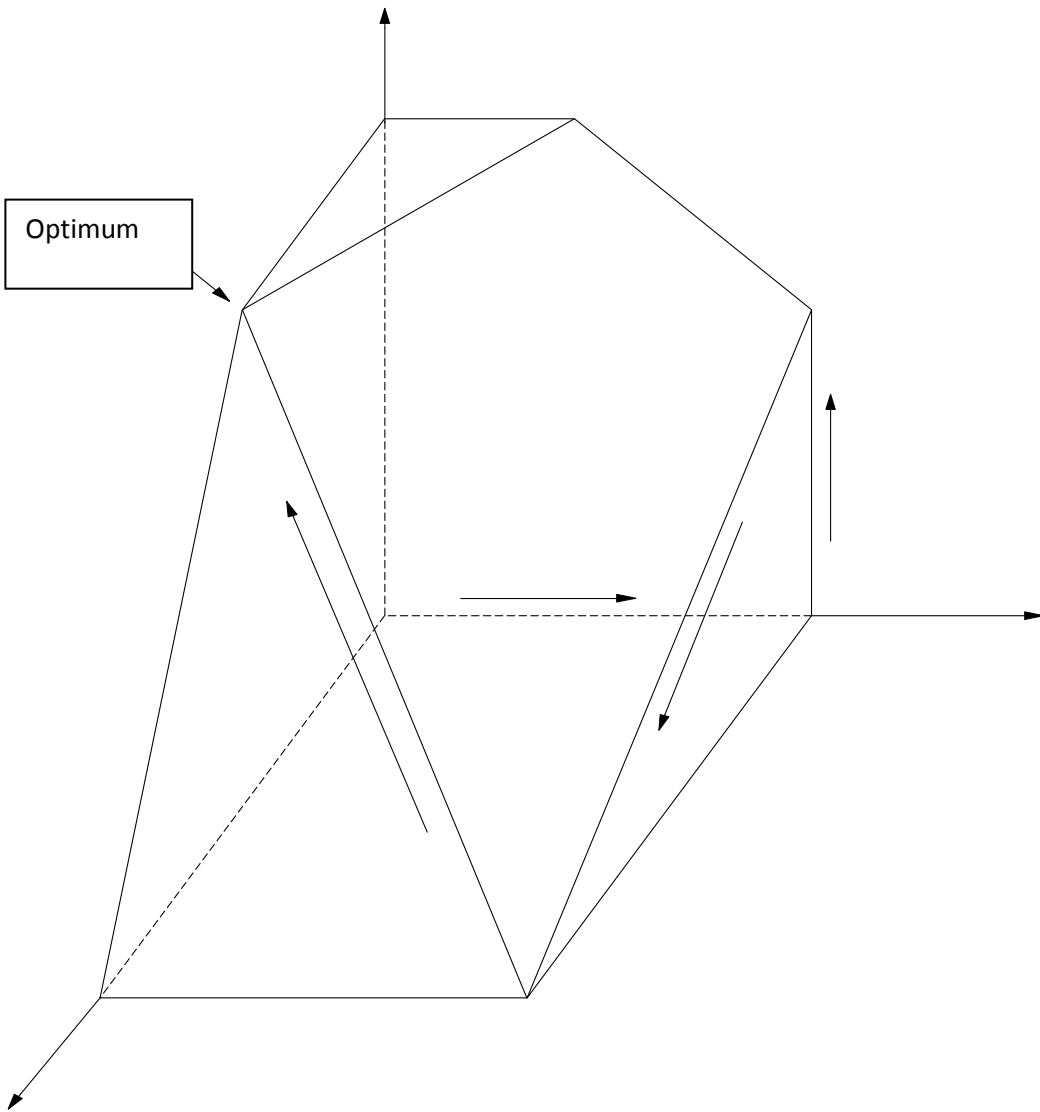
 4. Profit from P_1 : \$1
 5. " " P_2 : \$6
 6. " " P_3 : \$13

 7. daily demand for P_1 : ≤ 200
 8. " " " P_2 : ≤ 300

 9. at most a total of 400 boxes can be produced per day

 10. P_3 requires 3x more packaging than P_2 and together $3 \cdot P_3 + P_2 \leq 600$
-

Q: How many boxes of P_1, P_2, P_3 should we produce to maximize our profit?



Objective function

$$\text{max : } | 1 \cdot x_1 + 6 \cdot x_2 + 13x_3 |$$

subject to:

$$\bullet x_1, x_2, x_3 \geq 0$$

$$\bullet x_1 \leq 200$$

$$\bullet x_3 \leq 300$$

$$\bullet x_1 + x_2 + x_3 \leq 400$$

$$\bullet x_2 + 3x_3 \leq 600$$

} constraints

optimal : (0, 300, 100)

total profit \$3100

Exercise: check how simplex
would examine the vertices.

Main Elements of LP

(in fact of any constrained optimization problem)

- Variables

values to be found by the solution
they find/determine the best values for
the objective function

- Objective function

expression defined on the variables
to express the goal for the optimization

- Constraints

expresses limits on the possible
solutions via expressions using the variables

- Variable bounds

typically variables are bounded

In our 1st example:

- o variables x_1, x_2
- o objective function $\max x_1 + 6x_2$
- o Constraints $x_1 + x_2 \leq 400$
- o variable bounds $0 \leq x_1 \leq 200$
 $0 \leq x_2 \leq 300$

We will group constraints +
variable bounds under
heading constraints

Brown et al. use the term
"formulette"

LP Formulettes

For each formulette, we write constraints in terms of non-negative, continuous variables x_1, \dots

L1) "For each unit of x_1 , there must be at least 5 units of x_2 "

$$5x_1 \leq x_2$$

Frequent wrong answer: $x_1 \leq 5x_2$

When in doubt make an example.

$$x_1 = 3 \quad \Rightarrow \quad x_2 \geq 15$$

L2) "A port can load 11 x_1 's per week, or 45 x_2 's or 30 x_3 's.

What combination of x_1 , x_2 and x_3 can be loaded in 10 weeks?"

$$\left| \frac{1}{11} x_1 + \frac{1}{45} x_2 + \frac{1}{30} x_3 \leq 10 \right|$$

Most frequent wrong answer:

$$11x_1 + 45x_2 + 30x_3 \leq 10$$

Sanity check immediate shows that this is wrong.

$$11x_1 + 45x_2 + 30x_3 = 10$$

is also wrong: $x_1 = x_2 = x_3 = 0$ should be a feasible solution

L3) "There must be exactly 7 units of x_1 for every 9 units of x_2 "

$$\boxed{\frac{1}{7} x_1 = \frac{1}{9} x_2}$$

$$(or\ 9x_1 = 7x_2)$$

Again a simple example will tell you if you got it wrong.

L4) "Elements of type x_1 must constitute at most 33% of all elements of type x_1, x_2 and x_3 ."

$$0.67 x_1 \leq 0.33 x_2 + 0.33 x_3$$

How to obtain this?

$$\frac{x_1}{x_1 + x_2 + x_3} \leq \frac{33}{100}$$

But what is wrong with this constraint?

it is not a linear expression
(Linear Programming)

So, we need to convert it.

$$100x_1 \leq 33x_1 + 33x_2 + 33x_3$$

$$100x_1 - 33x_1 \leq 33x_2 + 33x_3$$

$$67x_1 \leq 33x_2 + 33x_3$$

25) "What mixtures of punch from 80-proof x_1 , 100-proof x_2 , and 0-proof x_3 are at least 30-proof?"

again, to derive it we state a non-linear inequality first.

$$80x_1 + 100x_2 + 0x_3 \geq 30(x_1 + x_2 + x_3)$$

$$\boxed{(80 - 30)x_1 + (100 - 30)x_2 + (0 - 30)x_3 \geq 0}$$

L6) "Process x_1 produces 24.5 units of x_2 and 73.1 units of x_3 per hour"

$$\begin{aligned}x_2 &= 24.5 x_1 \\x_3 &= 73.1 x_1\end{aligned}$$

Process x_1 produces simultaneous
at fixed rates 2 outputs.

These formulations are the basic building blocks for the LP solver input.

PRACTICE

LP solvers

① Simplex Method

In this method the algorithm go from corner to corner (of feasible region) always improving for an exponential number of steps.

In practice it runs fast on almost all inputs.

② 1975 a young Soviet mathematician Leonid Khachiyan designed the ellipsoid method.

to solve any LP in polynomial time. Theoretically better than the Simplex Method, but not practically.

③ 1984 Narendra Karmarker
grad student Uni. of California,
Berkeley

interior point method

Goes to corners through polyhedron
in a clever way.

Behaves well in practice.

Postscript: circuit evaluation

"The ultimate application"

Given a boolean circuit, i.e., a dag of gates of the following types.

- INPUT GATES

in degree = 0

values \in {true, false}

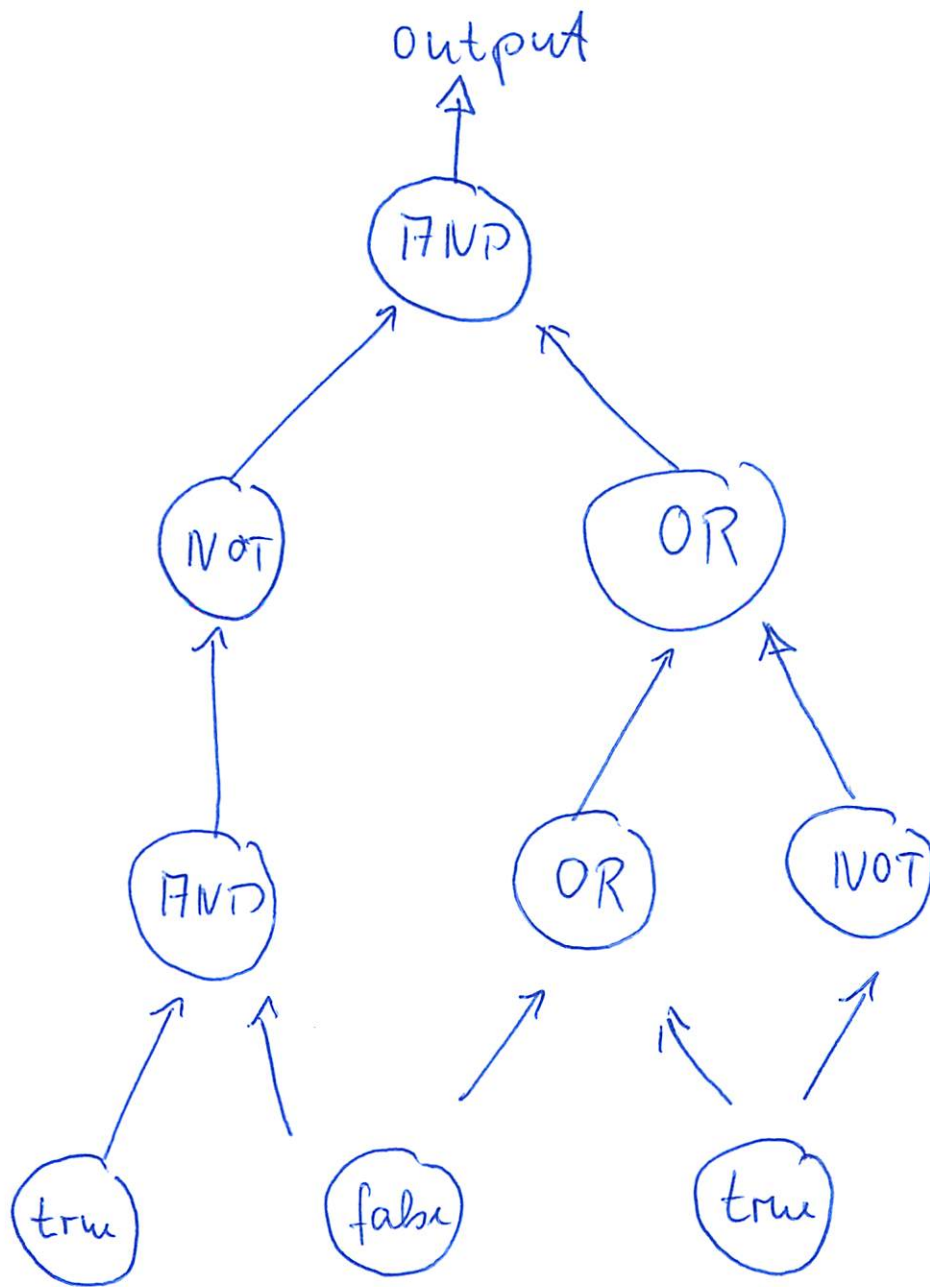
- AND GATES

- OR GATES

- NOT GATES

- One OUTPUT GATE

which is a designated gate



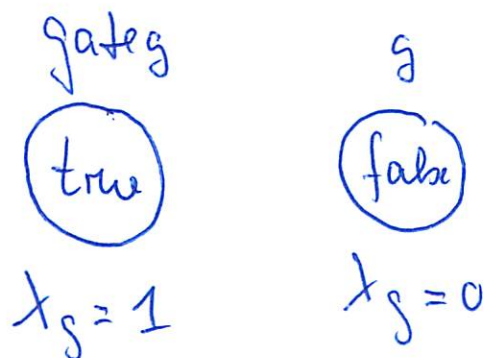
CIRCUIT Value Problem:

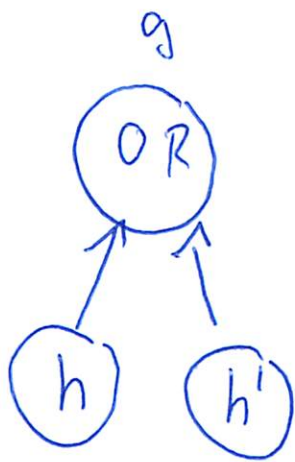
when laws of Boolean logic are applied to the gates in topological order, does the output gate "report" true?

↪ LP

For each gate g , create variable x_g ,
with constraints $0 \leq x_g \leq 1$.

add additional constraints
for each gate according to:

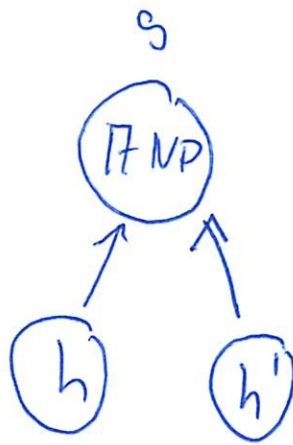




$$x_g \geq x_h$$

$$x_g \geq x_{h'}$$

$$x_g \leq x_h + x_{h'}$$



$$x_s \leq x_h$$

$$x_s \leq x_{h'}$$

$$x_s \geq x_h + x_{h'} - 1$$



$$x_g = 1 - x_h$$

No max or min function required.

The circuit value problem is (in a sense) the most general problem solvable in polynomial time.

Any algorithm will be executed on gates.

If an algorithm runs in polynomial time, take the computer's gates copy them a polynomial times and use the gate values layer by layer.

This together with:
CIRCUIT value reduces to LP

⇒ All problems that can
be solved in polynomial
time do.

In the final chapter of this
course, NP-completeness,
we will see that many
"hard" problems reduce
(in a similar way) to
INTEGER PROGRAMMING.

(your text book calls
integer programming
the difficult twin of LP.

Chapter 8

NP-complete Problems

Towards the biggest open problem
in Computer Science !

We have seen polynomial-time solutions for problems that could easily be solved via an exponential algorithm.

E.g., Dijkstra's shortest path algorithm
min. spanning tree
max increasing subsequence

These are polynomial-time solvable.

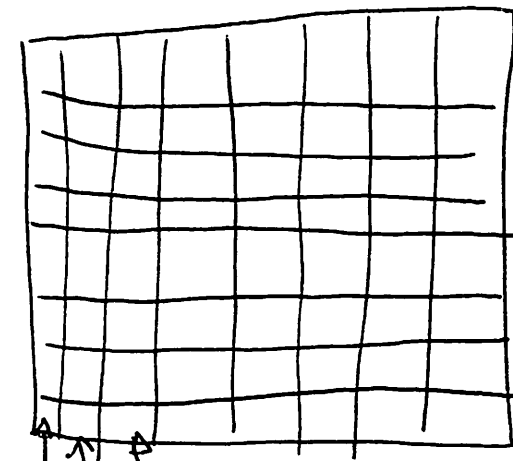
But e.g. - there are an exponential number of paths between a source and a destination node

- an exponential # of spanning trees
- subsequences

Is this always possible?

Is there always a clever technique
like dynamic programming, LP, ...
to get a polynomial-time algorithm
for any computable problem?

How bad is exponential?



chess board
Brahmin Sissa

always double
the previous
#

1 grain 2 grains 4 grains of rice

$$2^0, 2^1, 2^2, \dots, 2^{64-1}$$

$$2^{64-1} = 18,446,744,073,709,551,615$$

quin. billion quad. billion trillion bill. ion million
trillion billion

Thus far, we designed algorithm whose run-time is polynomial, i.e., $O(n^k)$ for some constant k and input size n .

The class of all problems for which a polynomial time algorithm exists is called:

\mathcal{P}

Now, we look at problems for which no polynomial-time algorithm has been discovered as yet.

However, nobody has proven that no such algorithm exists either for these problems.

We first study a particular class of problems.

NP

defined informally as those problems that can be verified in polynomial time.

Verified means that a certificate of a solution can be verified to be correct in polynomial time in the input size to the problem.

(There is a little bit more to the NP-class, but good enough for now.)

Clearly,

$$P \subseteq NP$$

Take a problem in P , solve it in polynomial time even without a certificate.

OPEN: $P \stackrel{?}{=} NP$

biggest CS open problem.

Examples :

P

① Shortest path
(weighted)
in a directed graph
with or without negative
weights.

② Eulerian tour

An Eulerian tour of a
connected, directed graph
 $G=(V, E)$ is a cycle
that traverses each
edge of G exactly once.

(Vertices may be visited
more than once.)

NP

Longest simple path

(even determining
whether a graph
contains a simple
path with at least a
given # of edges is
NP-complete)

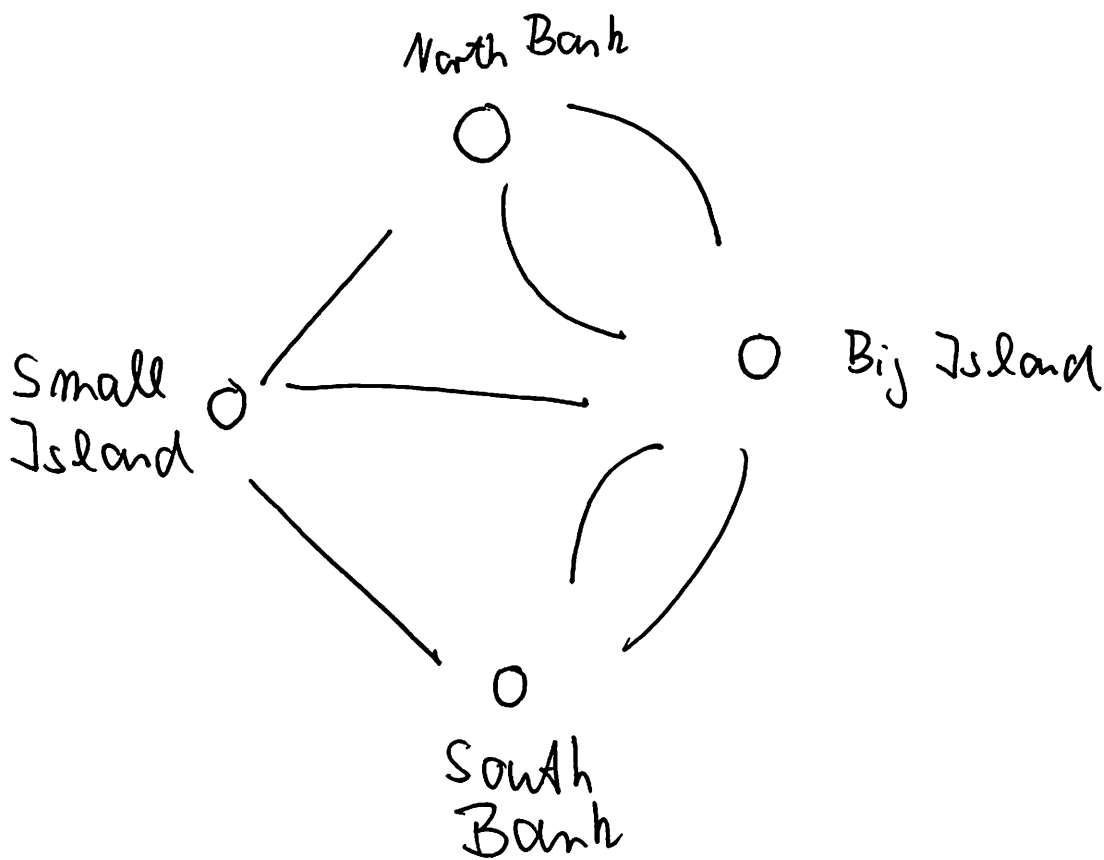
Hamiltonian cycle

(your book: Rudrata cycle)

A Hamiltonian cycle of
a directed graph $G=(V, E)$
is a simple cycle
that contains each
vertex in G .

Leonhard Euler

Swiss Mathematician studied a problem called, Königsberg Bridge Problem, 1735, which lead to Eulerian (or Euler) tours.



edges are bridge | vertices land

Can one find a start location to visit or a tour each bridge exactly once?

Note: the graph has multiple edges
— between nodes (or vertices).

The Euler tour through Königsberg
can be stated as:

Can the graph be drawn without
lifting the pencil from the paper
(and drawing edges twice)

Euler, being a mathematician, solved a more general problem than just for the Königsberg problem instance.

Theorem: There is an Eulerian tour for G

—
 \Leftrightarrow (1) G is connected

(2) every vertex, with the possible exception of two vertices (start and end) has even degree.

In the Königsberg Bridge Problem
all 4 vertices have odd degree \Rightarrow no solution

P

NP

③ Linear Programming

We saw that, although the Simplex Method, was not polynomial-time, there were Polynomial-time Solutions.

④ Minimum Spanning Tree

Find a tree (spanning G) of minimum cost

Integer Linear Programs

here we require the variables of the Linear Program solutions to be integers.

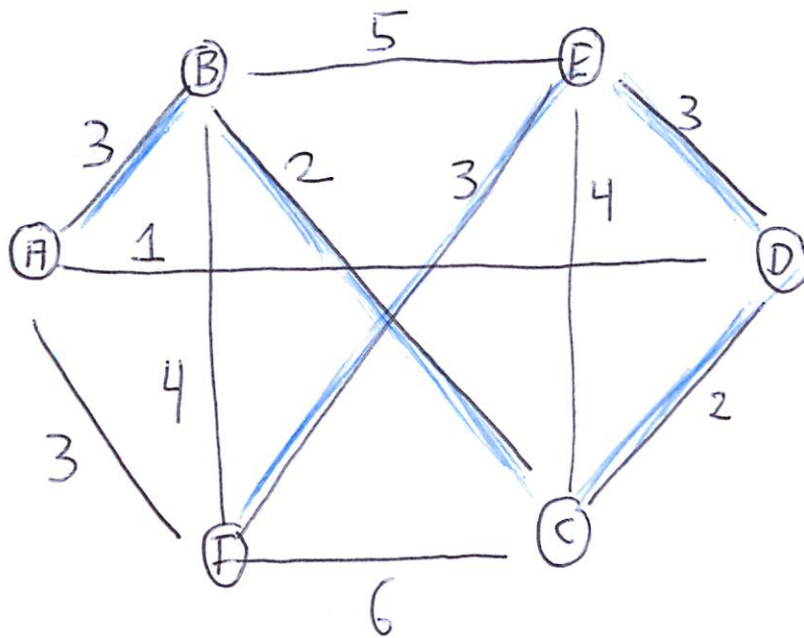
Traveling Salesman

(the problem remains to be called salesman not salesperson, sorry)

Here, we find a special tree, i.e., a tour through every vertex exactly once of cost b , for a Graph in which every pair of vertices has a connected edge of a particular cost.

The cost of the tour is the sum of the individual costs per edge.

Note, we are not asking for the minimum cost. (see later for a discussion on this)



Optimal tow: A, B, C, D, E, F

Cost: 13
(length)

There are $(n-1)!$ tours.

Checking if a tour cost $\leq b$
is clearly polynomial time.

(in fact, it is linear)

So, it is in NP.

A boolean formula is in
Conjunctive Normal Form
 CNF

if it is given as a collection of
 clauses (parentheses), each
 consisting of disjunctions (logical \vee)
 of several literals.

Literals: boolean variables or
 negations of boolean variables.

ex: $(x \vee y \vee z)(\bar{x} \vee \bar{y})(z \vee \bar{x})(\bar{y} \vee \bar{z})(\bar{x} \vee \bar{y} \vee \bar{z})$

\nearrow \uparrow \uparrow \uparrow \uparrow \uparrow \uparrow
 literal or $\underbrace{\hspace{10em}}_{\text{(and } \wedge \text{)}}$ negated

Say $x = \text{false}$ $y = \text{true}$ $z = \text{true}$ then

the expression is false

The question is: Satisfiability

Is there a true/false assignment to the literals so that the boolean formula given in CNF is true?

For this example:

Here, no why?

$$(x \vee \bar{y})(z \vee \bar{x})(y \vee \bar{z})$$

force all three to have the same value

$$\begin{aligned} &\text{either } x=y=z = \text{true} \\ &\text{or } x=y=z = \text{false} \end{aligned}$$

but, then

$$(x \vee y \vee z)(\bar{x} \vee \bar{y} \vee \bar{z}) = \text{false}$$

\Rightarrow the entire formula = false

We call the problem CNF - Satisfiability.

When each clause has exactly k clauses it is called

$\{k\text{-CNF Satisfiability}\}$

For $k=2$, we obtain

2-CNF satisfiability

for $k=3$, we obtain

3-CNF satisfiability.

ex:

$$(x_1 \vee \bar{x}_2)(\bar{x}_1 \vee x_3)(\bar{x}_2 \vee \bar{x}_3)$$

is 2-CNF

$$(x_1 \vee \bar{x}_2 \vee x_3)(\bar{x}_1 \vee x_2 \vee x_3)$$

is 3-CNF

P

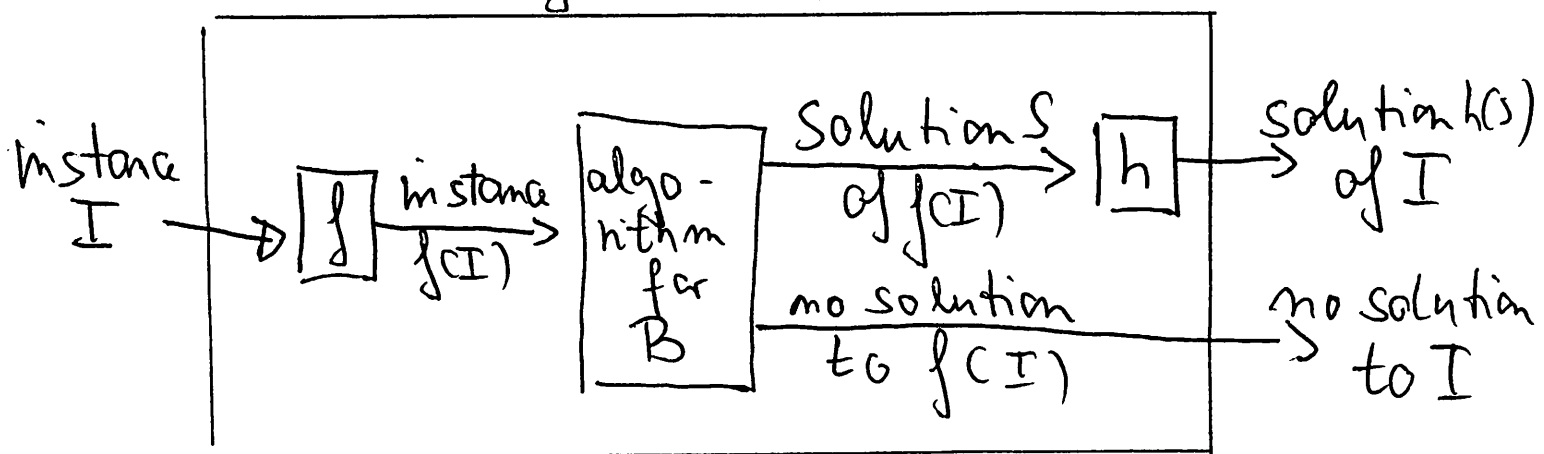
NP

2 - CNF
satu fiabilitas

3 - CNF
satu fiabilitas

A reduction from a search problem A to a search problem B is a polynomial time algorithm that transforms any instance I of A into an instance $f(I)$ of B , together with a polynomial time algorithm h that maps any solution S of $f(I)$ back into a solution $h(S)$ of I .

Algorithm for A



Let us look at 2 problems

Hamiltonian (s,t) -path and
Hamiltonian Cycle problem

Hamiltonian (s,t) -path problem

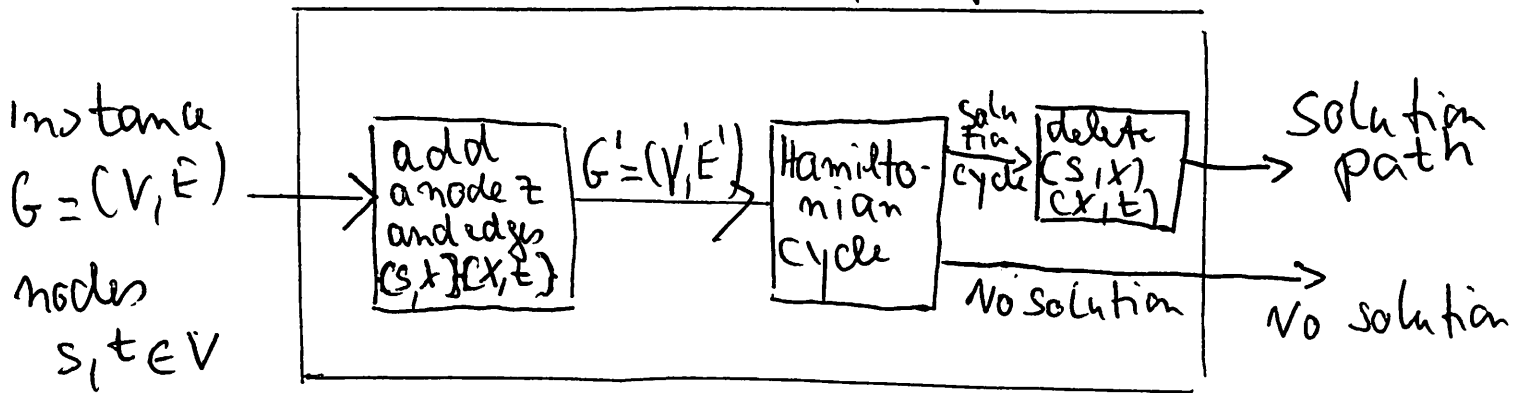
Given a graph and 2 vertices s, t .

Is there a path starting at s
and ending at t that goes
through each vertex exactly once?

Is it possible that Hamiltonian
cycle is easier than Hamiltonian
 (s,t) -path problem?

NO

Hamiltonian (s, t) -path



Case

1. The instance of Hamiltonian Cycle has a Solution

Vertex x has exactly 2 neighbours s and t .

Any Hamiltonian cycle c' in G' must therefore traverse $\{x, s\}$ and $\{x, t\}$

(undirected $\{x, s\} = \{s, x\}$)

The remainder from s to t must traverse all other vertices in G' .

So deleting $\{x, s\}$ and $\{x, t\}$ from the cycle leaves an Hamiltonian path for G connecting s to t .

2. Hamiltonian Cycle does not exist

It is easy to show that in this case the original Hamiltonian Path problem between s and t cannot have a solution either.

Assume it did have a solution.

Then just add $\{x, s\}$ and $\{t, x\}$ and we get a Hamiltonian cycle for G .
A contradiction.

Important!

we need to check that pre and post-processing can be done in polynomial time!

Here, easy as we just add 2 edges or delete 2 edges.

Reduction from A to B

$A \rightarrow B$

A search problem is NP-complete if all other search problems reduce to it.

These problems are in some sense the hardest problems in NP.

Assume A reduces to B

$$A \rightarrow B$$

"the difficulty" flows in the direction of the arrow.

While

"efficient algorithms" move the other way.

if $A \rightarrow B$ and $B \rightarrow C$

then $A \rightarrow C$

"Composition"
of the input of outputs functions

Fact: Once we know that a problem, A , is NP-Complete, we can use it to prove that another search problem, B , is also NP-Complete by reducing A to B .

Follows from composition.

Formal definitions: (p.244 your book)

P : set of ^{search} problems that have a polynomial time solution.

NP : Set of search problems that can be

- (1) found
- and
- (2) verified

in polynomial time by a non-deterministic (sort of) algorithm.

[Guesses correctly at every time.]

I do not like the definition. Original def. via formal languages, as decision problems $w \in L$ or $w \notin L$, $\forall L$ some language. 829

Decision problems vs. optimization problems

In optimization problems, feasible solutions have values and we want to determine the optimal value.

In decision problems, answers are yes or no.

Usually, we can state an optimization problem as a decision problem.

E.g., is there a shortest path with at most k edges, between s and t .

(k here is an integer)

(Q: Find shortest path with fewest edges.)

If a decision problem is hard
its related optimization problem
is then hard too.

Reductions cont'a

3SAT \rightarrow Independent Set

\uparrow

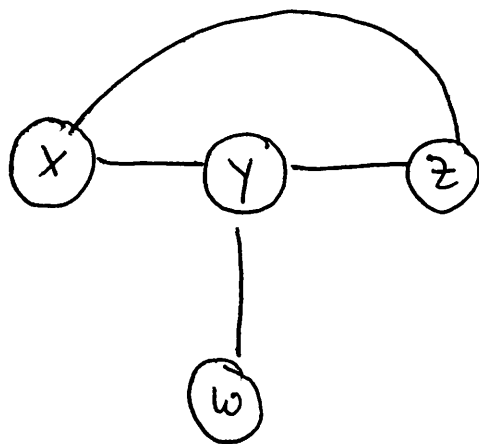
$(\bar{x} \vee y \vee \bar{z})(x \vee \bar{y} \vee z)(x \vee y \vee z)(\bar{x} \vee \bar{y})$

Independent Set Problem:

input: a graph G and a number g

output .. find a set of g pairwise non-adjacent vertices in G or report that no such set exists

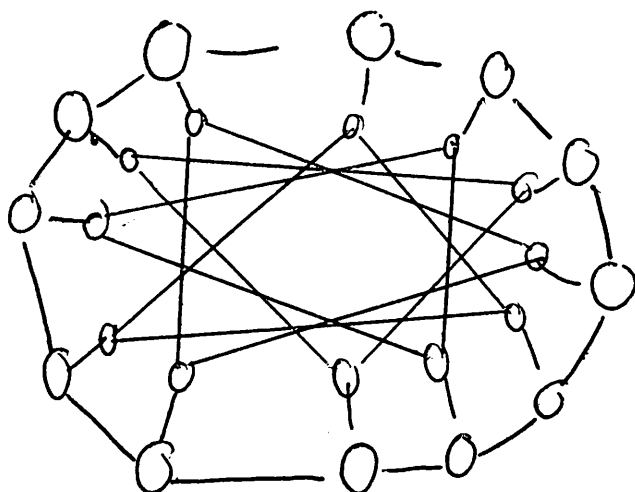
Ex:



$g=2$

indep. set
 $\{x, w\}$

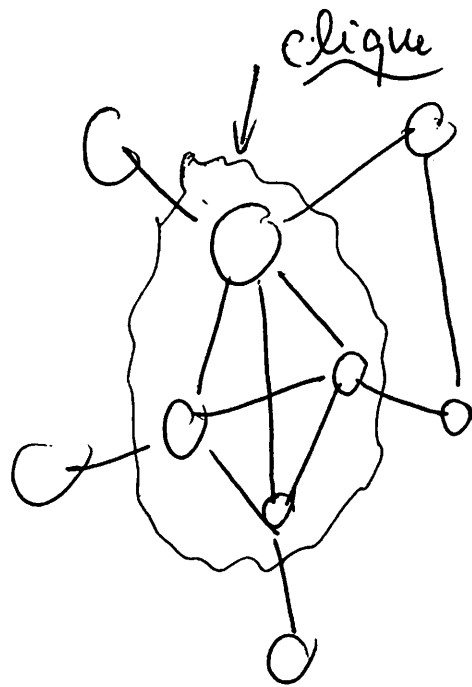
The optimization problem corresponding to this is maximum independent set



What is the size of the maximum independent set? 9

Find them!

A clique in a graph G is a subset of G 's vertices in which every two vertices are connected by an edge in G .



A set in G is independent \Leftrightarrow
it forms a clique in the
Complement of G .

Complement : if $(x,y) \in E$ then $(x,y) \notin E^c$
if $(x,y) \notin E$ then $(x,y) \in E^c$

$G = (V, E)$ $G^c = (V^c, E^c)$

Sufficiently large graphs
with no large cliques have
large independent sets
(\rightarrow Ramsey theory)

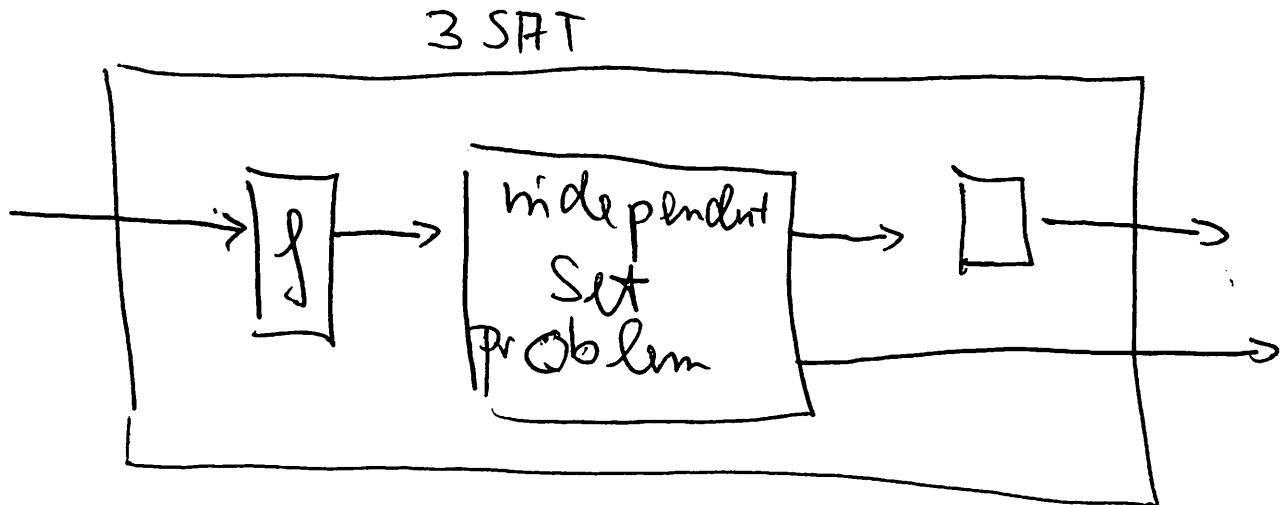
Cliques are useful as
they have very strong (optimal,
connectivity).

back to the reduction

3 SAT \rightarrow Independent Set

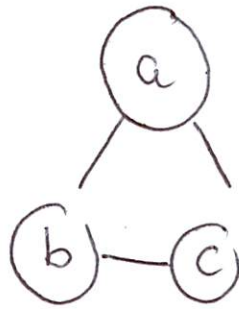
So, we need to find f

f . Instance I into an instance $f(I)$ for
(3-SAT) independent set



Construction: \boxed{f}

For each clause: $(a \vee b \vee c)$
construct a triangle



vertices: a, b, c

edges: $a-b$
 $a-c$
 $b-c$

(a, b, c can be negated say $b = \bar{y}$)

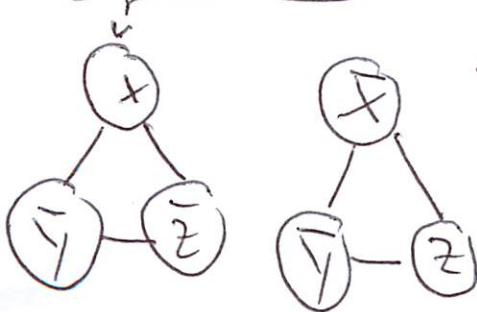
For each literal: x

connect $(x) - (\bar{x})$ in all occurrences

\Rightarrow The construction takes polynomial time.

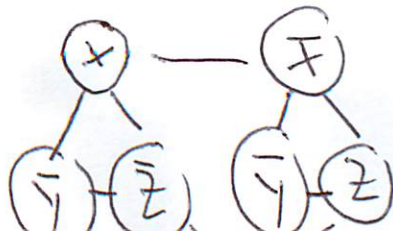
Ex:

$(x \vee \bar{y} \vee \bar{z})$ $(\bar{x} \vee \bar{y} \vee z)$



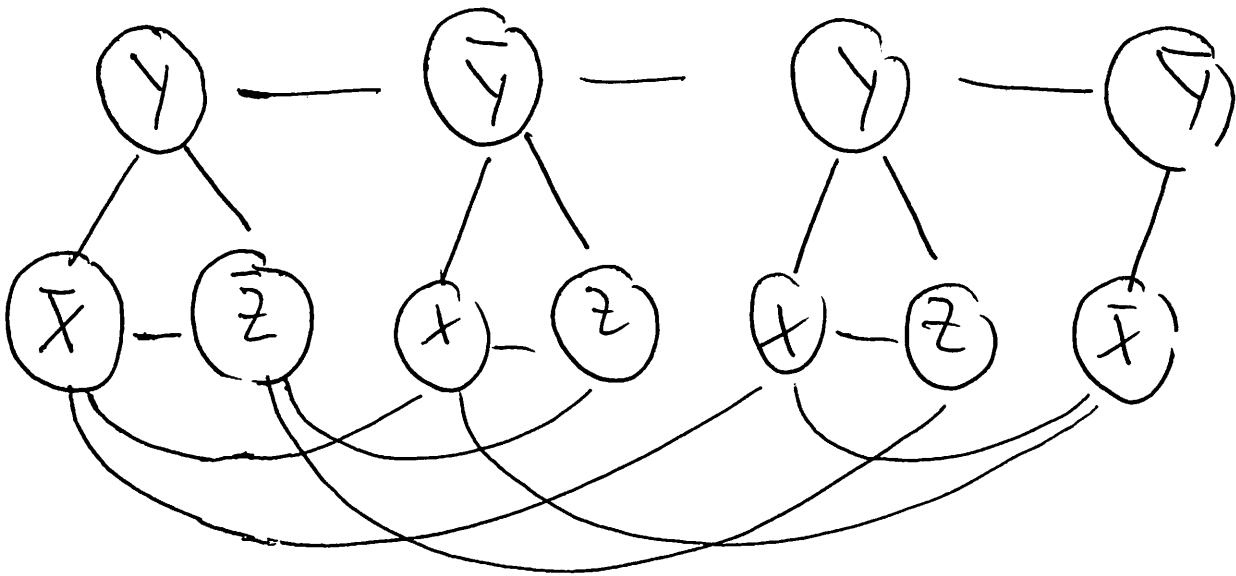
now add

$x - \bar{x}$
 $y - \bar{y}$
 $z - \bar{z}$



Ex:

$$(\bar{x} \vee y \vee \bar{z})(x \vee \bar{y} \vee z)(x \vee y \vee z)(\bar{x} \vee \bar{y} \vee z)$$



(Note, your book allows clauses with 2-literals)

The construction is then simply
an edge $(\bar{y} \vee \bar{x})$



The goal $g = \#$ of clauses

in our example $g = 4$.

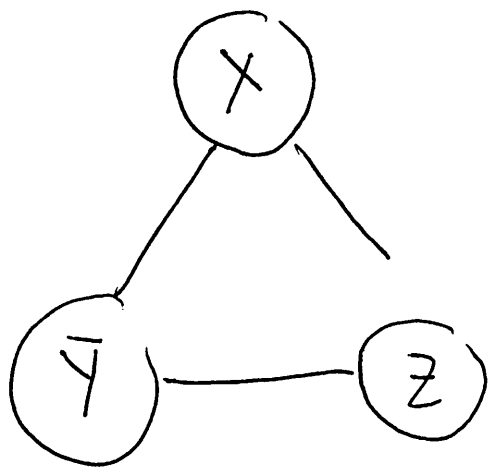
The "output", i.e. second conversion function h .

Because in G $(x) - (\bar{x})$
for any literal x , and any
occurrence of x, \bar{x} .

\Rightarrow either x or \bar{x} is in the
independent set, S , not both.

h : assign: $x = \text{true}$ if $x \in S$
 $x = \text{false}$ if $\bar{x} \in S$

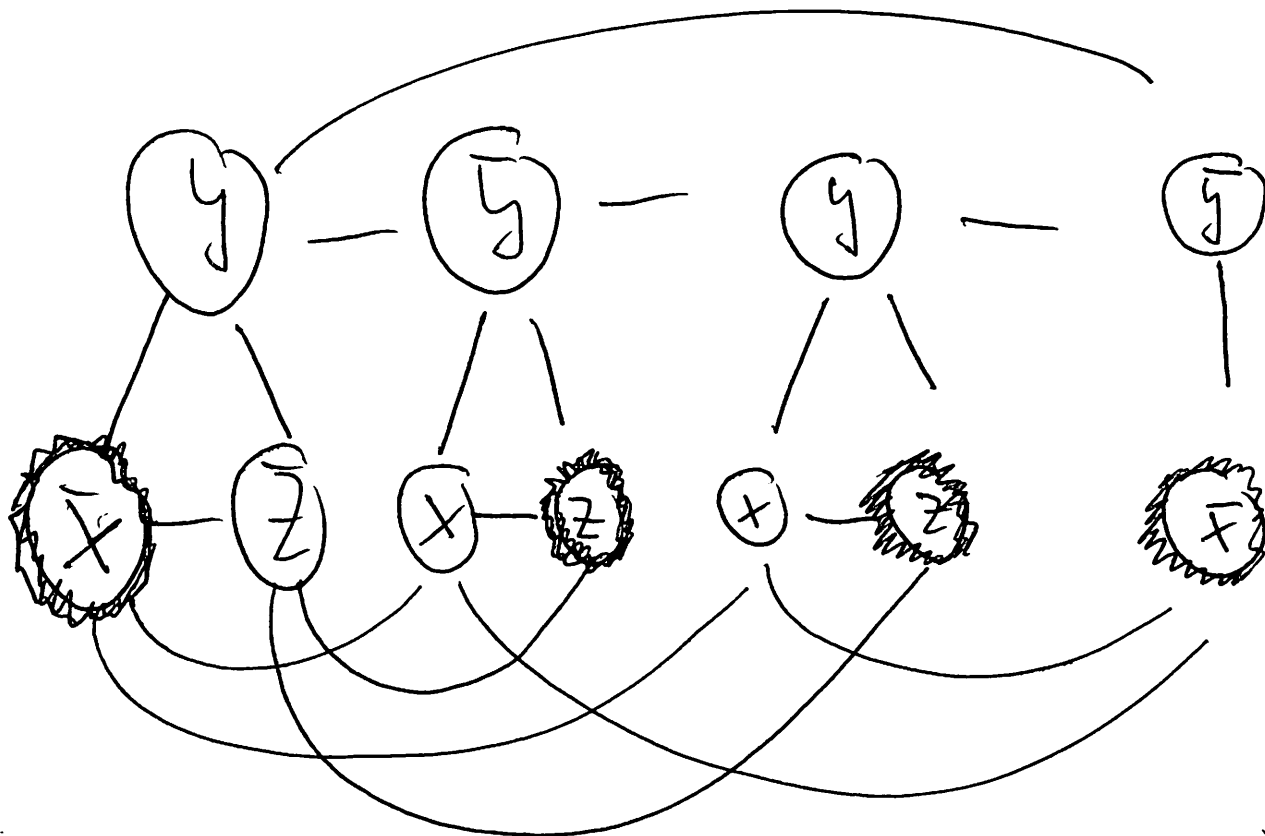
Why does this work?



in an independent set, we can select at most one vertex per triangle. But, since $g = \# \text{ of clauses} = \# \text{ triangles}$ we must select one per triangle.

We cannot choose both x and \bar{x} .

Choose the vertices as mid.
 Set as indicated



in each triangle (or \square) exactly
 one vertex is chosen.
 No literal can be chosen as both
 negated and unnegated.

chose \bar{x} : \Rightarrow $x = \text{false}$
 z \Rightarrow $z = \text{true}$
 y arbitrary

Fact If Graph G has no ind. set of size g , then the Boolean formula I is unsatisfiable.

Proof: Assume the contrary, i.e. I is satisfiable. Then, for each clause pick any literal whose value under the satisfying assignment is true.

Clearly, in each clause there is one.

They must be independent set. because we never pick a literal and its negation. \square

Independent Set \rightarrow Clique

We observed already

Lemma : A set of nodes S is
an independent set of G
if and only if S is a clique of
 G^c . (G^c is the complement
Graph of G)

Thus, the above reduction
is easy.

SAT \rightarrow 3-SAT

This is a reduction from a general problem to a special case of itself. So, the problem is just as hard.

How do we shorten clauses with more than 3 literals?

Here is the "trick":

$$\left. \begin{array}{l} (a_1 \vee a_2 \vee a_3 \vee \dots \vee a_k) \\ \text{is satisfied} \end{array} \right\} \Rightarrow \text{there is a setting} \\ \text{of the } y_i\text{'s for which} \\ (a_1 \vee y_1)(\bar{y}_1 \vee a_2 \vee y_2) \dots (\bar{y}_{k-1} \vee a_k) \\ \text{are all satisfied} \quad k-3$$

Why is this true?

(1) Assume all clauses on right are satisfied

Claim: at least one of a_1, \dots, a_n is true.

Assume otherwise. Then

$$(a_1 \vee a_2 \vee \gamma_1) = \text{false} \text{ forces } \gamma_1 = \text{true}$$

$$\begin{array}{ccc} (\overline{\gamma_1} \vee a_3 \vee \gamma_2) = \text{false} & \text{forces} & \gamma_2 = \text{true} \\ \uparrow & & \uparrow \\ \text{false} & & \text{false} \end{array}$$

\vdots

$$\gamma_{k-3} = \text{true}$$

$$(\overline{\gamma_{k-3}} \vee a_{k-1} \vee a_n) = \text{false} \quad \Downarrow$$

Thus claim holds. \Rightarrow one of a_1, \dots, a_n is true $\Rightarrow (a_1 \vee \dots \vee a_n) = \text{true}$.

(2) $\exists (a_1 \vee a_2 \vee \dots \vee a_n)$ is satisfied \Rightarrow one of

$a_1, \dots, a_n = \text{true}$, say $a_i = \text{true}$

Then set $\gamma_1, \dots, \gamma_{i-2} = \text{true}$

$\gamma_{i-1}, \dots, \gamma_{k-3} = \text{false}$

$$(a_1 \vee a_2 \vee \underset{\uparrow t}{y_1}) = t$$

⋮

$$(\bar{y}_{i-3} \vee a_{i-1} \vee \underset{\uparrow t}{y_{i-2}}) = t$$

$$(\bar{y}_{i-2} \vee \underset{\uparrow t}{a_i} \vee y_{i-1}) = t$$

$$(\bar{y}_{i-1} \vee a_{i+1} \vee \underset{\uparrow t}{y_i}) = t$$

⋮

$$(\bar{y}_{k-3} \vee a_{k-1} \vee a_k) = t$$

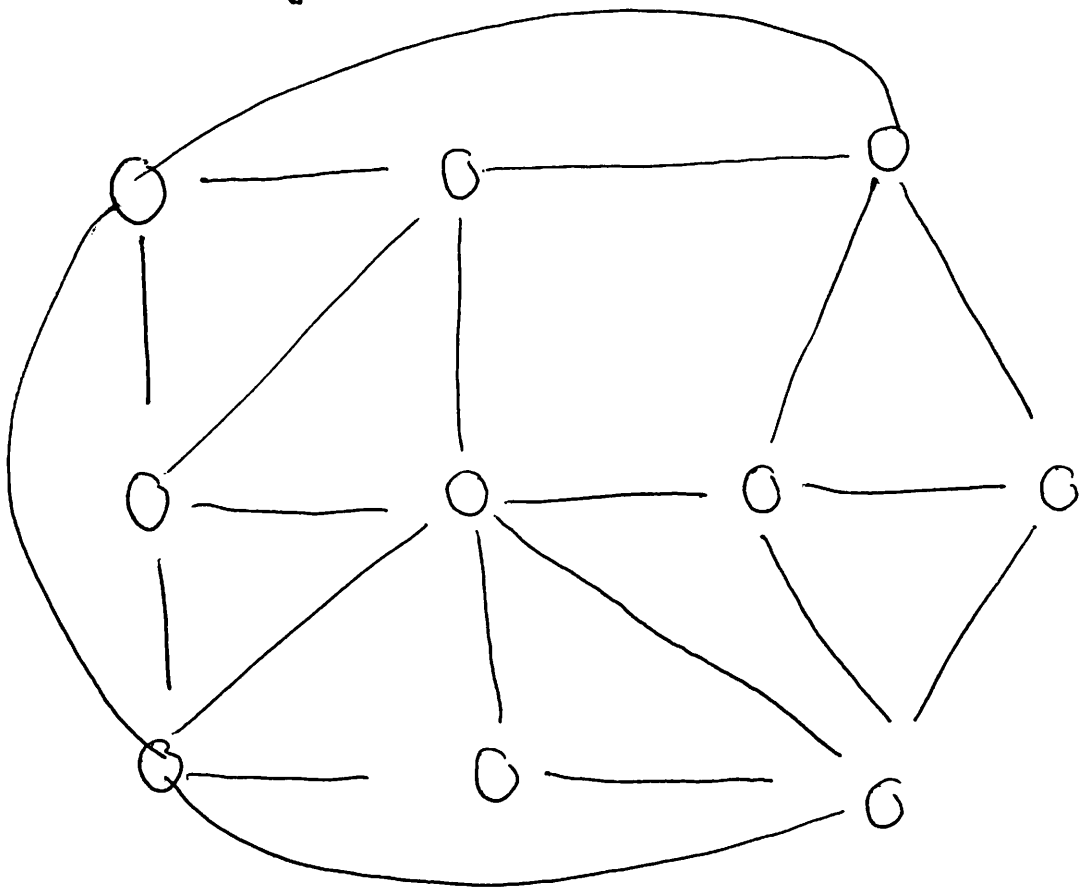
So what are f and h ?

Is it polynomial time for f, h ?

Vertex Cover Problem

Input: graph and budget b

Task: Find b vertices that touch every edge.

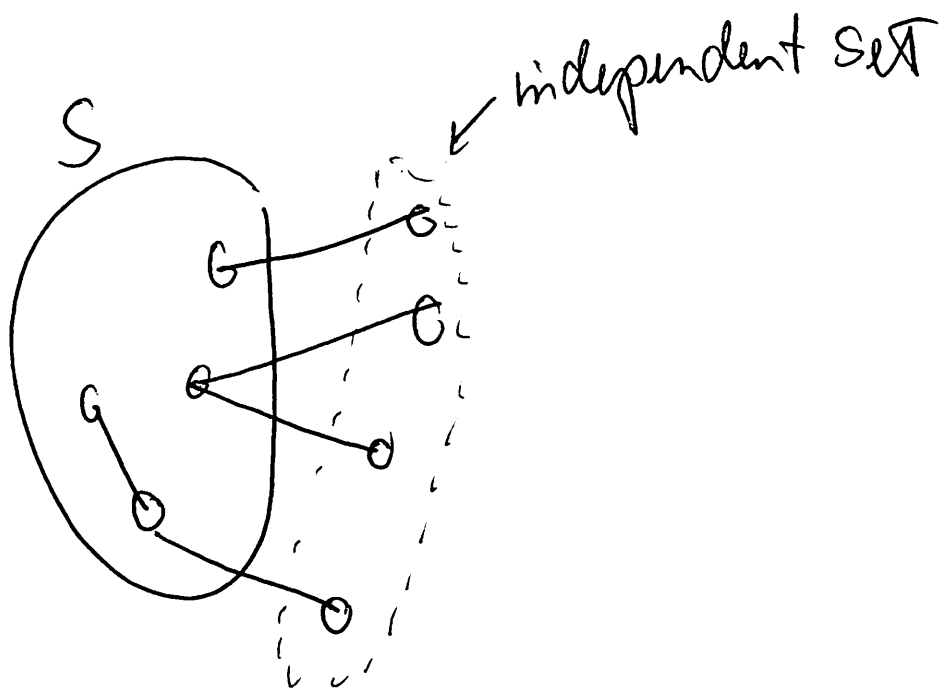


Independent Set \rightarrow Vertex Cover

Observe,

S is a vertex cover of G (\Leftrightarrow)

$V - S$ is an independent set of G .



To solve the independent set problem for (G, g) , look for a vertex cover of G with $|V| - g$ nodes.

If such a cover exists \Rightarrow take all nodes NOT in it.

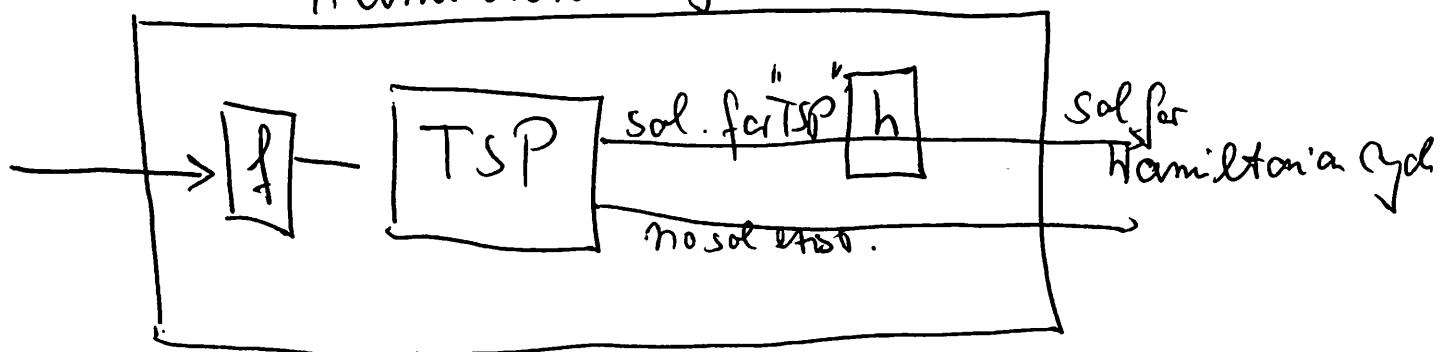
Otherwise, G cannot have an independent set of size g .

Reductions cont'd

Hamiltonian Cycle \rightarrow TSP

Given a graph $G=(V, E)$ construct
An instance of TSP:

Hamiltonian Cycle



f : we construct a special TSP
 $V \rightarrow$ vertices of the TSP

distances: $(u, v) \in E \Rightarrow \text{dist}(u, v) = 1$
 $(u, v) \notin E \Rightarrow \text{dist}(u, v) = 1 + \alpha$
for some $\alpha \geq 1$
to be determined

budget for TSP := $|V| (M = n)$

Note: \exists G has a Hamiltonian cycle
then the same cycle can be
made a tour within the budget.

\exists G has no Hamiltonian cycle
then there is no solution.

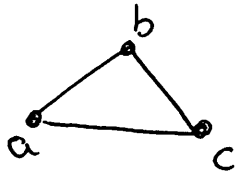
Cheapest TSP cost $\geq n + \alpha$

(at least one edge not in E
would have to be used, its
cost is $1 + \alpha$)

Why $\alpha \geq 2$? Interesting special cases!

$$\boxed{\alpha = 1} \Rightarrow \text{dist}(u, v) = \begin{cases} 1 & (u, v) \in E \\ 2 & (u, v) \notin E \end{cases}$$

in this case, the distances of TSP
satisfy the triangle inequality



$$d(a, c) \leq d(a, b) + d(b, c)$$

This is the case for Euclidean distances

Why satisfied here? if $1 \leq i, j, k \leq 2$
then $i + j \geq k$

TSP can in this case be well
approximated.

α large \Rightarrow TSP may not satisfy
the triangle inequality

Then: either there is a solution
of cost n (or less)
or all its solutions have cost
at least $n + \alpha$
but α can be arbitrarily large.
Compared to n

n $\xrightarrow{\text{huge gap}}$ $n + \alpha$

This gap property implies
(not shown here) that
unless $P = NP$, no approximation
for TSP can exist.

ZOE - Problem

Let A be an $m \times n$ matrix

$$A[i, j] = \begin{cases} 0 \\ 1 \end{cases} \text{ or}$$

Find a $\{0,1\}$ -vector $x = (x_1, \dots, x_n)$ so that

$$Ax = \mathbf{1} \quad \mathbf{1} = \begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix}$$

Ex: $A = \begin{bmatrix} 1 & 0 & 1 \\ 1 & 1 & 1 \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \\ x_3 \end{bmatrix}$ 3 equations $\begin{matrix} x_1 + x_3 = 1 \\ x_1 + x_2 + x_3 = 1 \\ x_3 = 1 \end{matrix}$

Many combinatorial problems can be expressed as a ZOE.

E.g., matching problems.

We will show

ZOE \rightarrow (Rudrata Cycle)
Hamiltonian Cycle

The reduction goes in 2 stages

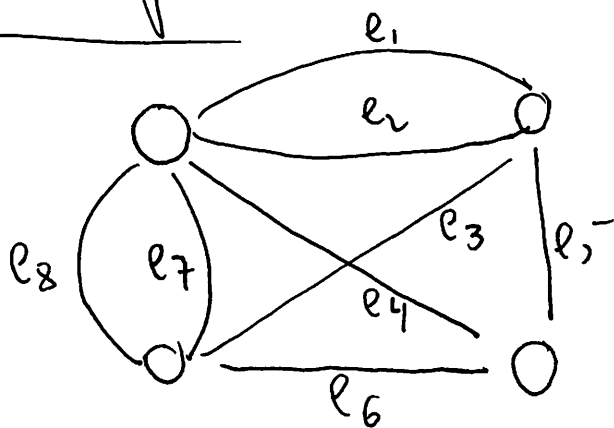
① $ZOE \rightarrow$ special Hamiltonian Cycle problem

② remove "special"

① $ZOE \rightarrow$ special Hamiltonian Cycle
Pair-edges

Paired-edges Hamiltonian Cycle

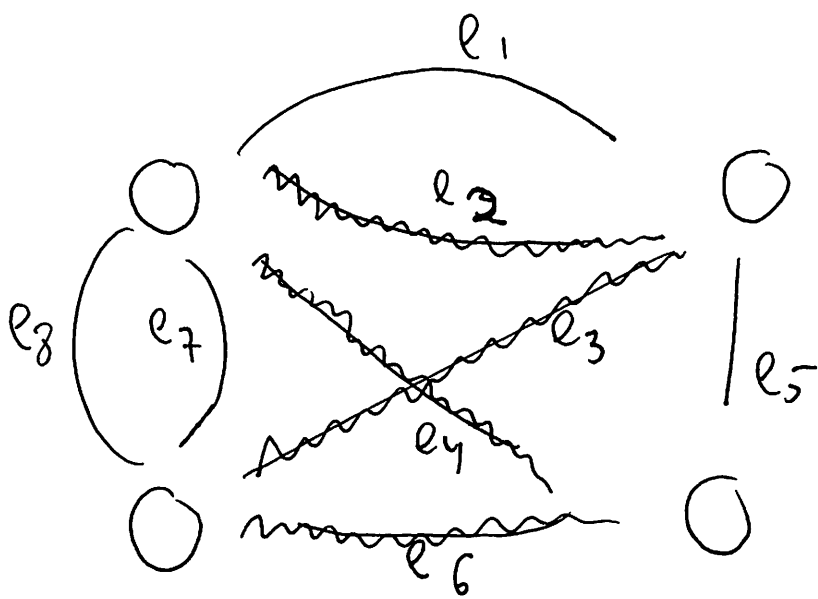
Example: $G = (V, E)$



$$C \subseteq E \times E$$

$$C = \{(e_1, e_3), (e_5, e_6), (e_4, e_5), (e_3, e_7), (e_3, e_8)\}$$

for each pair (e_i, e_j) exactly one edge is traversed by the cycle (visiting all nodes)



$(e_1, e_3) \in C$

So, if we pick e_1 we cannot pick e_3 ,
but we must pick one of e_1, e_3

e_2, e_3, e_6, e_4 is a pair-edge
Hamiltonian cycle in G .

e_2 is not paired

e_3 is paired with $\underbrace{e_1, e_7 \text{ and } e_8}_{\text{they are not on the cycle}}$

e_4 is paired with e_5
↑ not on the cycle

e_6 " " " " " " " "

w.r.t. the vertices we have a cycle.

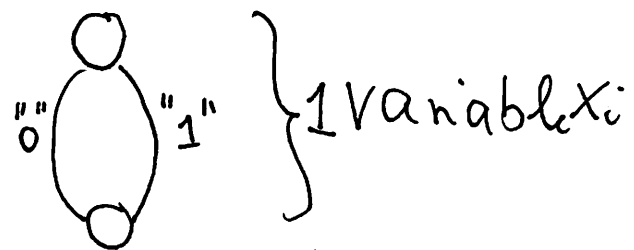
ZOE

instance is a matrix A ($n \times n$)
with 0,1 entries

\Rightarrow n equations in n 0-1 variables

We need to create an input instance
for paired Hamiltonian cycle problem.
2 STEPS

STEP 1: For each variable x_i
Create 2 edges



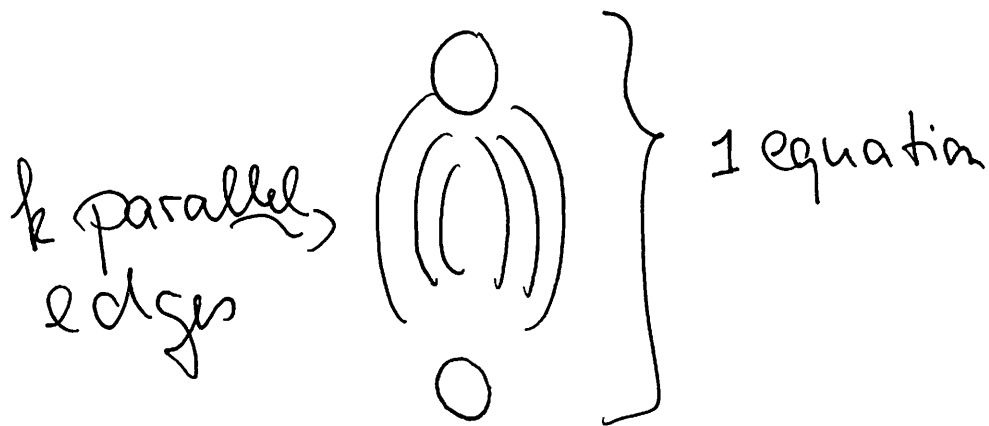
Corresponding to

$x_i = 0$ and $x_i = 1$

For each equation

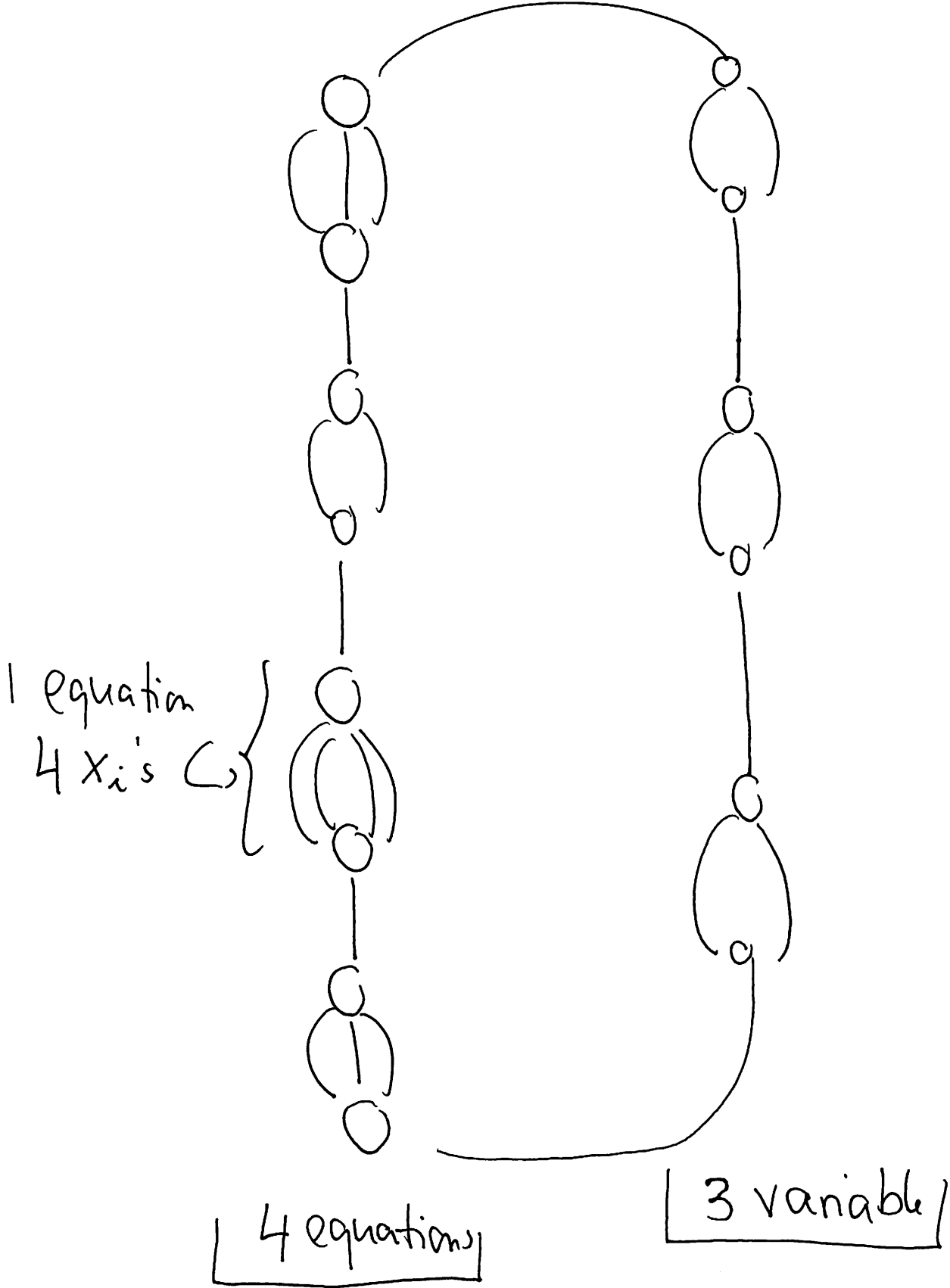
$$x_{j_1} + \dots + x_{j_k} = 1$$

if this involves k variables
Create

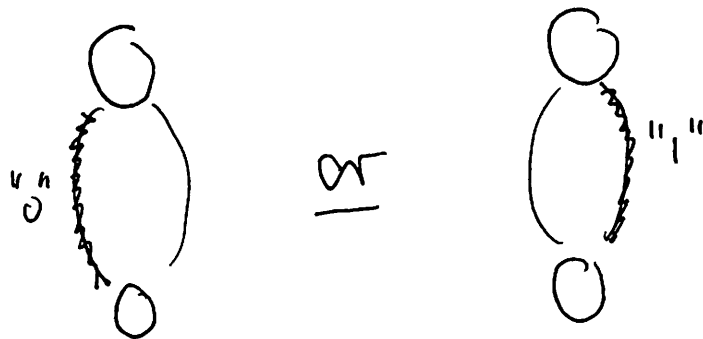
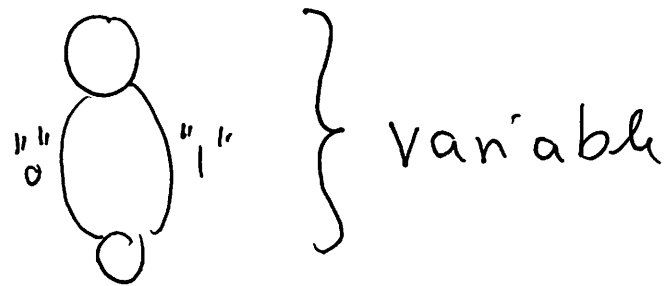


Balana. STEP 1

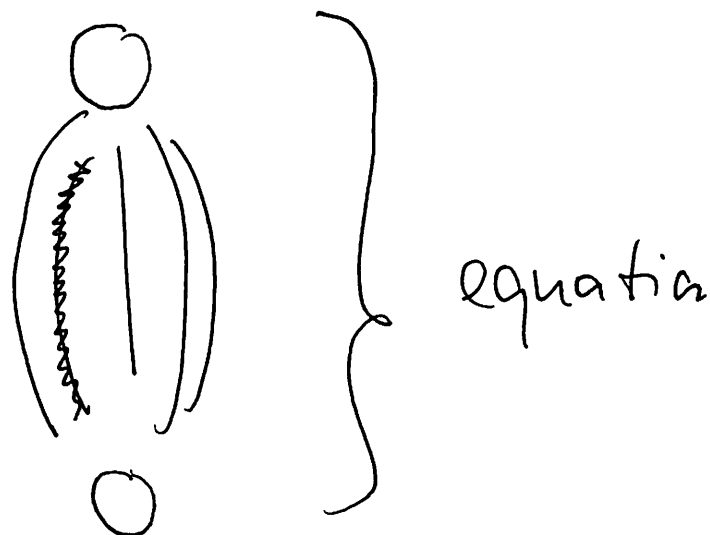
we created a graph
with $m + m$ parallel edges



In any Hamiltonian cycle solution the solution uses one of $x_i = \begin{cases} 0 \\ 1 \end{cases}$ or not both



and picks one variable per equation



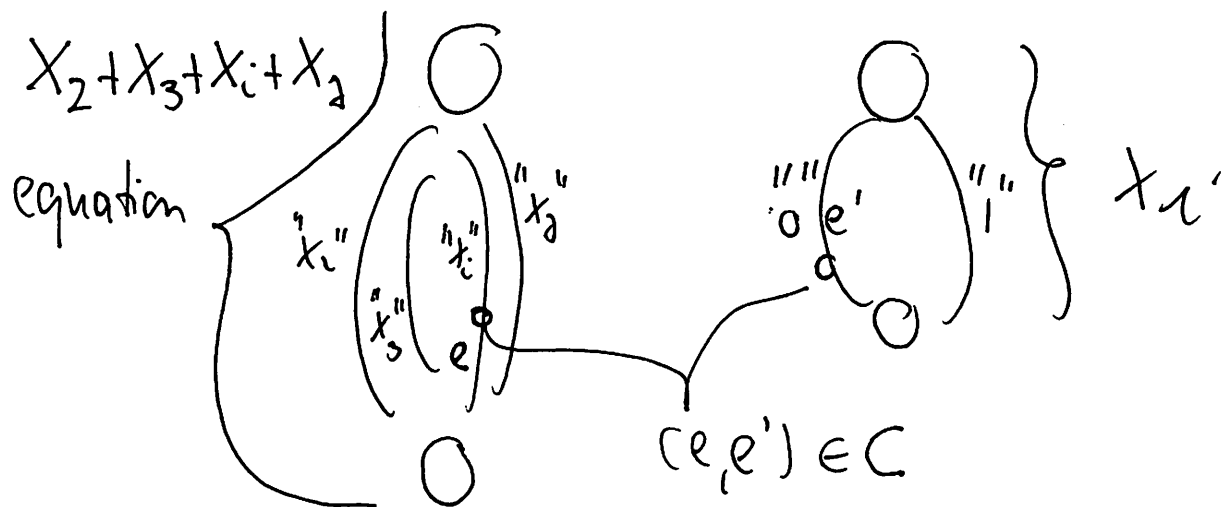
STEP 2:

For every equation
and every variable x_i
appearing in the equation

add (e, e') to C

where e is edge corresponding
to x_i in the equation

and e' is the edge corresponding
to $x_i = 0$



What does this do?

if the cycle picks/uses the edge corresponding to x_i

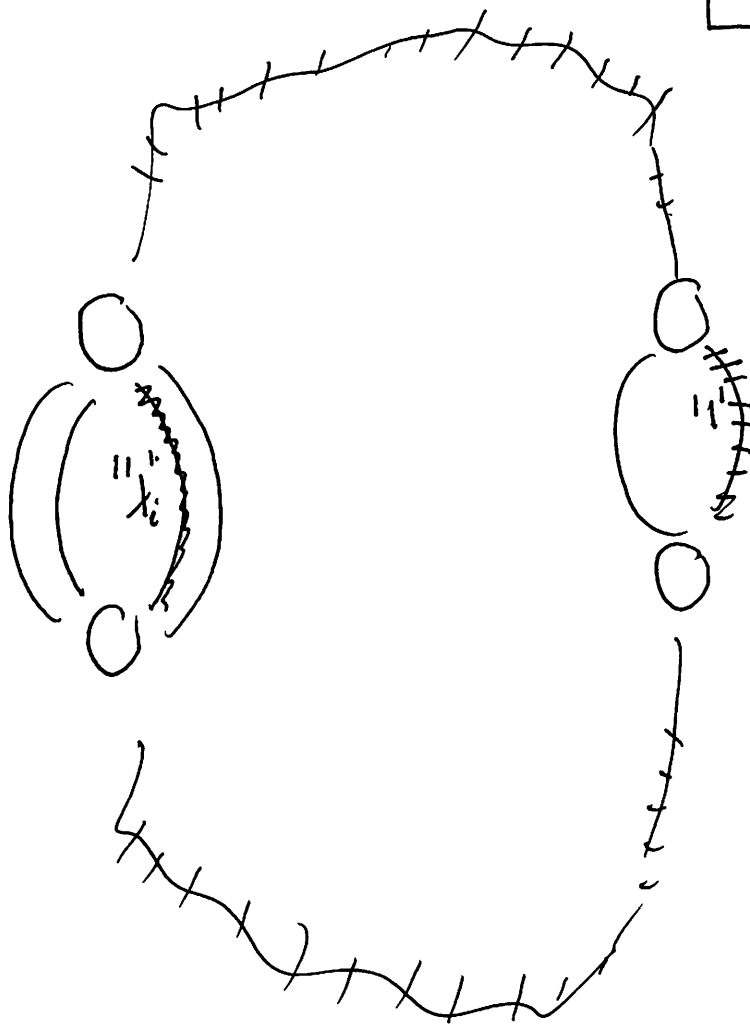
it cannot pick (" $x_i = 0$ ")

\Rightarrow " $x_i = 1$ "

is picked

but then in every equation x_i is picked. (if it appears)

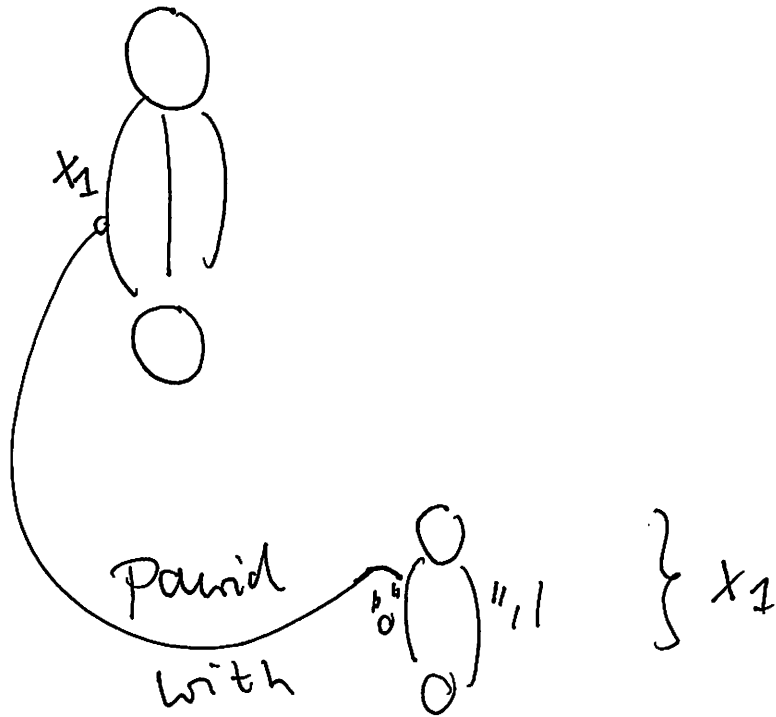
because they are paired with " $x_i = 0$ ".



On the "left", the equation side, always one edge is picked (i.e., is on the cycle).

For this variable x_i the pairedness ensures that " $x_i = 1$ " on the variable side. and " $x_i = 1$ " is then ensuring that x_i is always chosen

Every where



Example:

$$A = \begin{bmatrix} 1 & 1 & 1 & 0 \\ 1 & 0 & 1 & 1 \\ 1 & 1 & 0 & 0 \\ 1 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \\ x_3 \\ x_4 \end{bmatrix} = \mathbf{1}$$

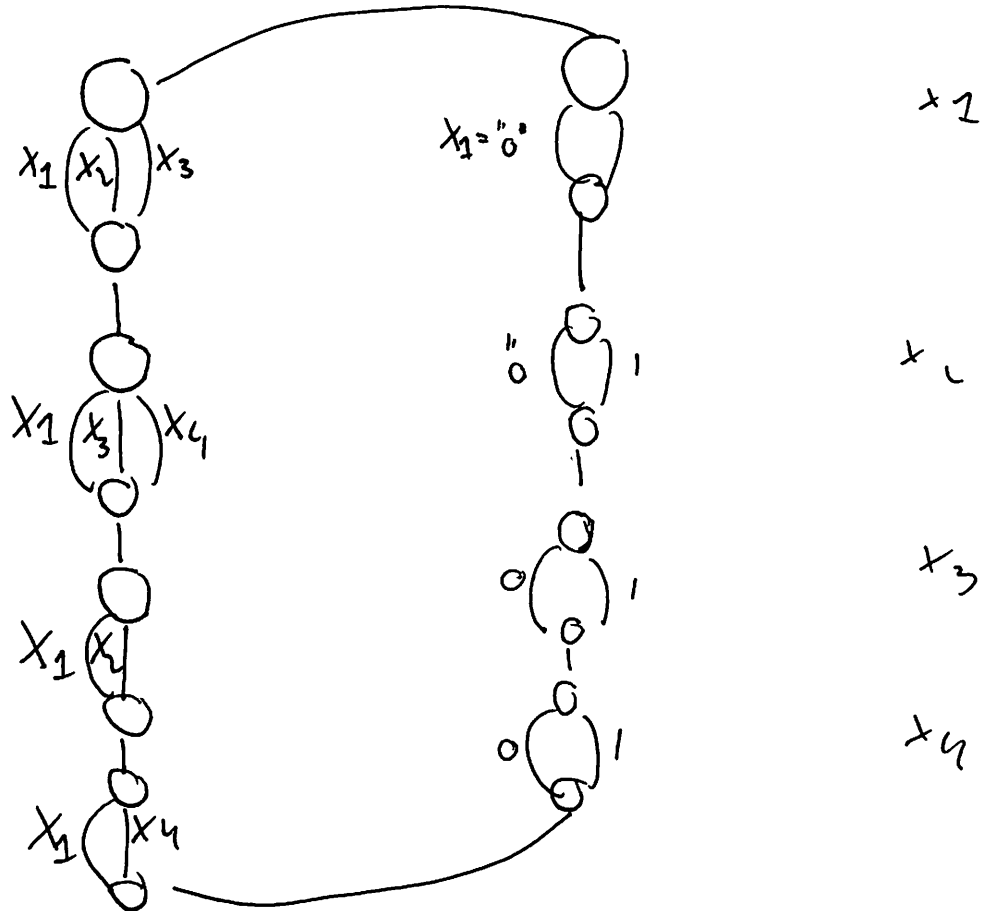
Equation

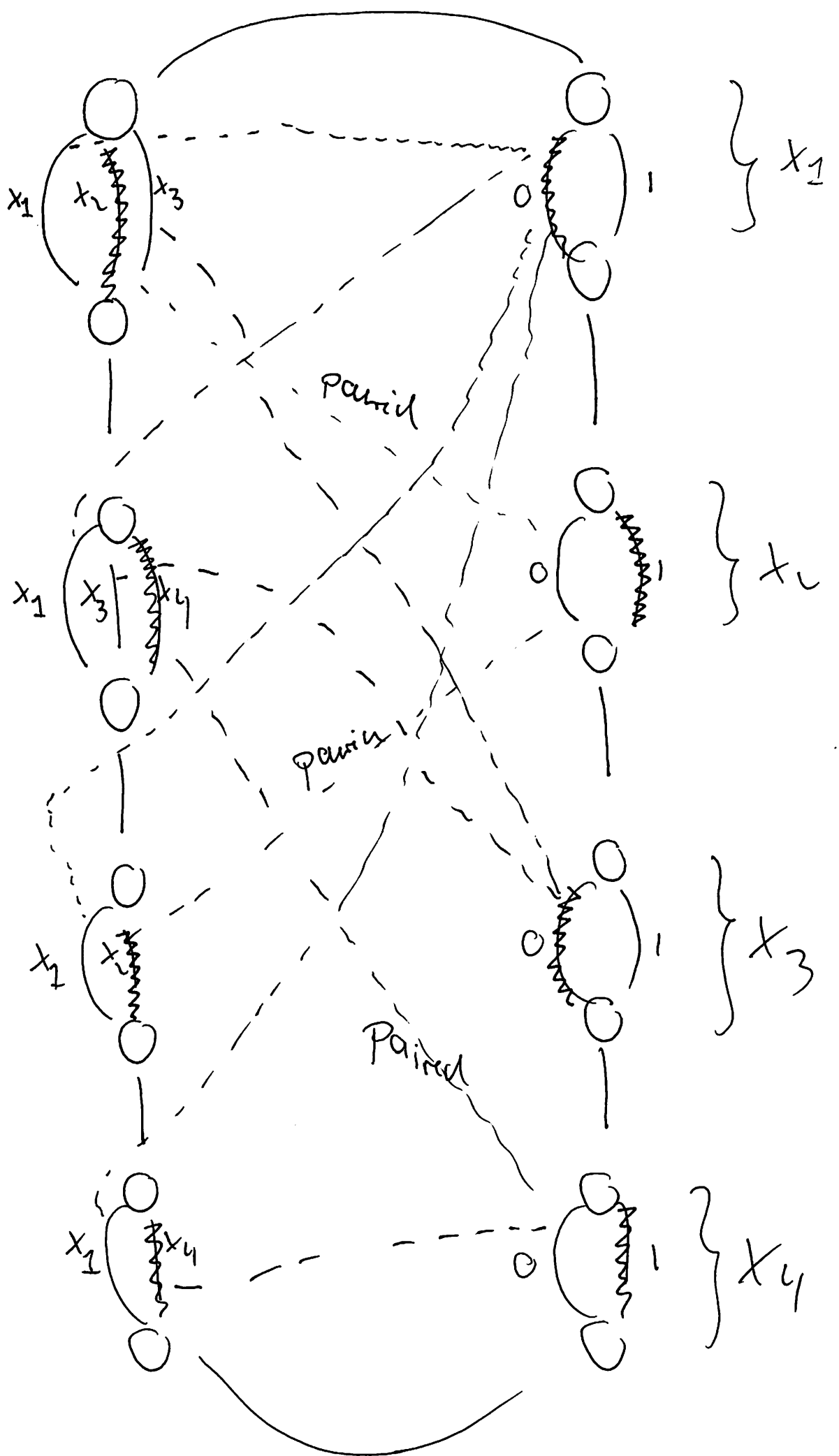
$$x_1 + x_2 + x_3 = 1$$

$$x_1 + x_3 + x_4 = 1$$

$$x_1 + x_2 = 1$$

$$x_1 + x_4 = 1$$





$$x_2 = 1$$

$$x_4 = 1$$

$$x_1 = 0$$

$$x_3 = 0$$

paired Ham.

Cycle

\Rightarrow

correct solution to

$$\overline{A}x = \underline{1}$$

Also, correct solution to $\overline{A}x = \underline{1}$

gives a solution to the paired Hamiltonian cycle problem.

② We wanted

$ZOE \rightarrow$ Hamiltonian
Cycle

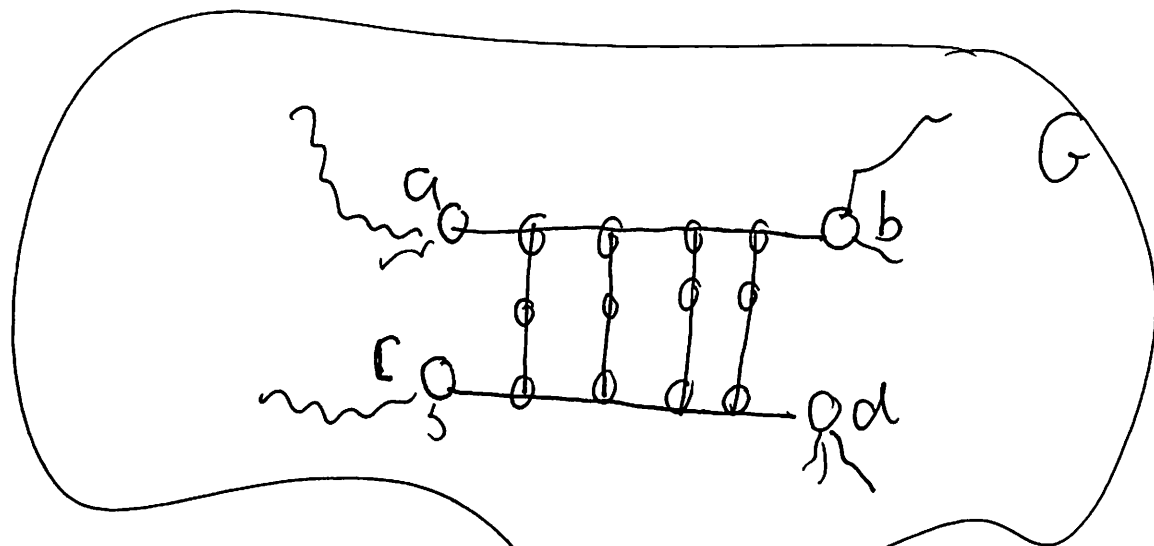
So far, we have

$ZOE \rightarrow$ Paired-edges
Hamiltonian
Cycle.

Hamiltonian cycle is a
special case of paired-
edges version for $C = \emptyset$.

But, this is not good enough
(wrong special case)
Way ↓

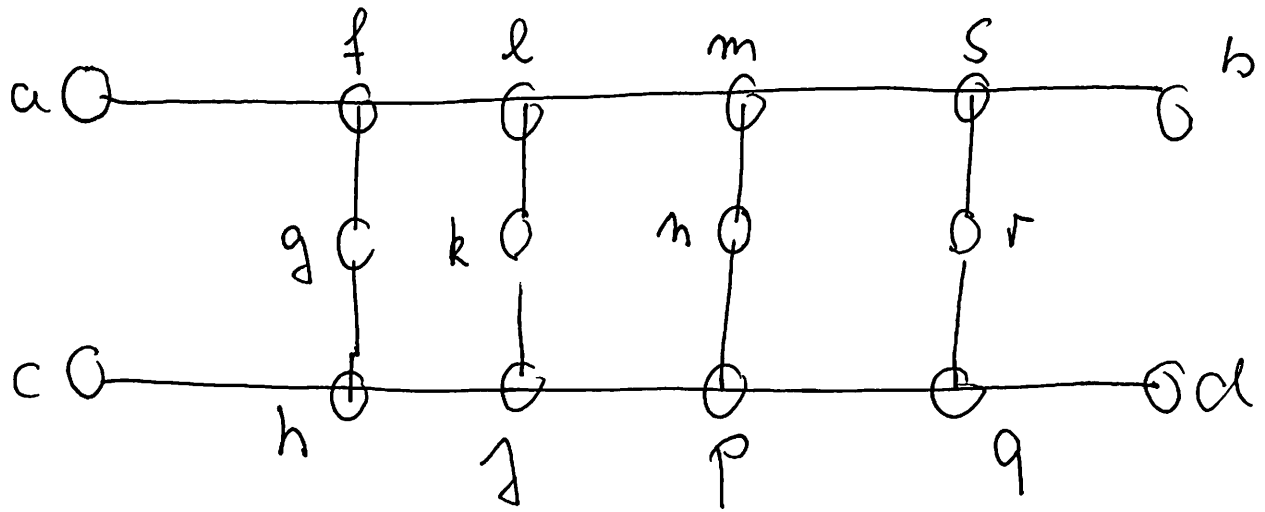
A gadget for enforcing
paired behavior



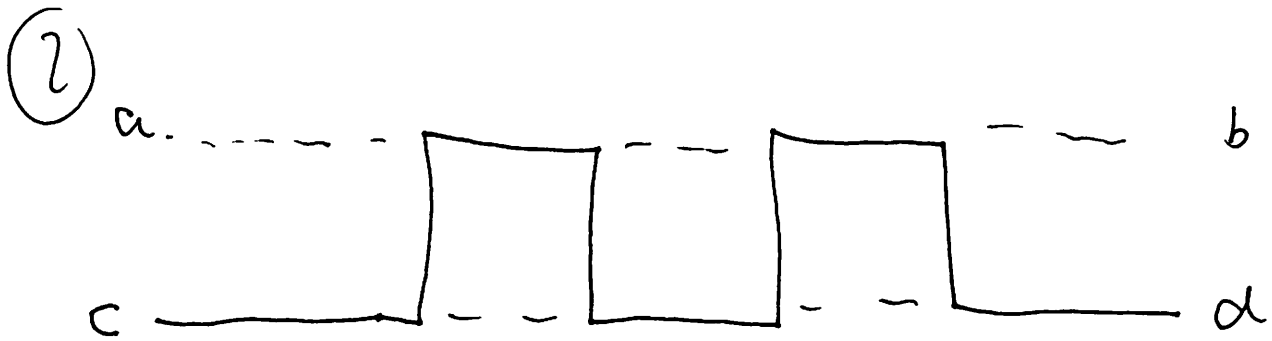
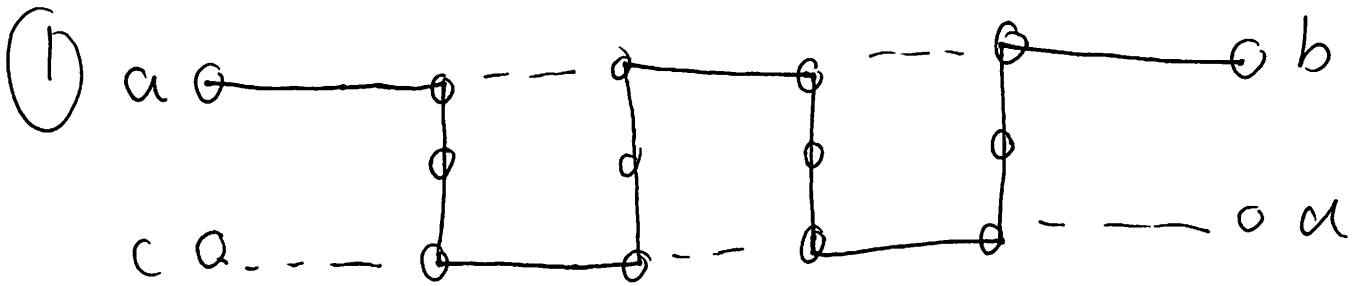
a, b, c, d connect the gadget to G 's gadget
no other connections.

Now study the gadget
in more detail.

We will see that there are
exactly 2 ways to traverse it.



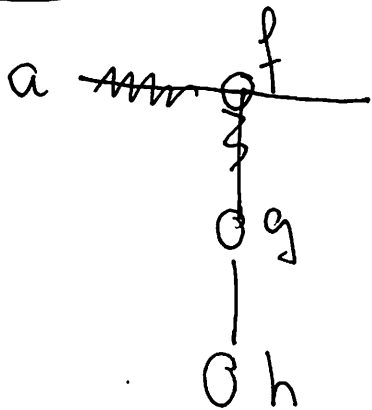
The 2 ways are



Why? see class

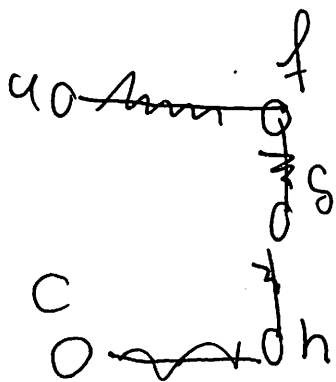
2 observations

(1)



if at f must go
to g at $\text{deg}(f) = 2$

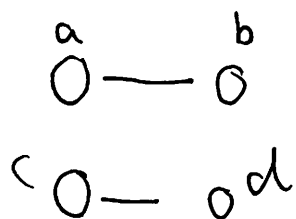
(2)



if a, f, g, h, c
is chosen the
rest of the gadget
cannot be traversed.
by a Hamiltonian
cycle.

pair of edges
Hamiltonian
cycle \rightarrow Hamiltonian
cycle

For each pair of pair of edges
($\{a, b\}, \{c, d\}$)



Construct the gadget

For any other pair that uses
($\{a, b\}, \{x, y\}$) replace b, y

($\{a, y\}, \{x, b\}$)

if $\{a, y\}$ is traversed in the gadget
then in the old graph
 $\{a, b\}$ would be traversed.

analogously $\{c, b\}$ replaces $\{c, d\}$

Each gadget adds 12 vertices
to the graph. + Polynomial time.

Is a Hamiltonian path in the
Pairwise problem instances

\Leftrightarrow

P' is a Hamiltonian path in
the so construct graph

(P' as described above)

Concludes reduction \square
 $20E \rightarrow$ Hamiltonian
Path

Recall,

SAT \rightarrow 3-SAT

Now

Any problem in NP \rightarrow SAT

We study Circuit sat defined as

given a Boolean circuit,
a dag with vertices : gates
of five different types

1) AND gates

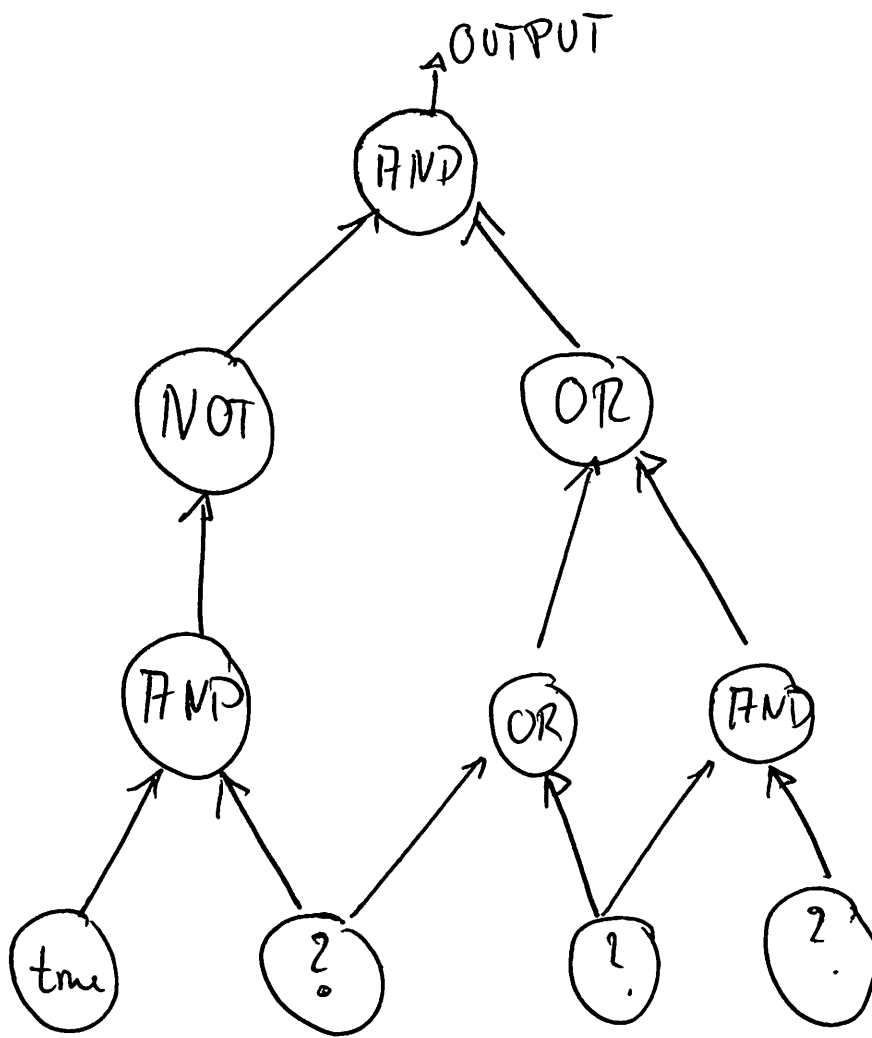
2) OR gates

3) NOT gates

4) Known input gate, (true / false)

5) Unknown input gates "2"

One of the DAG's sink is called "OUTPUT" gate.



After assigning true/false values to the "?"s the gates can be evaluated in topological order.

CIRCUIT SAT

Given a circuit, find a truth assignment for the unknown gates so that the circuit outputs "true".

- CIRCUIT SAT is a generalization of SAT
- a SAT formula is really a special circuit
 - $(\) (\)$
↑
and gate
 - $(\vee \vee)$ or gate
 - output is the value of the formula
 - each literal is an unknown input gate or its negation
 - no known input gates

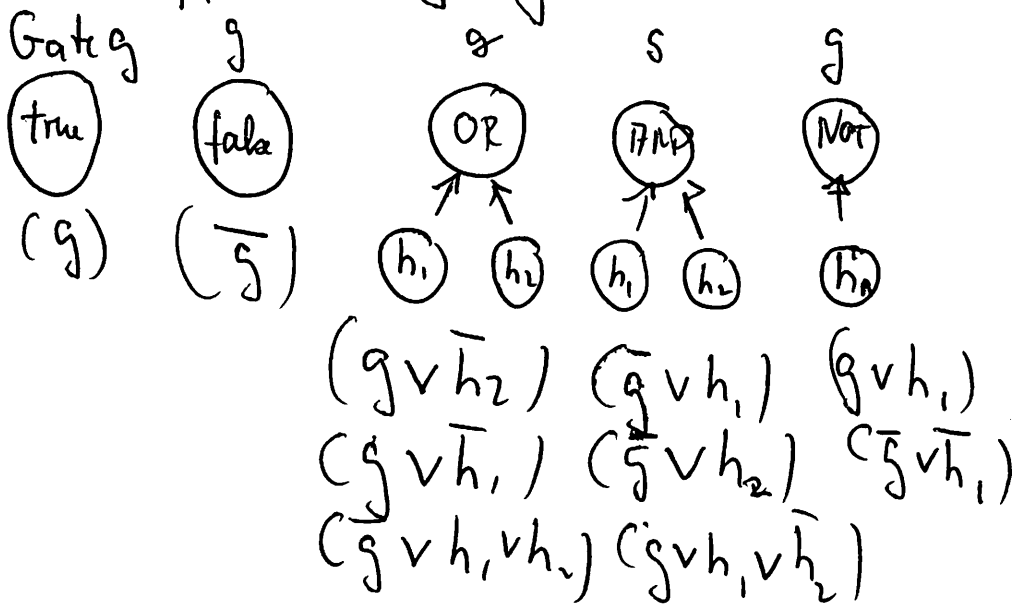
CIRCUIT SAT \rightarrow SAT

take a circuit \rightarrow write it in CNF

for each gate g in the circuit

\hookrightarrow create a variable g

- model effects of gates as:



[recall this construction in LP formulations]

- add ~~to~~ ^{to} output gate the clause (g)
value true is forced

So, CIRCUIT-SAT \rightarrow SAT

I now show that all search

problems in NP \rightarrow CIRCUIT-SAT

Any search problem's (in NP) solution can be checked in polynomial time.

(def. search problem in NP)

\Rightarrow there is a checking algorithm which takes an instance of the search problem, I , and a solution, S , and tests if S is a solution for I .

For the checking algorithm
 \bar{I} : is encoded as a binary
string of polynomial
length $o(\bar{I})$

Recall any polynomial algorithm
can be expressed (in a computer)
as a circuit, input gates
are the input to the algorithm
of gates polynomial in
the input.

Checking algorithm :

S is a solution to \bar{I} \rightarrow yes
 S is not a " " " \rightarrow no

The output gate then gives
true / false correspondingly.

Conclusion :

Instance I of problem A

\hookrightarrow in polynomial time circuit

known inputs \rightarrow bits encoding I

unknown input \rightarrow bits " S

Output: true S is a solution to I

false S is not a solution to I

NP - Complete Problems
Can be solved by "some"
algorithm.
Runtime likely exponential!

Some problems cannot be
solved at all!

There is NO ALGORITHM
that solves the problem.

Example:

are there integer values x, y, z
that satisfy

$$x^3 y z + 2 y^4 z^2 - 7 x y^5 z = 6?$$

No algorithm at all can solve this
polynomial, exponential,
whatever

Alan M. Turing 1936

Let P be a program
 X an input to P

Can we write a Program?

$\text{Terminates}(P, X)$

Input: any program P

X : input to P

Output: yes if P terminates
on input ~~X~~

no if P does not
terminate on X

Answer we cannot
write such a program!

Theorem No algorithm can exist which tests if any program terminates on a given input

Proof. Suppose there is such an algorithm $\text{terminates}(P, x)$

Then write this:

function $\text{paradox}(z: \text{file})$

1. if $\text{terminates}(z, z)$ goto 1

You see the "infinite loop"

if z applied on itself as input terminates

then paradox does not

if it does not then paradox will.

Now apply paradox to itself
(it is a program so that is legal)

Paradox stops when it does not ? 2.

Contradiction

So, no algorithm can exist
to decide if any algorithm stops.

↳ unsolvable
