WS 2021/22

Intro to Linear Programming

Harald Räcke

Fakultät für Informatik TU München

http://www14.in.tum.de/lehre/2021WS/ea/

Winter Term 2021/22

Organizational Matters

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► Modul: IN2003

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Name: "Efficient Algorithms and Data Structures"

"Effiziente Algorithmen und Datenstrukturen"

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ECTS: 8 Credit points

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Lectures:

4 SWS

Mon 10:00–12:00 (Room Interim2) Fri 10:00–12:00 (Room Interim2)

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Required knowledge:

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 - ► IN0001, IN0003
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 - "Diskrete Strukturen" (DS)



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 - ► IN0015
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 - "Diskrete Strukturen" (DS)
 - ► IN0018
 - "Discrete Probability Theory"
 - "Diskrete Wahrscheinlichkeitstheorie" (DWT)



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The Lecturer

- ► Harald Räcke
- Email: raecke@in.tum.de
- Room: 03.09.044
- Office hours: (by appointment)

Tutorials

```
Monday,
              12:00-14:00, 00.08.038
                                        (Michael Laraia)
  Monday,
              14:00-16:00, 02.09.023
                                        (Ruslan Zabrodin)
4 Tuesday,
              10:00-12:00, 00.08.053
                                        (Letian Shi)
  Tuesday,
              14:00-16:00, 00.08.038
                                        (Arnor Kristmundsson)
  Wednesday, 10:00-12:00, 03.11.018
                                        (Abdelrahman Metwally)
2 Wednesday, 12:00-14:00, online
                                        (Arnor Kristmundsson)
  Wednesday, 14:00-16:00, online
                                        (Abdelrahman Metwally)
 Thursday,
              16:00-18:00, online
                                        (Michael Laraia)
7 Friday.
              12:00-14:00. 00.13.009A (Ruslan Zabrodin)
```



Registration for Tutorials

Registration Period for Tutorial Sessions:

Saturday, 23 Oct- Tuesday, 26 Oct

via TUMonline; you have to choose at least 3 options...

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- This is done via Moodle.

Assignment sheets

In order to pass the module you need to pass an exam.

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- ➤ You should submit solutions in groups of up to 2 people.

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Submissions must be handwritten by a member of the group. Please indicate who wrote the submission.

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- Don't forget name and student id number for each group member.



Assignment can be used to improve you grade

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$$f(x) = \begin{cases} \frac{1}{10} \text{round} \left(10 \left(\frac{\text{round}(3x) - 1}{3} \right) \right) & 1 < x \le 4 \\ x & \text{otw.} \end{cases}$$

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If you obtain a bonus your grade will improve according to the following function

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 - > 4.0 no improvement

Assessment

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Requirements for Bonus

- 50% of the points are achieved on submissions 2-8,
- ▶ 50% of the points are achieved on submissions 9-14,
- each group member has written at least 4 solutions.

- Foundations
 - Machine models
 - Efficiency measures
 - Asymptotic notation
 - Recursion

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- Cuts/Flows
- Matchings



2 Literatur



Thomas H. Cormen, Charles E. Leiserson, Ron L. Rivest, Clifford Stein:

Introduction to algorithms,

McGraw-Hill, 1990

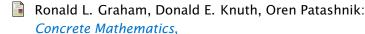
Michael T. Goodrich, Roberto Tamassia:

Algorithm design: Foundations, analysis, and internet examples,
John Wiley & Sons, 2002



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2 Literatur



2. Auflage, Addison-Wesley, 1994

Volker Heun:

Grundlegende Algorithmen: Einführung in den Entwurf und die Analyse effizienter Algorithmen,

2. Auflage, Vieweg, 2003

Jon Kleinberg, Eva Tardos:

Algorithm Design,

Addison-Wesley, 2005

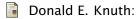
Donald F. Knuth:

The art of computer programming. Vol. 1: Fundamental Algorithms,

3. Auflage, Addison-Wesley, 1997



2 Literatur



The art of computer programming. Vol. 3: Sorting and Searching,

3. Auflage, Addison-Wesley, 1997

Christos H. Papadimitriou, Kenneth Steiglitz:

Combinatorial Optimization: Algorithms and Complexity,
Prentice Hall, 1982

Uwe Schöning: Algorithmik, Spektrum Akademischer Verlag, 2001

Steven S. Skiena:
The Algorithm Design Manual,
Springer, 1998

Part II

Foundations

3 Goals

Gain knowledge about efficient algorithms for important problems, i.e., learn how to solve certain types of problems efficiently.

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- Learn how to analyze and judge the efficiency of algorithms.
- Learn how to design efficient algorithms.

What do you measure?

Memory requirement

- Memory requirement
- Running time

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- Implementing and testing on representative inputs
 - How do you choose your inputs?
 - May be very time-consuming.
 - Very reliable results if done correctly.
 - Results only hold for a specific machine and for a specific set of inputs.

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- Implementing and testing on representative inputs
 - How do you choose your inputs?
 - May be very time-consuming.
 - Very reliable results if done correctly.
 - Results only hold for a specific machine and for a specific set of inputs.
- Theoretical analysis in a specific model of computation.
 - Gives asymptotic bounds like "this algorithm always runs in time $\mathcal{O}(n^2)$ ".
 - Typically focuses on the worst case.
 - Can give lower bounds like "any comparison-based sorting algorithm needs at least $\Omega(n \log n)$ comparisons in the worst case".



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Example 1

Suppose n numbers from the interval $\{1,\ldots,N\}$ have to be sorted. In this case we usually say that the input length is n instead of e.g. $n\log N$, which would be the number of bits required to encode the input.

How to measure performance

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 Calculate running time and storage space etc. on a simplified, idealized model of computation, e.g. Random Access Machine (RAM), Turing Machine (TM), . . .

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How to measure performance

- 1. Calculate running time and storage space etc. on a simplified, idealized model of computation, e.g. Random Access Machine (RAM), Turing Machine (TM), ...
- 2. Calculate number of certain basic operations: comparisons, multiplications, harddisc accesses, ...

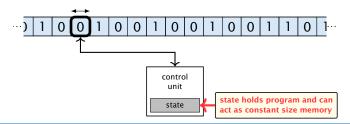
How to measure performance

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Version 2. is often easier, but focusing on one type of operation makes it more difficult to obtain meaningful results.

Turing Machine

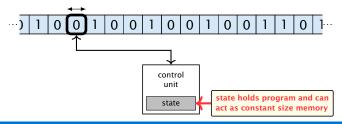
Very simple model of computation.



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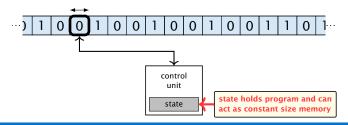
Turing Machine

- Very simple model of computation.
- Only the "current" memory location can be altered.



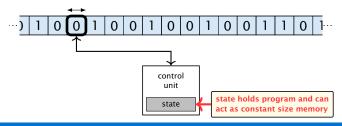
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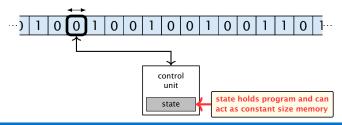
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- Some simple problems like recognizing whether input is of the form xx, where x is a string, have quadratic lower bound.

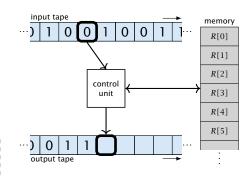


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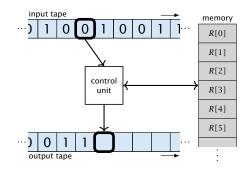
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- Very good model for discussing computability, or polynomial vs. exponential time.
- Some simple problems like recognizing whether input is of the form xx, where x is a string, have quadratic lower bound.
- \Rightarrow Not a good model for developing efficient algorithms.



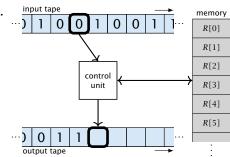
Input tape and output tape (sequences of zeros and ones; unbounded length).



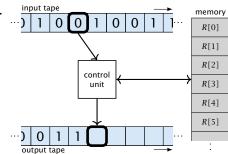
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- Indirect addressing.



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The jump-directives are very close to the jump-instructions contained in the assembler language of real machines.

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 - ► R[i] := R[j] + R[k];R[i] := -R[k];

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The latter model is quite realistic as the word-size of a standard computer that handles a problem of size n must be at least $\log_2 n$ as otherwise the computer could either not store the problem instance or not address all its memory.

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Bounded word RAM model: cost is uniform but the largest value stored in a register may not exceed 2^w , where usually $w = \log_2 n$.

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6. Feb. 2022

Example 2

Algorithm 1 RepeatedSquaring(n)

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- space requirement:
 - uniform model: $\mathcal{O}(1)$
 - logarithmic model: $\mathcal{O}(2^n)$



There are different types of complexity bounds:

best-case complexity:

$$C_{\rm bc}(n) := \min\{C(x) \mid |x| = n\}$$

Usually easy to analyze, but not very meaningful.

 $C(x) \begin{cases} cost \text{ of instance } \\ x \end{cases}$ $|x| \quad \begin{array}{c} \text{input length of instance } \\ x \\ \text{instance } x \\ \end{array}$ $I_n \quad \begin{array}{c} \text{set of instances} \\ \text{of length } n \\ \end{array}$

 μ is a probability distribution over inputs of length n.

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Usually moderately easy to analyze; sometimes too pessimistic.

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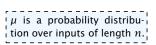
average case complexity:

$$C_{\text{avg}}(n) := \frac{1}{|I_n|} \sum_{|x|=n} C(x)$$

C(x) cost of instance input length of instance x

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more general: probability measure μ

$$\mu$$
 is a probability distribution over inputs of length n .

$$C_{\operatorname{avg}}(n) := \sum_{x \in I_n} \mu(x) \cdot C(x)$$

$$C(x) \begin{cases} \cos t \text{ of instance} \\ x \end{cases}$$

$$|x| \text{ input length of instance } x$$

 I_n set of instances of length n

amortized complexity:

The average cost of data structure operations over a worst case sequence of operations.

 $\begin{array}{c|c} C(x) & C(x) \\ \hline x & \\ \hline |x| & \text{in} \\ \text{in} \\ \text{in over inputs of length } n. \\ \end{array}$

x |x| input length of instance x I_n set of instances of length n

cost of instance



- amortized complexity:
 - The average cost of data structure operations over a worst case sequence of operations.
- randomized complexity:

The algorithm may use random bits. Expected running time (over all possible choices of random bits) for a fixed input x.

Then take the worst-case over all x with |x| = n.

	C(x)	cost of instance
1		X
1	x	input length of
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- Running time should be expressed by simple functions.



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Formal Definition

Let f, g denote functions from \mathbb{N} to \mathbb{R}^+ .

• $\mathcal{O}(f) = \{g \mid \exists c > 0 \ \exists n_0 \in \mathbb{N}_0 \ \forall n \geq n_0 : [g(n) \leq c \cdot f(n)] \}$ (set of functions that asymptotically grow not faster than f)

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32/616

There is an equivalent definition using limes notation (assuming that the respective limes exists). f and g are functions from \mathbb{N}_0 to \mathbb{R}_0^+ .

- Note that for the version of the Landau notation defined here, we assume that f and g are positive functions.
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Abuse of notation

1. People write $f = \mathcal{O}(g)$, when they mean $f \in \mathcal{O}(g)$. This is **not** an equality (how could a function be equal to a set of functions).

- **2.** In this context f(n) does **not** mean the function f evaluated at n, but instead it is a shorthand for the function itself (leaving out
 - 3. This is particularly useful if you do not want to ignore constant factors. For example the median of n elements can be determined usdomain and codomain and only giving the ing $\frac{3}{2}n + o(n)$ comparisons. rule of correspondence of the function).

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- 3. People write e.g. h(n) = f(n) + o(g(n)) when they mean that there exists a function $z: \mathbb{N} \to \mathbb{R}^+, n \mapsto z(n), z \in o(g)$ such that h(n) = f(n) + z(n).

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- **4.** People write $\mathcal{O}(f(n)) = \mathcal{O}(g(n))$, when they mean $\mathcal{O}(f(n)) \subseteq \mathcal{O}(g(n))$. Again this is not an equality.

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Note that $\Theta(n)$ is on the right hand side, otw. this interpretation is wrong.

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Regardless of how we choose the anonymous function $f(n) \in \mathcal{O}(n)$ there is an anonymous function $g(n) \in \Theta(n^2)$ that makes the expression true.

How do we interpret an expression like:

$$\sum_{i=1}^{n} \Theta(i) = \Theta(n^2)$$

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Careful!

The $\Theta(i)$ -symbol on the left represents one anonymous function $f: \mathbb{N} \to \mathbb{R}^+$, and then $\sum_i f(i)$ is computed.

How do we interpret an expression like:

$$\sum_{i=1}^{n} \Theta(i) = \Theta(n^2)$$

Careful!

"It is understood" that every occurrence of an \mathcal{O} -symbol (or $\Theta, \Omega, \rho, \omega$) on the left represents one anonymous function.

Hence, the left side is **not** equal to

$$\Theta(1) + \Theta(2) + \cdots + \Theta(n-1) + \Theta(n)$$

 $\Theta(1) + \Theta(2) + \cdots + \Theta(n-1) + \Theta(n)$ does not really have a reasonable interpreta-

We can view an expression containing asymptotic notation as generating a set:

$$n^2 \cdot \mathcal{O}(n) + \mathcal{O}(\log n)$$

represents

$$\left\{ f: \mathbb{N} \to \mathbb{R}^+ \mid f(n) = n^2 \cdot g(n) + h(n) \right.$$
 with $g(n) \in \mathcal{O}(n)$ and $h(n) \in \mathcal{O}(\log n) \right\}$ Recall that according to the previous slide e.g. the expressions $\sum_{i=1}^n \mathcal{O}(i)$ and $\sum_{i=1}^{n/2} \mathcal{O}(i) + \sum_{i=n/2+1}^n \mathcal{O}(i)$ generate differential $\sum_{i=1}^{n/2} \mathcal{O}(i) + \sum_{i=n/2+1}^n \mathcal{O}(i)$ generate differential $\sum_{i=1}^n \mathcal{O}(i) + \sum_{i=n/2+1}^n \mathcal{O}(i)$

Then an asymptotic equation can be interpreted as containement btw. two sets:

$$n^2 \cdot \mathcal{O}(n) + \mathcal{O}(\log n) = \Theta(n^2)$$

represents

$$n^2 \cdot \mathcal{O}(n) + \mathcal{O}(\log n) \subseteq \Theta(n^2)$$

Note that the equation does not hold.

Lemma 3

Let f, g be functions with the property

 $\exists n_0 > 0 \ \forall n \ge n_0 : f(n) > 0$ (the same for g). Then

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- $\mathcal{O}(f(n)) + \mathcal{O}(g(n)) = \mathcal{O}(\max\{f(n), g(n)\})$

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- $\triangleright \mathcal{O}(f(n)) + \mathcal{O}(g(n)) = \mathcal{O}(\max\{f(n),g(n)\})$

The expressions also hold for Ω . Note that this means that $f(n) + g(n) \in \Theta(\max\{f(n), g(n)\}).$

Comments

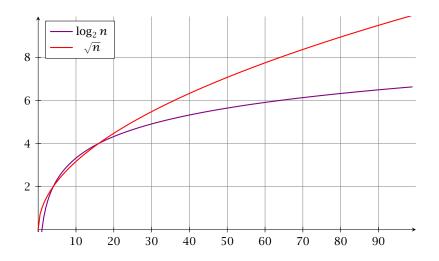
Do not use asymptotic notation within induction proofs.

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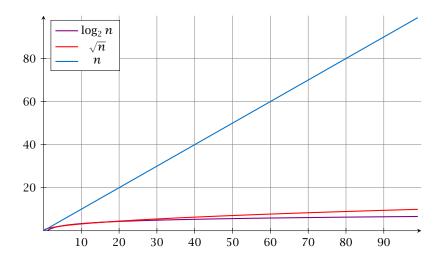
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- For any constants a, b we have $\log_a n = \Theta(\log_b n)$. Therefore, we will usually ignore the base of a logarithm within asymptotic notation.

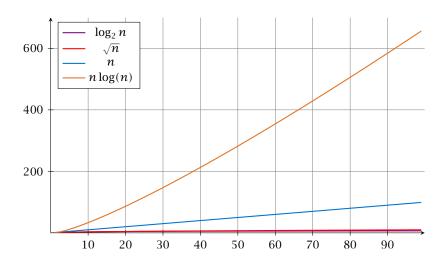
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- Do not use asymptotic notation within induction proofs.
- For any constants a, b we have $\log_a n = \Theta(\log_b n)$. Therefore, we will usually ignore the base of a logarithm within asymptotic notation.
- In general $\log n = \log_2 n$, i.e., we use 2 as the default base for the logarithm.

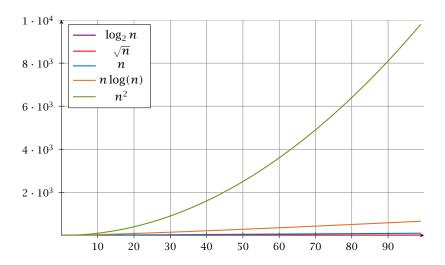




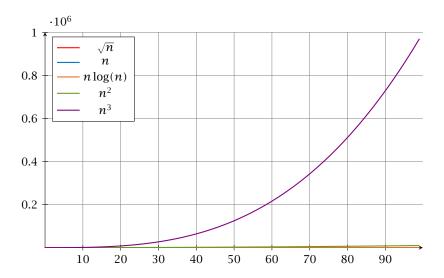




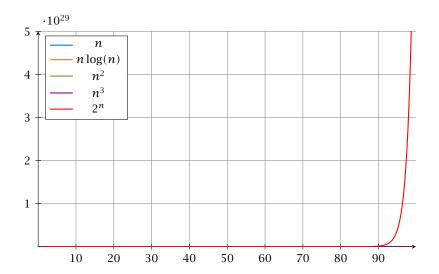












Laufzeiten

Funktion	Eingabelänge n							
f(n)	10	10^{2}	10^{3}	10^{4}	10^{5}	10^{6}	10 ⁷	10^{8}
$\log n$	33ns	66ns	0.1µs	0.1µs	0.2µs	0.2µs	0.2µs	0.3µs
\sqrt{n}	32ns	0.1µs	0.3µs	1µs	3.1 µs	10µs	31 µs	$0.1 \mathrm{ms}$
n	100ns	1µs	10µs	$0.1 \mathrm{ms}$	1ms	10ms	0.1s	1s
$n \log n$	0.3µs	6.6µs	0.1ms	1.3ms	16ms	0.2s	2.3s	27s
$n^{3/2}$	0.3µs	10µs	0.3ms	10ms	0.3s	10s	5.2min	2.7h
n^2	1µs	0.1 ms	10ms	1s	1.7min	2.8h	11 d	3.2 y
n^3	10µs	10ms	10s	2.8h	115 d	317 y	3.2·10 ⁵ y	
1.1^{n}	26ns	$0.1 \mathrm{ms}$	$7.8 \cdot 10^{25}$ y					
2^n	10µs	$4\cdot 10^{14}$ y						
n!	36ms	$3 \cdot 10^{142}$ y						

1 Operation = 10ns; 100MHz

Alter des Universums: ca. $13.8 \cdot 10^9 \mathrm{y}$

In general asymptotic classification of running times is a good measure for comparing algorithms:

▶ If the running time analysis is tight and actually occurs in practise (i.e., the asymptotic bound is not a purely theoretical worst-case bound), then the algorithm that has better asymptotic running time will always outperform a weaker algorithm for large enough values of n.

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- However, suppose that I have two algorithms:

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Clearly f = o(g). However, as long as $\log n \le 1000$ Algorithm B will be more efficient.



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Formal Definition

Let f, g denote functions from \mathbb{N}^d to \mathbb{R}_0^+ .

 $\mathcal{O}(f) = \{g \mid \exists c > 0 \ \exists N \in \mathbb{N}_0 \ \forall \vec{n} \ \text{with} \ n_i \geq N \ \text{for some} \ i : \\ [g(\vec{n}) \leq c \cdot f(\vec{n})] \}$

(set of functions that asymptotically grow not faster than f)

Example 4

 $f: \mathbb{N} \to \mathbb{R}_0^+, f(n,m) = 1 \text{ und } g: \mathbb{N} \to \mathbb{R}_0^+, g(n,m) = n-1$

Example 4

▶ $f: \mathbb{N} \to \mathbb{R}_0^+$, f(n,m) = 1 und $g: \mathbb{N} \to \mathbb{R}_0^+$, g(n,m) = n-1 then $f = \mathcal{O}(g)$ does not hold

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Algorithm 2 mergesort(list *L*)

1: $n \leftarrow \text{size}(L)$

2: if $n \le 1$ return L

3: $L_1 \leftarrow L[1 \cdots \lfloor \frac{n}{2} \rfloor]$

4: $L_2 \leftarrow L[\lfloor \frac{n}{2} \rfloor + 1 \cdots n]$

5: $mergesort(L_1)$

6: mergesort(L_2)

7: $L \leftarrow \text{merge}(L_1, L_2)$

8: return L

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6: mergesort(L_2)
7: $L \leftarrow \text{merge}(L_1, L_2)$ 8: **return** L

This algorithm requires

$$T(n) = T(\left\lceil \frac{n}{2} \right\rceil) + T(\left\lfloor \frac{n}{2} \right\rfloor) + \mathcal{O}(n) \le 2T(\left\lceil \frac{n}{2} \right\rceil) + \mathcal{O}(n)$$

comparisons when n > 1 and 0 comparisons when $n \le 1$.

How do we bring the expression for the number of comparisons (≈ running time) into a closed form?

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For this we need to solve the recurrence.

Methods for Solving Recurrences

1. Guessing+Induction

Guess the right solution and prove that it is correct via induction. It needs experience to make the right guess.

2. Master Theorem

For a lot of recurrences that appear in the analysis of algorithms this theorem can be used to obtain tight asymptotic bounds. It does not provide exact solutions.

3. Characteristic Polynomial

Linear homogenous recurrences can be solved via this method.



Methods for Solving Recurrences

4. Generating Functions

A more general technique that allows to solve certain types of linear inhomogenous relations and also sometimes non-linear recurrence relations.

5. Transformation of the Recurrence

Sometimes one can transform the given recurrence relations so that it e.g. becomes linear and can therefore be solved with one of the other techniques.



6.1 Guessing+Induction

First we need to get rid of the \mathcal{O} -notation in our recurrence:

$$T(n) \le \begin{cases} 2T(\lceil \frac{n}{2} \rceil) + cn & n \ge 2\\ 0 & \text{otherwise} \end{cases}$$

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One way of solving such a recurrence is to guess a solution, and check that it is correct by plugging it in.

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Suppose we guess $T(n) \le dn \log n$ for a constant d. Then

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if we choose d > c.

Formally, this is not correct if n is not a power of 2. Also even in this case one would need to do an induction proof.

$$T(n) \le \begin{cases} 2T(\frac{n}{2}) + cn & n \ge 16 \\ b & \text{otw.} \end{cases}$$

- Note that this proves the statement for $n=2^k$, $k\in\mathbb{N}_{\geq 1}$, as the statement is wrong for n=1.
- The base case is usually omitted, as it is the same for different recurrences.

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base case $(2 \le n < 16)$:

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Note that we can do this as for constant-sized inputs the running time is always some constant (b in the above case).

We also make a guess of $T(n) \le dn \log n$ and get

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6.1 Guessina+Induction

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We also make a guess of $T(n) \le dn \log n$ and get

$$T(n) \leq 2T\left(\left\lceil\frac{n}{2}\right\rceil\right) + cn$$

$$\leq 2\left(d\left\lceil\frac{n}{2}\right\rceil\log\left\lceil\frac{n}{2}\right\rceil\right) + cn$$

$$\left\lceil\frac{n}{2}\right\rceil \leq \frac{n}{2} + 1\right\rceil \leq 2\left(d(n/2 + 1)\log(n/2 + 1)\right) + cn$$

$$\left\lceil\frac{n}{2} + 1 \leq \frac{9}{16}n\right\rceil \leq dn\log\left(\frac{9}{16}n\right) + 2d\log n + cn$$

$$\left\lceil\log\frac{9}{16}n\right\rceil = \log n + (\log 9 - 4)\right\rceil = dn\log n + (\log 9 - 4)dn + 2d\log n + cn$$

$$\left\lceil\log n \leq \frac{n}{4}\right\rceil \leq dn\log n + (\log 9 - 3.5)dn + cn$$

$$\leq dn\log n - 0.33dn + cn$$

$$\leq dn\log n$$

for a suitable choice of d.

Lemma 5

Let $a \ge 1, b \ge 1$ and $\epsilon > 0$ denote constants. Consider the recurrence

$$T(n) = aT\left(\frac{n}{h}\right) + f(n) .$$

Case 1.

If
$$f(n) = \mathcal{O}(n^{\log_b(a) - \epsilon})$$
 then $T(n) = \Theta(n^{\log_b a})$.

Case 2.

If
$$f(n) = \Theta(n^{\log_b(a)} \log^k n)$$
 then $T(n) = \Theta(n^{\log_b a} \log^{k+1} n)$, $k \ge 0$.

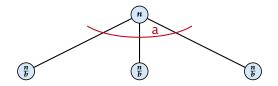
Case 3.

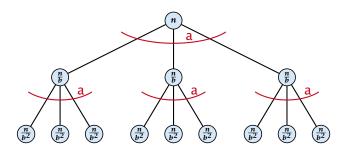
If
$$f(n) = \Omega(n^{\log_b(a) + \epsilon})$$
 and for sufficiently large n $af(\frac{n}{b}) \le cf(n)$ for some constant $c < 1$ then $T(n) = \Theta(f(n))$.

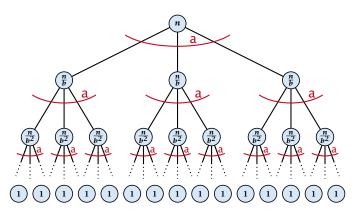
6.2 Master Theorem

We prove the Master Theorem for the case that n is of the form b^{ℓ} , and we assume that the non-recursive case occurs for problem size 1 and incurs cost 1.

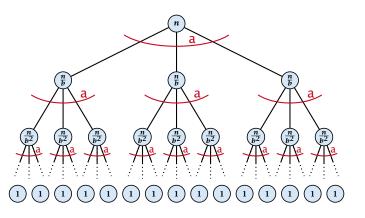






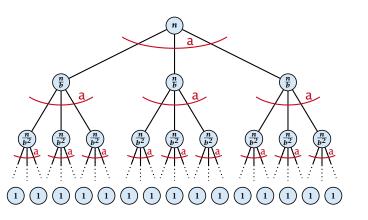


The running time of a recursive algorithm can be visualized by a recursion tree:



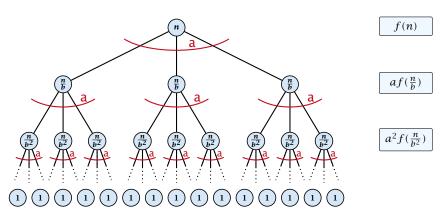
f(n)

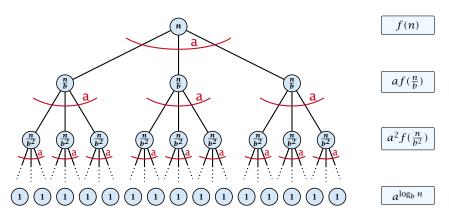
The running time of a recursive algorithm can be visualized by a recursion tree:

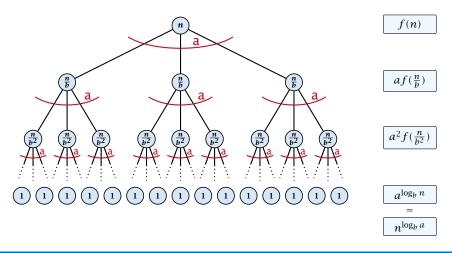


f(n)

 $af(\frac{n}{b})$







6.2 Master Theorem

This gives

$$T(n) = n^{\log_b a} + \sum_{i=0}^{\log_b n-1} a^i f\left(\frac{n}{b^i}\right) \ .$$

$$T(n) - n^{\log_b a}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$b^{-i(\log_b a - \epsilon)} = b^{\epsilon i} (b^{\log_b a})^{-i} = b^{\epsilon i} a^{-i}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$b^{-i(\log_b a - \epsilon)} = b^{\epsilon i} (b^{\log_b a})^{-i} = b^{\epsilon i} a^{-i} = c n^{\log_b a - \epsilon} \sum_{i=0}^{i=0} (b^{\epsilon})^i$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\underline{b^{-i(\log_b a - \epsilon)} = b^{\epsilon i}(b^{\log_b a})^{-i} = b^{\epsilon i}a^{-i}} = c n^{\log_b a - \epsilon} \sum_{i=0}^{\log_b n - 1} \left(b^{\epsilon}\right)^i$$

$$\sum_{i=0}^k q^i = \frac{q^{k+1} - 1}{a-1}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\underline{b^{-i(\log_b a - \epsilon)} = b^{\epsilon i}(b^{\log_b a})^{-i} = b^{\epsilon i}a^{-i}} = c n^{\log_b a - \epsilon} \sum_{i=0}^{\log_b n - 1} (b^{\epsilon})^i$$

$$\sum_{i=0}^k q^i = \frac{q^{k+1} - 1}{a^{-1}} = c n^{\log_b a - \epsilon} (b^{\epsilon \log_b n} - 1) / (b^{\epsilon} - 1)$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\underline{b^{-i(\log_b a - \epsilon)} = b^{\epsilon i}(b^{\log_b a})^{-i} = b^{\epsilon i}a^{-i}} = c n^{\log_b a - \epsilon} \sum_{i=0}^{\log_b n - 1} (b^{\epsilon})^i$$

$$\sum_{i=0}^k q^i = \frac{q^{k+1} - 1}{q - 1} = c n^{\log_b a - \epsilon} (b^{\epsilon \log_b n} - 1) / (b^{\epsilon} - 1)$$

$$= c n^{\log_b a - \epsilon} (n^{\epsilon} - 1) / (b^{\epsilon} - 1)$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\underline{b^{-i(\log_b a - \epsilon)} = b^{\epsilon i} (b^{\log_b a})^{-i} = b^{\epsilon i} a^{-i}} = c n^{\log_b a - \epsilon} \sum_{i=0}^{\log_b n - 1} (b^{\epsilon})^i$$

$$\underline{\sum_{i=0}^k q^i = \frac{q^{k+1} - 1}{q - 1}} = c n^{\log_b a - \epsilon} (b^{\epsilon \log_b n} - 1) / (b^{\epsilon} - 1)$$

$$= c n^{\log_b a - \epsilon} (n^{\epsilon} - 1) / (b^{\epsilon} - 1)$$

$$= \frac{c}{b^{\epsilon} - 1} n^{\log_b a} (n^{\epsilon} - 1) / (n^{\epsilon})$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\begin{bmatrix}
b^{-i(\log_b a - \epsilon)} = b^{\epsilon i}(b^{\log_b a})^{-i} = b^{\epsilon i}a^{-i}
\end{bmatrix} = cn^{\log_b a - \epsilon} \sum_{i=0} (b^{\epsilon})^i$$

$$\begin{bmatrix}
\sum_{i=0}^k q^i = \frac{q^{k+1} - 1}{q - 1}
\end{bmatrix} = cn^{\log_b a - \epsilon} (b^{\epsilon \log_b n} - 1)/(b^{\epsilon} - 1)$$

$$= cn^{\log_b a - \epsilon} (n^{\epsilon} - 1)/(b^{\epsilon} - 1)$$

$$= \frac{c}{b^{\epsilon} - 1} n^{\log_b a} (n^{\epsilon} - 1)/(n^{\epsilon})$$

Hence,

$$T(n) \le \left(\frac{c}{h^{\epsilon} - 1} + 1\right) n^{\log_b(a)}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a - \epsilon}$$

$$\frac{b^{-i(\log_b a - \epsilon)} = b^{\epsilon i}(b^{\log_b a})^{-i} = b^{\epsilon i}a^{-i}}{b^{-i(\log_b a - \epsilon)}} = cn^{\log_b a - \epsilon} \sum_{i=0}^{\log_b n - 1} (b^{\epsilon})^i$$

$$\sum_{i=0}^k q^i = \frac{q^{k+1} - 1}{q - 1} = cn^{\log_b a - \epsilon} (b^{\epsilon \log_b n} - 1)/(b^{\epsilon} - 1)$$

$$= cn^{\log_b a - \epsilon} (n^{\epsilon} - 1)/(b^{\epsilon} - 1)$$

$$= c n^{\log_b a} { (n^{\epsilon} - 1)/(b^{\epsilon} - 1)}$$
$$= \frac{c}{b^{\epsilon} - 1} n^{\log_b a} (n^{\epsilon} - 1)/(n^{\epsilon})$$

Hence,

$$T(n) \le \left(\frac{c}{h^{\epsilon} - 1} + 1\right) n^{\log_b(a)}$$

$$\Rightarrow T(n) = \mathcal{O}(n^{\log_b a}).$$

$$T(n) - n^{\log_b a}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

Hence,

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

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

Hence,

$$T(n) = \mathcal{O}(n^{\log_b a} \log_b n)$$
 $\Rightarrow T(n) = \mathcal{O}(n^{\log_b a} \log n).$

$$\Rightarrow T(n) = \mathcal{O}(n^{\log_b a} \log n).$$

$$T(n) - n^{\log_b a}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\begin{split} T(n) - n^{\log_b a} &= \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right) \\ &\geq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \end{split}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\geq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\geq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\geq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

Hence,

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

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\geq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a}$$

$$= c n^{\log_b a} \sum_{i=0}^{\log_b n - 1} 1$$

$$= c n^{\log_b a} \log_b n$$

Hence,

$$T(n) = \mathbf{\Omega}(n^{\log_b a} \log_b n)$$
 $\Rightarrow T(n) = \mathbf{\Omega}(n^{\log_b a} \log n).$

$$\Rightarrow T(n) = \mathbf{\Omega}(n^{\log_b a} \log n).$$

$$T(n) - n^{\log_b a}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$n=b^\ell\Rightarrow \ell=\log_b n$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$\boxed{n = b^\ell \Rightarrow \ell = \log_b n} = c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^\ell}{b^i}\right)\right)^k$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$\boxed{n = b^\ell \Rightarrow \ell = \log_b n} = c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^\ell}{b^i}\right)\right)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} (\ell - i)^k$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$\boxed{n = b^\ell \Rightarrow \ell = \log_b n} = c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^\ell}{b^i}\right)\right)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} (\ell - i)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell} i^k$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$n = b^{\ell} \Rightarrow \ell = \log_b n$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^{\ell}}{b^i}\right)\right)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} (\ell - i)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell} i^k \approx \frac{1}{k} \ell^{k+1}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$\boxed{n = b^{\ell} \Rightarrow \ell = \log_b n} = c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^{\ell}}{b^i}\right)\right)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} (\ell - i)^k$$

$$= c n^{\log_b a} \sum_{i=1}^{\ell} i^k$$

$$\approx \frac{c}{b} n^{\log_b a} \ell^{k+1}$$

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq c \sum_{i=0}^{\log_b n - 1} a^i \left(\frac{n}{b^i}\right)^{\log_b a} \cdot \left(\log_b \left(\frac{n}{b^i}\right)\right)^k$$

$$n = b^{\ell} \Rightarrow \ell = \log_b n$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} \left(\log_b \left(\frac{b^{\ell}}{b^i}\right)\right)^k$$

$$= c n^{\log_b a} \sum_{i=0}^{\ell - 1} (\ell - i)^k$$

$$= c n^{\log_b a} \sum_{i=1}^{\ell} i^k$$

$$\approx \frac{c}{\nu} n^{\log_b a} \ell^{k+1}$$

$$\Rightarrow T(n) = \mathcal{O}(n^{\log_b a} \log^{k+1} n).$$

From this we get $a^i f(n/b^i) \le c^i f(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

From this we get $a^if(n/b^i) \le c^if(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

From this we get $a^if(n/b^i) \le c^if(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq \sum_{i=0}^{\log_b n - 1} c^i f(n) + \mathcal{O}(n^{\log_b a})$$

From this we get $a^i f(n/b^i) \le c^i f(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq \sum_{i=0}^{\log_b n - 1} c^i f(n) + \mathcal{O}(n^{\log_b a})$$

$$q < 1: \sum_{i=0}^{n} q^i = \frac{1-q^{n+1}}{1-q} \le \frac{1}{1-q}$$

Where did we use $f(n) \ge \Omega(n^{\log_b a + \epsilon})$?

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From this we get $a^if(n/b^i) \le c^if(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq \sum_{i=0}^{\log_b n - 1} c^i f(n) + \mathcal{O}(n^{\log_b a})$$

$$q < 1: \sum_{i=0}^n q^i = \frac{1 - q^{n+1}}{1 - q} \leq \frac{1}{1 - c} f(n) + \mathcal{O}(n^{\log_b a})$$

Where did we use $f(n) \ge \Omega(n^{\log_b a + \epsilon})$?

From this we get $a^i f(n/b^i) \le c^i f(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq \sum_{i=0}^{\log_b n - 1} c^i f(n) + \mathcal{O}(n^{\log_b a})$$

$$q < 1: \sum_{i=0}^n q^i = \frac{1 - q^{n+1}}{1 - q} \leq \frac{1}{1 - c} f(n) + \mathcal{O}(n^{\log_b a})$$

Hence,

$$T(n) \le \mathcal{O}(f(n))$$

From this we get $a^i f(n/b^i) \le c^i f(n)$, where we assume that $n/b^{i-1} \ge n_0$ is still sufficiently large.

$$T(n) - n^{\log_b a} = \sum_{i=0}^{\log_b n - 1} a^i f\left(\frac{n}{b^i}\right)$$

$$\leq \sum_{i=0}^{\log_b n - 1} c^i f(n) + \mathcal{O}(n^{\log_b a})$$

$$q < 1: \sum_{i=0}^n q^i = \frac{1 - q^{n+1}}{1 - q} \leq \frac{1}{1 - c} f(n) + \mathcal{O}(n^{\log_b a})$$

Hence,

$$T(n) \le \mathcal{O}(f(n))$$

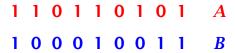
$$\Rightarrow T(n) = \Theta(f(n)).$$

Where did we use $f(n) \ge \Omega(n^{\log_b a + \epsilon})$?

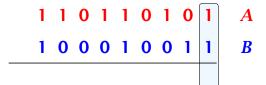
Suppose we want to multiply two n-bit Integers, but our registers can only perform operations on integers of constant size.

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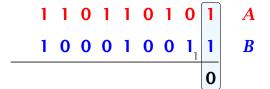
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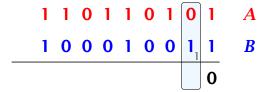
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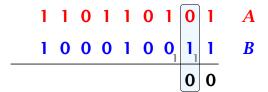
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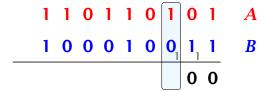
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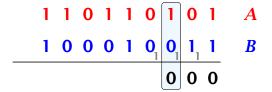
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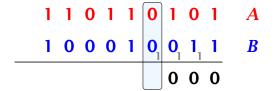
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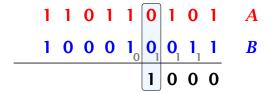
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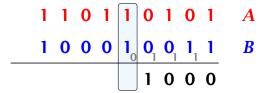
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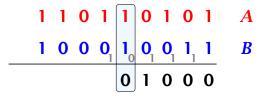
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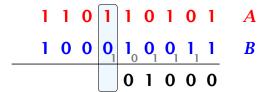


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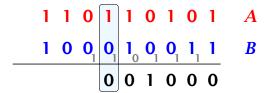
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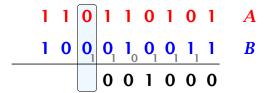


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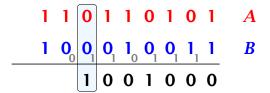


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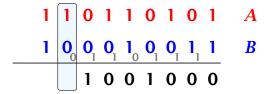
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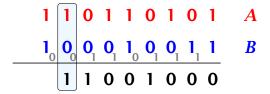


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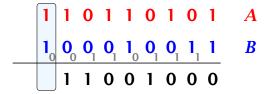
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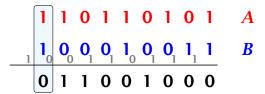
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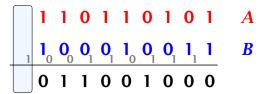


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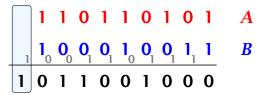


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For this we first need to be able to add two integers \mathbf{A} and \mathbf{B} :

This gives that two n-bit integers can be added in time $\mathcal{O}(n)$.

- This is also nown as the "school method" for multiplying integers.
- Note that the intermediate numbers that are generated can have at most $m + n \le 2n$ bits.

Suppose that we want to multiply an n-bit integer A and an m-bit integer B ($m \le n$).

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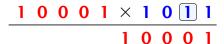
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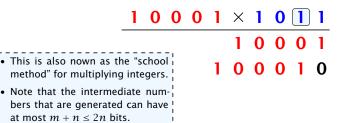
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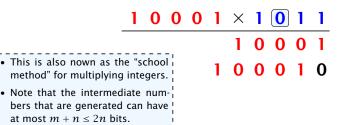


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_	1	0	0	0	1	×	1	0	1	1
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Note that the intermed bers that are generate				0	0	0	0	0	0	0
at most $m + n \le 2n$ b								0	0	0

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	1	0	0	0	1	\times	1	0	1	1
						1	0	0	0	1
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-			1	n	1	1	1	n	1	1

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	1	0	0	0	1	X	1	0	1	1	
						1	0	0	0	1	
 This is also nown as th method" for multiplying 					1	0	0	0	1	0	
 Note that the intermed bers that are generated 				0	0	0	0	0	0	0	
at most $m + n \le 2n$ bits			1	0	0	0	1	0	0	0	
-			1	0	1	1	1	0	1	1	

Time requirement:

Suppose that we want to multiply an n-bit integer A and an m-bit integer B ($m \le n$).

	1	0	0	0	1	×	1	0	1	1	
-						1	0	0	0	1	
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 Note that the intermed bers that are generated 				0	0	0	0	0	0	0	
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▶ Computing intermediate results: O(nm).

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-			1	0	1	1	1	0	1	1

Time requirement:

- ▶ Computing intermediate results: O(nm).
- ▶ Adding m numbers of length $\leq 2n$: $\mathcal{O}((m+n)m) = \mathcal{O}(nm)$.

A recursive approach:

Suppose that integers **A** and **B** are of length $n = 2^k$, for some k.

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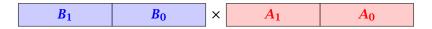
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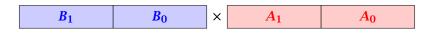
$$b_{n-1} \cdots b_{\frac{n}{2}} b_{\frac{n}{2}-1} \cdots b_0 \times a_{n-1} \cdots a_{\frac{n}{2}} a_{\frac{n}{2}-1} \cdots a_0$$

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Suppose that integers **A** and **B** are of length $n = 2^k$, for some k.



Then it holds that

$$A = A_1 \cdot 2^{\frac{n}{2}} + A_0$$
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Hence,

$$A \cdot B = A_1 B_1 \cdot 2^n + (A_1 B_0 + A_0 B_1) \cdot 2^{\frac{n}{2}} + A_0 B_0$$

Algorithm 3 mult(A, B)

1: **if** |A| = |B| = 1 **then** 2: **return** $a_0 \cdot b_0$

3: split A into A_0 and A_1 4: split B into B_0 and B_1 5: $Z_2 \leftarrow \text{mult}(A_1, B_1)$ 6: $Z_1 \leftarrow \text{mult}(A_1, B_0) + \text{mult}(A_0, B_1)$ 7: $Z_0 \leftarrow \text{mult}(A_0, B_0)$ 8: **return** $Z_2 \cdot 2^n + Z_1 \cdot 2^{\frac{n}{2}} + Z_0$

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6: $Z_1 \leftarrow \operatorname{mult}(A_1, B_0) + \operatorname{mult}(A_0, B_1)$	$2T(\frac{n}{2}) + \mathcal{O}(n)$
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We get the following recurrence:

$$T(n) = 4T\left(\frac{n}{2}\right) + \mathcal{O}(n) .$$

Master Theorem: Recurrence: $T[n] = aT(\frac{n}{b}) + f(n)$.

- ► Case 1: $f(n) = O(n^{\log_b a \epsilon})$ $T(n) = O(n^{\log_b a})$
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⇒ Not better then the "school method".

We can use the following identity to compute Z_1 :

A more precise (correct) analysis would say that computing Z_1 needs time $T(\frac{n}{2}+1)+\mathcal{O}(n)$.

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Hence,

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A huge improvement over the "school method".

6.3 The Characteristic Polynomial

Consider the recurrence relation:

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Note that we ignore boundary conditions for the moment.



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Approach:

- First determine all solutions that satisfy recurrence relation.
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- First consider the homogenous case.

The solution space

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How do we find a non-trivial solution?

We guess that the solution is of the form λ^n , $\lambda \neq 0$, and see what happens. In order for this guess to fulfill the recurrence we need

$$c_0\lambda^n + c_1\lambda^{n-1} + c_2 \cdot \lambda^{n-2} + \dots + c_k \cdot \lambda^{n-k} = 0$$

for all n > k.



Dividing by λ^{n-k} gives that all these constraints are identical to

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Let $\lambda_1, \ldots, \lambda_k$ be the k (complex) roots of $P[\lambda]$. Then, because of the vector space property

$$\alpha_1\lambda_1^n + \alpha_2\lambda_2^n + \cdots + \alpha_k\lambda_k^n$$

is a solution for arbitrary values α_i .

Lemma 6

Assume that the characteristic polynomial has k distinct roots $\lambda_1, \ldots, \lambda_k$. Then all solutions to the recurrence relation are of the form

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There is one solution for every possible choice of boundary conditions for $T[1], \ldots, T[k]$.

We show that the above set of solutions contains one solution for every choice of boundary conditions.

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We show that the column vectors are linearly independent. Then the above equation has a solution.

$$\begin{vmatrix} \lambda_1 & \lambda_2 & \cdots & \lambda_{k-1} & \lambda_k \\ \lambda_1^2 & \lambda_2^2 & \cdots & \lambda_{k-1}^2 & \lambda_k^2 \\ \vdots & \vdots & & \vdots & \vdots \\ \lambda_1^k & \lambda_2^k & \cdots & \lambda_{k-1}^k & \lambda_k^k \end{vmatrix} =$$

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$$\begin{vmatrix} 1 & \lambda_1 - \lambda_1 \cdot 1 & \cdots & \lambda_1^{k-2} - \lambda_1 \cdot \lambda_1^{k-3} & \lambda_1^{k-1} - \lambda_1 \cdot \lambda_1^{k-2} \\ 1 & \lambda_2 - \lambda_1 \cdot 1 & \cdots & \lambda_2^{k-2} - \lambda_1 \cdot \lambda_2^{k-3} & \lambda_2^{k-1} - \lambda_1 \cdot \lambda_2^{k-2} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & \lambda_k - \lambda_1 \cdot 1 & \cdots & \lambda_k^{k-2} - \lambda_1 \cdot \lambda_k^{k-3} & \lambda_k^{k-1} - \lambda_1 \cdot \lambda_k^{k-2} \end{vmatrix} =$$

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$$\begin{vmatrix} \sum_{i=2}^{k} (\lambda_i - \lambda_1) \cdot \begin{pmatrix} 1 & \lambda_2 & \cdots & \lambda_2^{k-3} & \lambda_2^{k-2} \\ \vdots & \vdots & & \vdots & \vdots \\ 1 & \lambda_k & \cdots & \lambda_k^{k-3} & \lambda_k^{k-2} \end{pmatrix}$$

Repeating the above steps gives:

$$\begin{vmatrix} \lambda_1 & \lambda_2 & \cdots & \lambda_{k-1} & \lambda_k \\ \lambda_1^2 & \lambda_2^2 & \cdots & \lambda_{k-1}^2 & \lambda_k^2 \\ \vdots & \vdots & & \vdots & \vdots \\ \lambda_1^k & \lambda_2^k & \cdots & \lambda_{k-1}^k & \lambda_k^k \end{vmatrix} = \prod_{i=1}^k \lambda_i \cdot \prod_{i>\ell} (\lambda_i - \lambda_\ell)$$

Hence, if all λ_i 's are different, then the determinant is non-zero.

What happens if the roots are not all distinct?

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To see this consider the polynomial

$$P[\lambda] \cdot \lambda^{n-k} = c_0 \lambda^n + c_1 \lambda^{n-1} + c_2 \lambda^{n-2} + \dots + c_k \lambda^{n-k}$$



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Since λ_i is a root we can write this as $Q[\lambda] \cdot (\lambda - \lambda_i)^2$. Calculating the derivative gives a polynomial that still has root λ_i .

This means

$$c_0 n \lambda_i^{n-1} + c_1 (n-1) \lambda_i^{n-2} + \dots + c_k (n-k) \lambda_i^{n-k-1} = 0$$

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Hence,

$$c_0 \underbrace{n\lambda_i^n}_{T[n]} + c_1 \underbrace{(n-1)\lambda_i^{n-1}}_{T[n-1]} + \cdots + c_k \underbrace{(n-k)\lambda_i^{n-k}}_{T[n-k]} = 0$$

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Doing this again gives

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We can continue j-1 times.

Hence, $n^{\ell} \lambda_i^n$ is a solution for $\ell \in 0, ..., j-1$.

Lemma 7

Let $P[\lambda]$ denote the characteristic polynomial to the recurrence

$$c_0T[n] + c_1T[n-1] + \cdots + c_kT[n-k] = 0$$

Let λ_i , $i=1,\ldots,m$ be the (complex) roots of $P[\lambda]$ with multiplicities ℓ_i . Then the general solution to the recurrence is given by

$$T[n] = \sum_{i=1}^{m} \sum_{j=0}^{\ell_i - 1} \alpha_{ij} \cdot (n^j \lambda_i^n) .$$

The full proof is omitted. We have only shown that any choice of α_{ij} 's is a solution to the recurrence.

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 $T[1] = 1$
 $T[n] = T[n-1] + T[n-2]$ for $n \ge 2$

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Finding the roots, gives

$$\lambda_{1/2} = \frac{1}{2} \pm \sqrt{\frac{1}{4} + 1} = \frac{1}{2} \left(1 \pm \sqrt{5} \right)$$

$$\alpha \left(\frac{1+\sqrt{5}}{2}\right)^n + \beta \left(\frac{1-\sqrt{5}}{2}\right)^n$$

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 gives

$$\alpha\left(\frac{1+\sqrt{5}}{2}\right)+\beta\left(\frac{1-\sqrt{5}}{2}\right)=1 \Rightarrow \alpha-\beta=\frac{2}{\sqrt{5}}$$

Hence, the solution is

$$\frac{1}{\sqrt{5}} \left[\left(\frac{1+\sqrt{5}}{2} \right)^n - \left(\frac{1-\sqrt{5}}{2} \right)^n \right]$$

Consider the recurrence relation:

$$c_0T(n) + c_1T(n-1) + c_2T(n-2) + \cdots + c_kT(n-k) = f(n)$$

with $f(n) \neq 0$.

While we have a fairly general technique for solving homogeneous, linear recurrence relations the inhomogeneous case is different.

The general solution of the recurrence relation is

$$T(n) = T_h(n) + T_p(n) ,$$

where T_h is any solution to the homogeneous equation, and T_p is one particular solution to the inhomogeneous equation.

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There is no general method to find a particular solution.

Example:

$$T[n] = T[n-1] + 1$$
 $T[0] = 1$

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I get a completely determined recurrence if I add T[0] = 1 and T[1] = 2.

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$$\lambda^2 - 2\lambda + 1 = 0$$

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 gives $\alpha = 1$.

$$T[1] = 2$$
 gives $1 + \beta = 2 \Rightarrow \beta = 1$.

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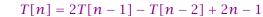
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$$T[n] = 3T[n-1] - 3T[n-2] + T[n-3] + 2$$
 and so on...

Definition 8 (Generating Function)

Let $(a_n)_{n\geq 0}$ be a sequence. The corresponding

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 exponential generating function (exponentielle Erzeugendenfunktion) is

$$F(z) := \sum_{n \ge 0} \frac{a_n}{n!} z^n .$$

Example 9

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There are no convergence issues here.

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Then, it is important to think about convergence/convergence radius etc.

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This is well-defined.

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Suppose we are given the generating function the known rules for differentiation

$$\sum_{n>0} z^n = \frac{1}{1-z}$$

We can compute the derivative:

$$\sum_{n \ge 1} nz^{n-1} = \frac{1}{(1-z)^2}$$

Hence, the generating function of the sequence $a_n = n + 1$ is $1/(1-z)^2$.

Formally the derivative of a formal power series
$$\sum_{n\geq 0} a_n z^n$$
 is defined as $\sum_{n\geq 0} na_n z^{n-1}$.

 $\sum_{n\geq 0} z^n = \frac{1}{1-z} \ . \ | \begin{array}{c} \text{ular, e.g. the derivative of } \frac{1}{1-z} \text{ is } \\ \frac{1}{(1-z)^2}. \end{array}$ work for this definition. In partic-

Note that this requires a proof if we consider power series as algebraic objects. However, we did not prove this in the lecture.

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Derivative:

$$\sum_{n>1} n(n+1)z^{n-1} = \frac{2}{(1-z)^3}$$

We can repeat this

$$\sum_{n \ge 0} (n+1) z^n = \frac{1}{(1-z)^2} \ .$$

Derivative:

$$\sum_{\substack{n \ge 1 \\ \sum_{n > 0} (n+1)(n+2)z^n}} n(n+1)z^{n-1} = \frac{2}{(1-z)^3}$$

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Hence, the generating function of the sequence $a_n = (n+1)(n+2)$ is $\frac{2}{(1-z)^3}$.

$$\sum_{n\geq k} n(n-1)\cdot\ldots\cdot(n-k+1)z^{n-k}$$

$$\sum_{n>k} n(n-1)\cdot\ldots\cdot(n-k+1)z^{n-k} = \sum_{n\geq 0} (n+k)\cdot\ldots\cdot(n+1)z^n$$

$$\sum_{n \ge k} n(n-1) \cdot \dots \cdot (n-k+1) z^{n-k} = \sum_{n \ge 0} (n+k) \cdot \dots \cdot (n+1) z^n$$
$$= \frac{k!}{(1-z)^{k+1}}.$$

Computing the k-th derivative of $\sum z^n$.

$$\sum_{n \ge k} n(n-1) \cdot \dots \cdot (n-k+1) z^{n-k} = \sum_{n \ge 0} (n+k) \cdot \dots \cdot (n+1) z^n$$
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The generating function of the sequence $a_n = \binom{n+k}{k}$ is $\frac{1}{(1-z)^{k+1}}$.

$$\sum_{n \ge 0} n z^n = \sum_{n \ge 0} (n+1) z^n - \sum_{n \ge 0} z^n$$

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The generating function of the sequence $a_n = n$ is $\frac{z}{(1-z)^2}$.

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The generating function of the sequence $f_n = a^n$ is $\frac{1}{1-az}$.

Suppose we have the recurrence $a_n = a_{n-1} + 1$ for $n \ge 1$ and $a_0 = 1$.

A(z)

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$$= zA(z) + \frac{1}{1 - z}$$

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Solving for A(z) gives

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Hence, $a_n = n + 1$.

n-th sequence element	generating function

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1	$\frac{1}{1-z}$
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n^2	$\frac{z(1+z)}{(1-z)^3}$
$\frac{1}{n!}$	e^z



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$\sum_{i=0}^{n} f_i$	$\frac{F(z)}{1-z}$

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- **6.** The coefficients of the resulting power series are the a_n .

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$$A(z) = \frac{(1-z)^2 + z}{(1-3z)(1-z)^2} = \frac{z^2 - z + 1}{(1-3z)(1-z)^2}$$

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This gives

$$z^{2} - z + 1 = A(1 - z)^{2} + B(1 - 3z)(1 - z) + C(1 - 3z)$$
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$$= (A + 3B)z^{2} + (-2A - 4B - 3C)z + (A + B + C)$$

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This leads to the following conditions:

$$A + B + C = 1$$
$$2A + 4B + 3C = 1$$
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Example:
$$a_n = 3a_{n-1} + n$$
, $a_0 = 1$

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This leads to the following conditions:

$$A + B + C = 1$$
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which gives

$$A = \frac{7}{4}$$
 $B = -\frac{1}{4}$ $C = -\frac{1}{2}$

$$A(z) = \frac{7}{4} \cdot \frac{1}{1 - 3z} - \frac{1}{4} \cdot \frac{1}{1 - z} - \frac{1}{2} \cdot \frac{1}{(1 - z)^2}$$

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$$= \frac{7}{4} \cdot \sum_{n \ge 0} 3^n z^n - \frac{1}{4} \cdot \sum_{n \ge 0} z^n - \frac{1}{2} \cdot \sum_{n \ge 0} (n + 1) z^n$$

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$$= \sum_{n \ge 0} \left(\frac{7}{4} \cdot 3^n - \frac{1}{4} - \frac{1}{2} (n + 1) \right) z^n$$

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5. Write f(z) as a formal power series:

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6. This means $a_n = \frac{7}{4}3^n - \frac{1}{2}n - \frac{3}{4}$.

Example 10

$$f_0 = 1$$

$$f_1 = 2$$

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Define

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 $g_1 = \log 2 = 1$ (for $\log = \log_2$), $g_0 = 0$

Example 10

$$f_0 = 1$$

$$f_1 = 2$$

$$f_n = f_{n-1} \cdot f_{n-2} \text{ for } n \ge 2 .$$

Define

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Example 11

$$f_1 = 1$$

 $f_n = 3f_{\frac{n}{2}} + n$; for $n = 2^k$, $k \ge 1$;

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.

Example 11

$$\begin{split} f_1 &= 1 \\ f_n &= 3f_{\frac{n}{2}} + n; \text{ for } n = 2^k, \ k \geq 1 \ ; \end{split}$$

Define

$$g_k := f_{2^k}$$
.

Then:

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Example 11

$$f_1 = 1$$

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Define

$$g_k := f_{2^k}$$
.

Then:

$$g_0 = 1$$

 $g_k = 3g_{k-1} + 2^k, \ k \ge 1$

$$g_k = 3\left[g_{k-1}\right] + 2^k$$

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Let
$$n = 2^k$$
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$$g_k = 3^{k+1} - 2^{k+1}$$
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 $= 3(2^k)^{\log 3} - 2 \cdot 2^k$
 $= 3n^{\log 3} - 2n$.

Part III

Data Structures



Abstract Data Type

An abstract data type (ADT) is defined by an interface of operations or methods that can be performed and that have a defined behavior.

The data types in this lecture all operate on objects that are represented by a [key, value] pair.

- The key comes from a totally ordered set, and we assume that there is an efficient comparison function.
- The value can be anything; it usually carries satellite information important for the application that uses the ADT.

▶ *S.* search(k): Returns pointer to object x from S with key[x] = k or null.

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- S. delete(x): Given pointer to object x from S, delete x from the set.
- S. minimum(): Return pointer to object with smallest key-value in S.
- S. maximum(): Return pointer to object with largest key-value in S.
- ► *S.* successor(*x*): Return pointer to the next larger element in *S* or null if *x* is maximum.

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- S. minimum(): Return pointer to object with smallest key-value in S.
- S. maximum(): Return pointer to object with largest key-value in S.
- S. successor(x): Return pointer to the next larger element in S or null if x is maximum.
- S. predecessor(x): Return pointer to the next smaller element in S or null if x is minimum.



▶ *S.* union(S'): Sets $S := S \cup S'$. The set S' is destroyed.

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```
S := \{x \in S \mid \text{key}[x] \le k\}, S' := \{x \in S \mid \text{key}[x] > k\}.
```

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- ► S. concatenate(S'): $S := S \cup S'$. Requires $\text{key}[S. \text{maximum}()] \le \text{key}[S'. \text{minimum}()]$.

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- ► *S*. concatenate(S'): $S := S \cup S'$. Requires key[S. maximum()] \leq key[S'. minimum()].
- ► *S.* decrease-key(x, k): Replace key[x] by $k \le key[x]$.

Examples of ADTs

Stack:

- \triangleright S. push(x): Insert an element.
- ▶ *S.* pop(): Return the element from *S* that was inserted most recently; delete it from *S*.
- ► *S.* empty(): Tell if *S* contains any object.

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Queue:

- S. enqueue(x): Insert an element.
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Queue:

- S. enqueue(x): Insert an element.
- S. dequeue(): Return the element that is longest in the structure; delete it from S.
- ► *S.* empty(): Tell if *S* contains any object.

Priority-Queue:

- S. insert(x): Insert an element.
- S. delete-min(): Return the element with lowest key-value; delete it from S.

7 Dictionary

Dictionary:

- S. insert(x): Insert an element x.
- **S.** delete(x): Delete the element pointed to by x.
- ▶ *S.* search(k): Return a pointer to an element e with key[e] = k in S if it exists; otherwise return null.

7.1 Binary Search Trees

An (internal) binary search tree stores the elements in a binary tree. Each tree-node corresponds to an element. All elements in the left sub-tree of a node v have a smaller key-value than $\ker[v]$ and elements in the right sub-tree have a larger-key value. We assume that all key-values are different.

(External Search Trees store objects only at leaf-vertices)

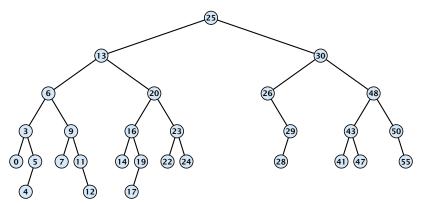
Examples:



7.1 Binary Search Trees

We consider the following operations on binary search trees. Note that this is a super-set of the dictionary-operations.

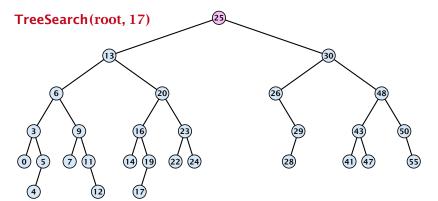
- ightharpoonup T. insert(x)
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Algorithm 1 TreeSearch(x, k)

1: **if** x = null or k = key[x] **return** x

2: **if** k < key[x] **return** TreeSearch(left[x], k)

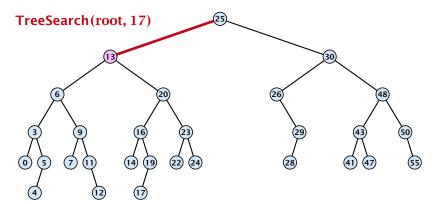


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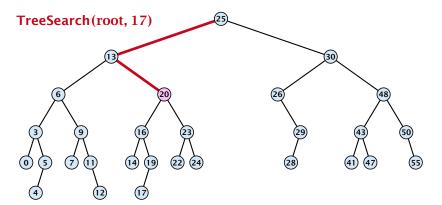


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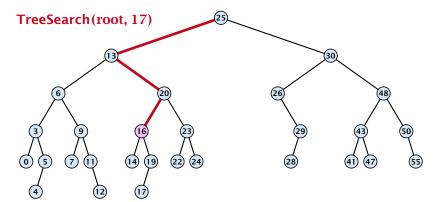


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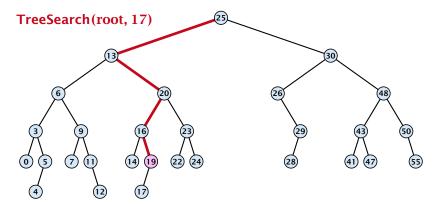


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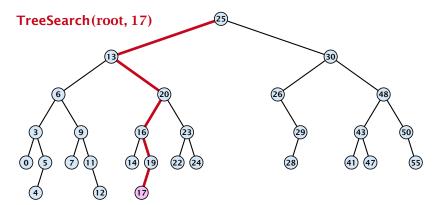


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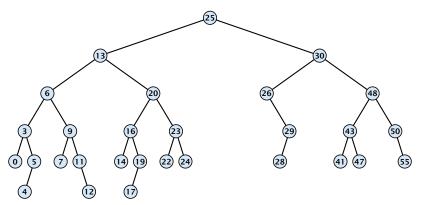


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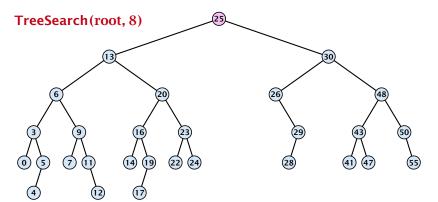




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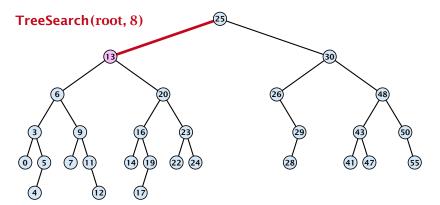
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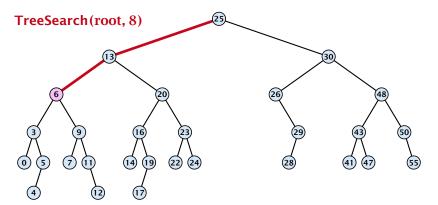
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- 3: **else return** TreeSearch(right[x], k)





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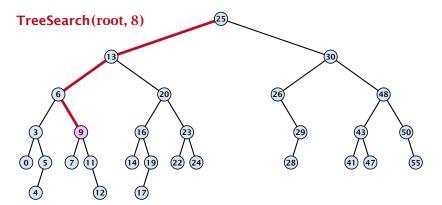


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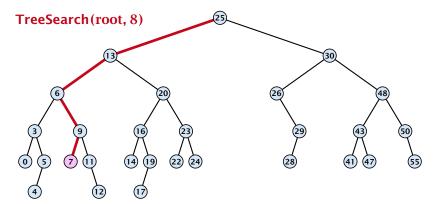


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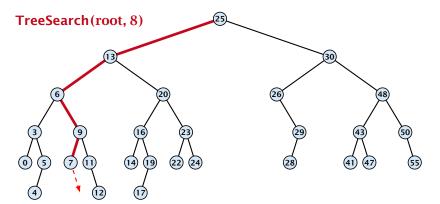


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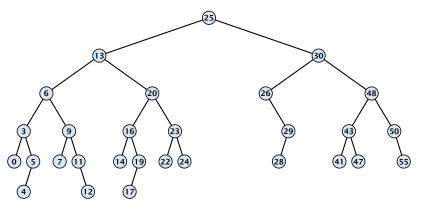


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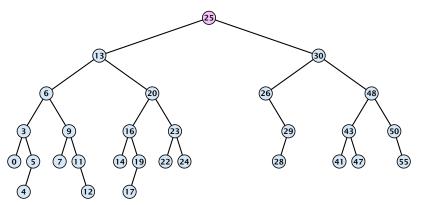
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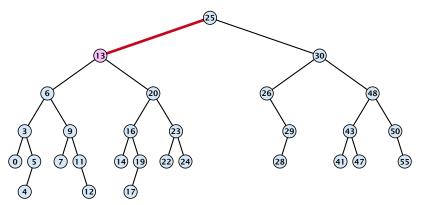
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- 2: **return** TreeMin(left[x])





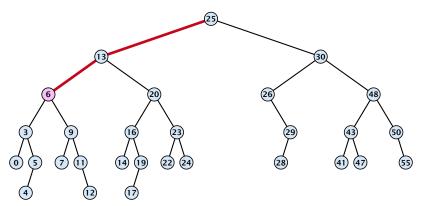
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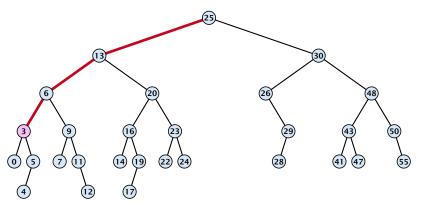
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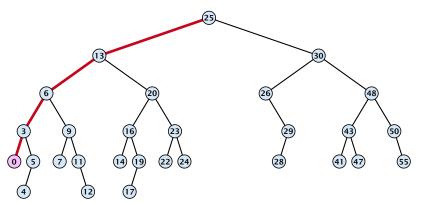


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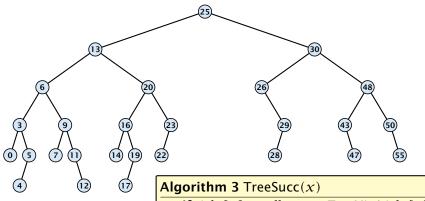


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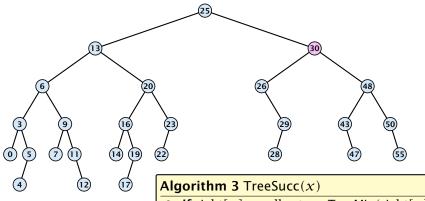
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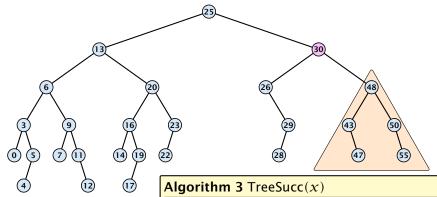
- 1: **if** right[x] \neq null **return** TreeMin(right[x])
- 2: $y \leftarrow \text{parent}[x]$
- 3: while $y \neq \text{null and } x = \text{right}[y]$ do
- 4: $x \leftarrow y; y \leftarrow \text{parent}[x]$
- 5: **return** y;





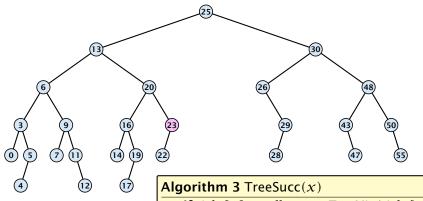
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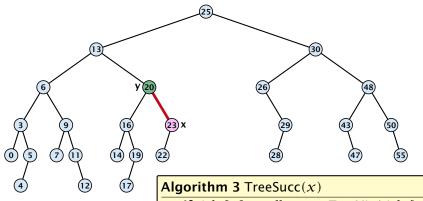
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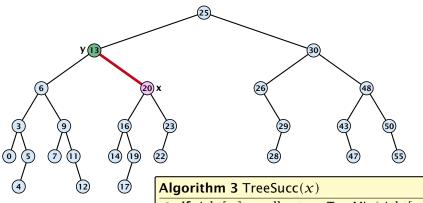
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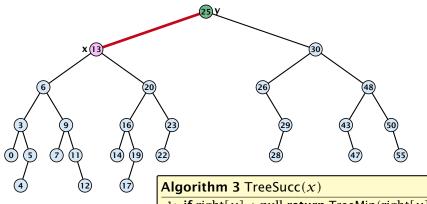
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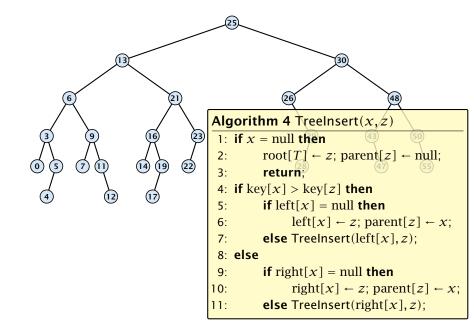
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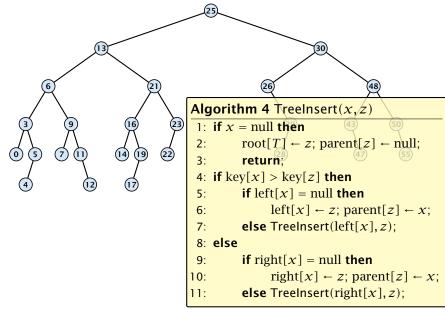


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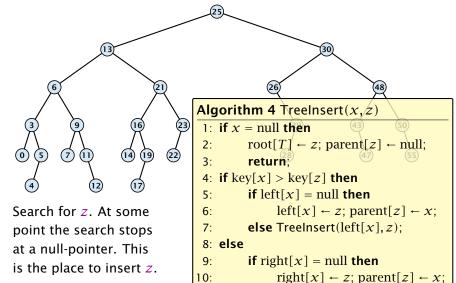




Insert element not in the tree.



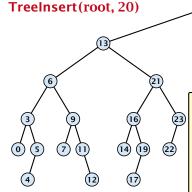
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11:

else Treelnsert(right[x], z);

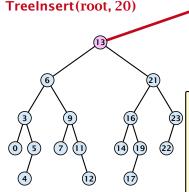
Insert element not in the tree.



Search for z. At some point the search stops at a null-pointer. This is the place to insert z.

- 1: if x = null then2: $\text{root}[T] \leftarrow z$; parent $[z] \leftarrow \text{null}$;
 - 3: return
- 4: **if** key[x] > key[z] **then**
- 5: **if** left[x] = null **then**
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- 6: $\operatorname{left}[x] \leftarrow z$; $\operatorname{parent}[z] \leftarrow x$; 7: $\operatorname{else} \operatorname{TreeInsert}(\operatorname{left}[x], z)$;
- 8: **else**
- 9: **if** right[x] = null **then**
- 10: $\operatorname{right}[x] \leftarrow z$; $\operatorname{parent}[z] \leftarrow x$;
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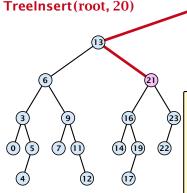
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- 1: if x = null then
 - root[T] $\leftarrow z$; parent[z] \leftarrow null; return:
- 4: **if** key[x] > key[z] **then**
- 5: **if** left[x] = null **then**
- 6: $\operatorname{left}[x] \leftarrow z$; parent $[z] \leftarrow x$;
- 7: **else** TreeInsert(left[x], z);
- 8: else
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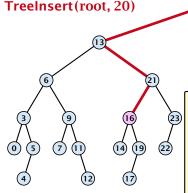
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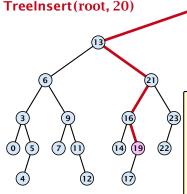
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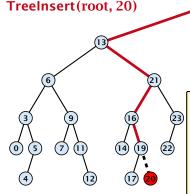
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Search for z. At some point the search stops at a null-pointer. This is the place to insert z.

Algorithm 4 TreeInsert(x,z)

1: if x = null then2: $\text{root}[T] \leftarrow z$; parent $[z] \leftarrow \text{null}$;

3: return

4: **if** key[x] > key[z] **then**

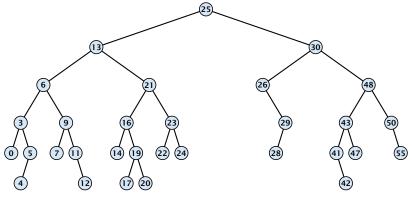
5: **if** left[x] = null **then**

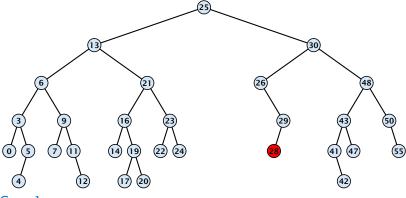
6: $\operatorname{left}[x] \leftarrow z$; parent $[z] \leftarrow x$;

7: **else** Treelnsert(left[x], z); 8: **else**

9: **if** right[x] = null **then**

10: $\operatorname{right}[x] \leftarrow z$; $\operatorname{parent}[z] \leftarrow x$; 11: $\operatorname{else} \operatorname{TreeInsert}(\operatorname{right}[x], z)$;

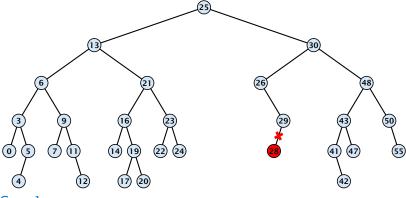




Case 1:

Element does not have any children

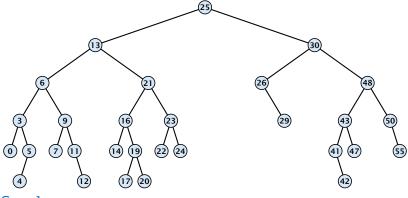
Simply go to the parent and set the corresponding pointer to null.



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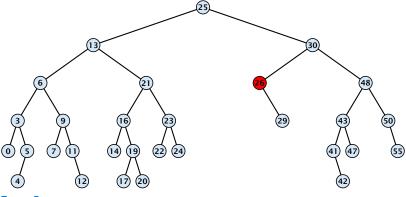
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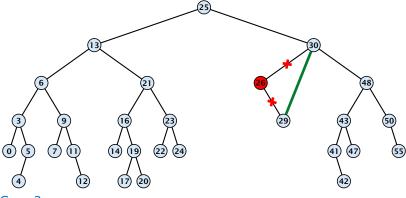
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Case 2:

Element has exactly one child

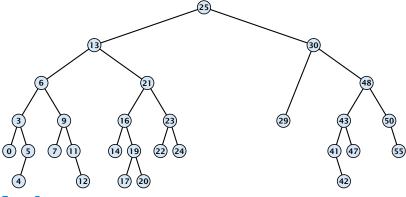
Splice the element out of the tree by connecting its parent to its successor.



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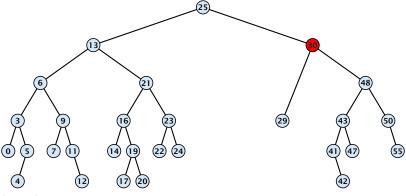
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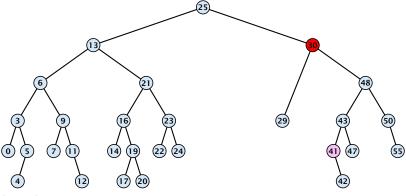
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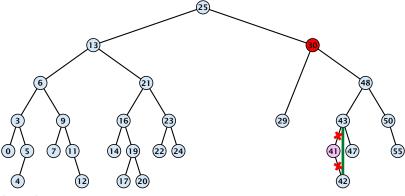
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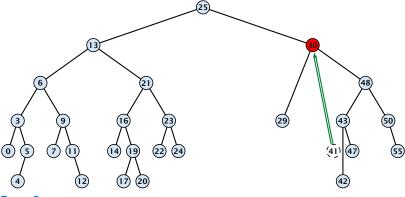
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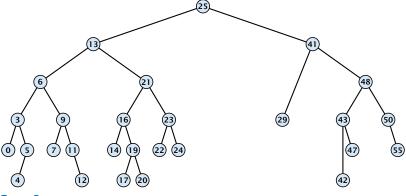
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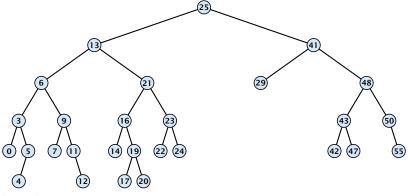
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```
Algorithm 9 TreeDelete(z)
 1: if left[z] = null or right[z] = null
          then \gamma \leftarrow z else \gamma \leftarrow \text{TreeSucc}(z); select \gamma to splice out
 3: if left[\gamma] \neq null
         then x \leftarrow \text{left}[y] else x \leftarrow \text{right}[y]; x is child of y (or null)
 5: if x \neq \text{null then parent}[x] \leftarrow \text{parent}[y]; parent[x] is correct
 6: if parent[\gamma] = null then
 7: root[T] \leftarrow x
 8: else
 9: if \gamma = \text{left[parent}[\gamma]] then
                                                                  fix pointer to x
10:
                left[parent[v]] \leftarrow x
    else
11:
12.
        right[parent[y]] \leftarrow x
13: if y \neq z then copy y-data to z
```

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With each insert- and delete-operation perform local adjustments to guarantee a height of $\mathcal{O}(\log n)$.

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AVL-trees, Red-black trees, Scapegoat trees, 2-3 trees, B-trees, AA trees, Treaps

similar: SPLAY trees.

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A red black tree is a balanced binary search tree in which each internal node has two children. Each internal node has a color, such that

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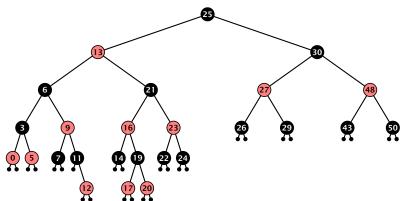
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Red Black Trees: Example



Lemma 13

A red-black tree with n internal nodes has height at most $\mathcal{O}(\log n)$.

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Lemma 13

A red-black tree with n internal nodes has height at most $O(\log n)$.

Definition 14

The black height bh(v) of a node v in a red black tree is the number of black nodes on a path from v to a leaf vertex (not counting v).

We first show:

Lemma 15

A sub-tree of black height bh(v) in a red black tree contains at least $2^{bh(v)}-1$ internal vertices.

Proof of Lemma 15.

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Induction on the height of v.

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- ▶ If height(v) (maximum distance btw. v and a node in the sub-tree rooted at v) is 0 then v is a leaf.
- ▶ The black height of v is 0.
- ► The sub-tree rooted at v contains $0 = 2^{bh(v)} 1$ inner vertices.

Proof (cont.)

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induction step

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- ► Then T_v contains at least $2(2^{\text{bh}(v)-1}-1)+1 \ge 2^{\text{bh}(v)}-1$ vertices.



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Hence, $h \le 2\log(n+1) = \mathcal{O}(\log n)$.



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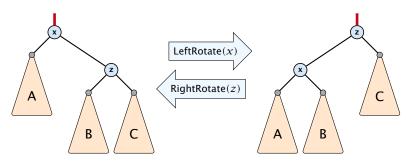
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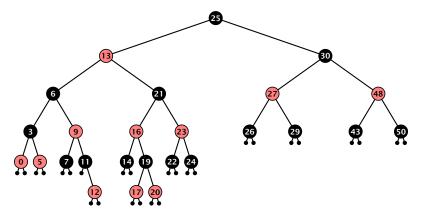
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We need to adapt the insert and delete operations so that the red black properties are maintained.

Rotations

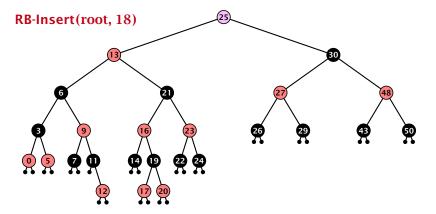
The properties will be maintained through rotations:





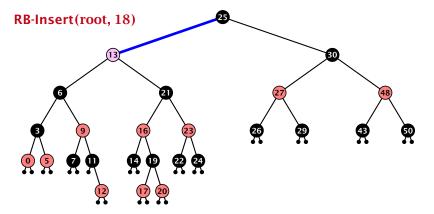
- first make a normal insert into a binary search tree
- then fix red-black properties





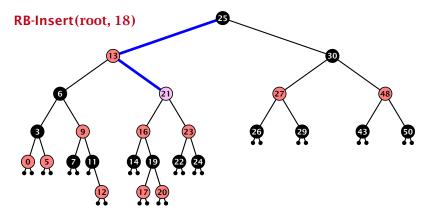
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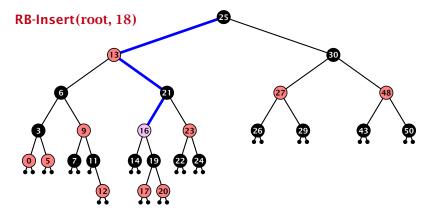
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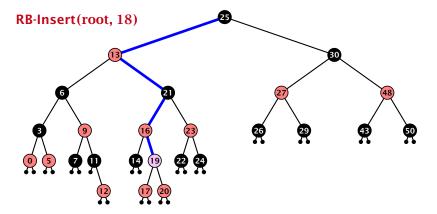
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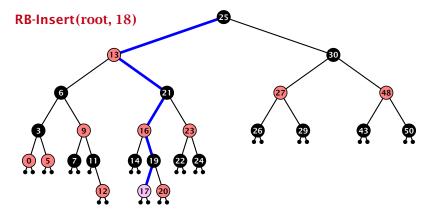
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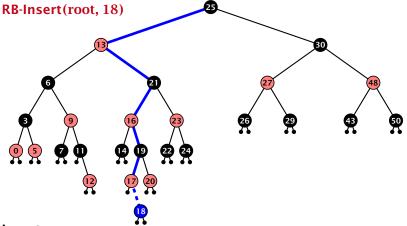
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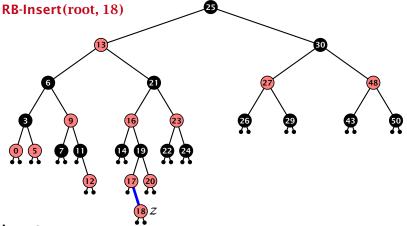
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- the only violation of red-black properties occurs at z and parent[z]
 - either both of them are red (most important case)
 - or the parent does not exist (violation since root must be black)

If z has a parent but no grand-parent we could simply color the parent/root black; however this case never happens.

```
Algorithm 10 InsertFix(z)
 1: while parent[z] \neq null and col[parent[z]] = red do
         if parent[z] = left[gp[z]] then
 2:
 3:
              uncle \leftarrow right[grandparent[z]]
             if col[uncle] = red then
 4:
                  col[p[z]] \leftarrow black; col[u] \leftarrow black;
 5:
                  col[gp[z]] \leftarrow red; z \leftarrow grandparent[z];
 6:
 7:
             else
                  if z = right[parent[z]] then
 8:
                       z \leftarrow p[z]; LeftRotate(z);
 9:
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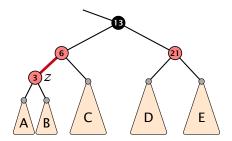
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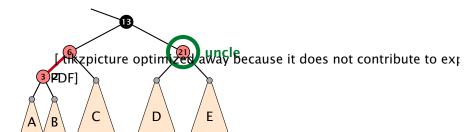
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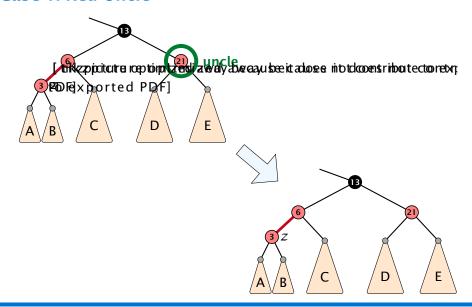
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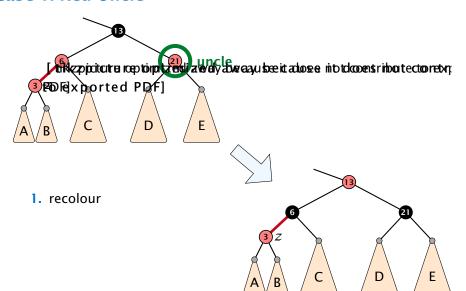
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 3:
              uncle \leftarrow right[grandparent[z]]
              if col[uncle] = red then
 4:
                  col[p[z]] \leftarrow black; col[u] \leftarrow black;
 5:
                  col[gp[z]] \leftarrow red; z \leftarrow grandparent[z];
 6:
              else
 7:
                  if z = right[parent[z]] then
 8:
                                                              2a: z right child
                       z \leftarrow p[z]; LeftRotate(z);
 9:
                  col[p[z]] \leftarrow black; col[gp[z]] \leftarrow red;
10:
                  RightRotate(gp[z]);
11:
12:
         else same as then-clause but right and left exchanged
13: col(root[T]) \leftarrow black;
```

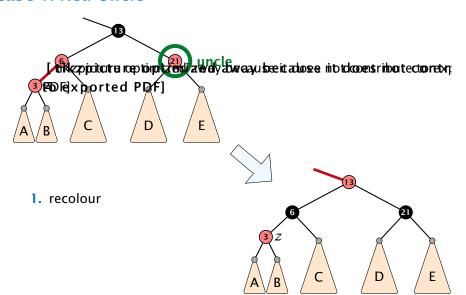
```
Algorithm 10 InsertFix(z)
 1: while parent[z] \neq null and col[parent[z]] = red do
         if parent[z] = left[gp[z]] then
 2:
 3:
              uncle \leftarrow right[grandparent[z]]
              if col[uncle] = red then
 4:
                  col[p[z]] \leftarrow black; col[u] \leftarrow black;
 5:
                  col[gp[z]] \leftarrow red; z \leftarrow grandparent[z];
 6:
             else
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                  if z = right[parent[z]] then
 8:
                       z \leftarrow p[z]; LeftRotate(z);
 9:
10:
                  col[p[z]] \leftarrow black; col[gp[z]] \leftarrow red; 2b: z left child
                  RightRotate(gp[z]);
11:
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13: col(root[T]) \leftarrow black;
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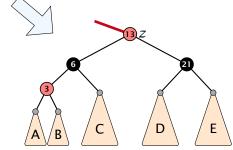








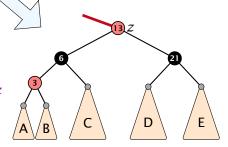
- 1. recolour
- 2. move z to grand-parent



Case 1: Red Uncle



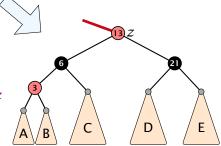
- 1. recolour
- 2. move z to grand-parent
- 3. invariant is fulfilled for new z

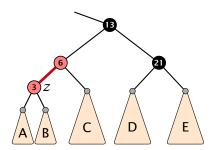


Case 1: Red Uncle

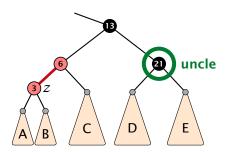


- 1. recolour
- 2. move z to grand-parent
- 3. invariant is fulfilled for new z
- 4. you made progress





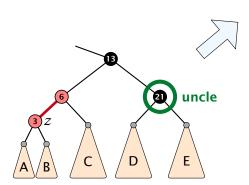
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1. rotate around grandparent

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7.2 Red Black Trees

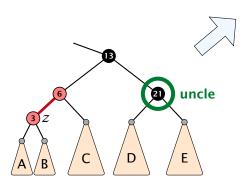
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7.2 Red Black Trees

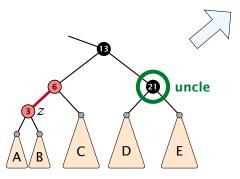
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1. rotate around grandparent

2. [reikzphoùchto repromptei eloliaterely aberaayu bee it actions en into toloroes into toloroes into toloroes into toloroes.

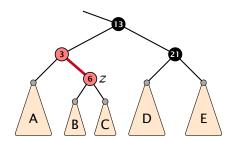
3. you have a red black tree



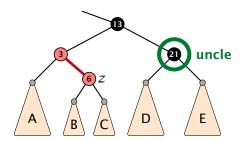


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1. rotate around parent [tikzpjatare reptino tżedi zwely zwezyu by tładoce into takcens riporecto retx Pole[xported PDF] D Ε uncle

1. rotate around parent 2. [minitezpizitalnewapvanpatzi erali zwedy abwezyu speritasto se ni o tako ensrindi ecto netr Pole xported PDF] D Ε uncle

1. rotate around parent 2. [minitezpizitaltrawapvanpatzi endi zverely abverazu soprifactore into taladens rindat ectoreter 3. POFFINE OF TEST PET D Ε uncle Α

Running time:

▶ Only Case 1 may repeat; but only h/2 many steps, where h is the height of the tree.

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Performing Case 1 at most $\mathcal{O}(\log n)$ times and every other case at most once, we get a red-black tree. Hence $\mathcal{O}(\log n)$ re-colorings and at most 2 rotations.

First do a standard delete.

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If the spliced out node \boldsymbol{x} was red everything is fine.

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If it was black there may be the following problems.

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Parent and child of x were red; two adjacent red vertices.

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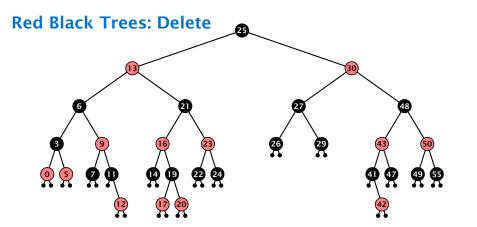
- Parent and child of x were red; two adjacent red vertices.
- If you delete the root, the root may now be red.

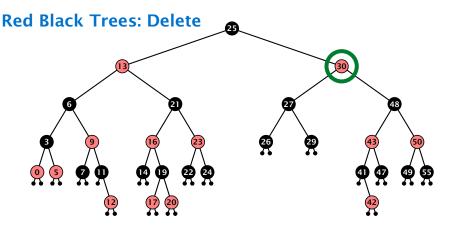
First do a standard delete.

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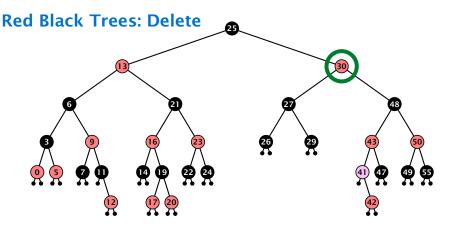
- Parent and child of x were red; two adjacent red vertices.
- If you delete the root, the root may now be red.
- Every path from an ancestor of x to a descendant leaf of x changes the number of black nodes. Black height property might be violated.





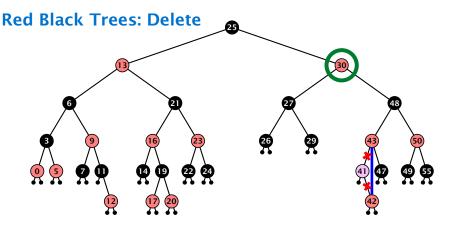
Case 3:

- do normal delete
- when replacing content by content of successor, don't change color of node



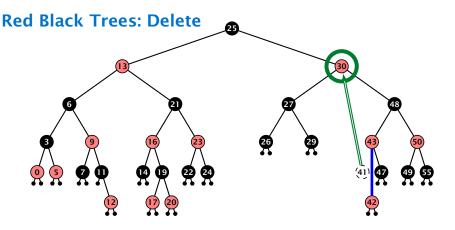
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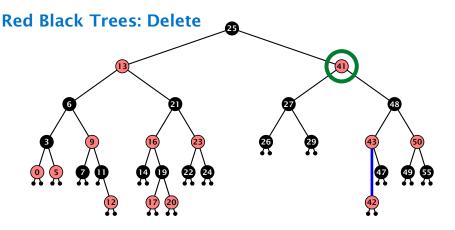
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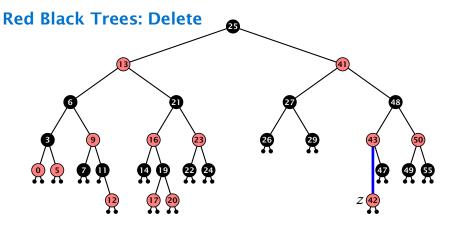
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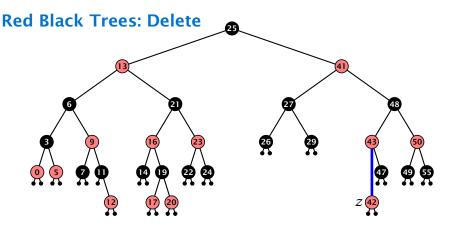
Case 3:

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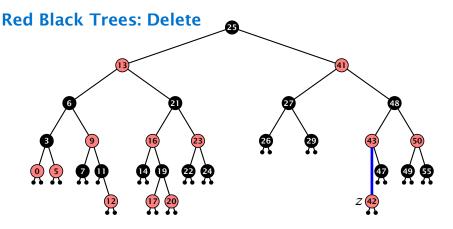
Delete:

deleting black node messes up black-height property



Delete:

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- ightharpoonup if z is red, we can simply color it black and everything is fine



Delete:

- deleting black node messes up black-height property
- ightharpoonup if z is red, we can simply color it black and everything is fine
- the problem is if z is black (e.g. a dummy-leaf); we call a fix-up procedure to fix the problem.

Invariant of the fix-up algorithm

► the node z is black

Invariant of the fix-up algorithm

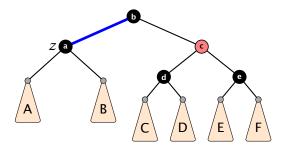
- ► the node z is black
- if we "assign" a fake black unit to the edge from z to its parent then the black-height property is fulfilled

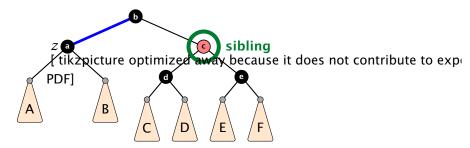
Invariant of the fix-up algorithm

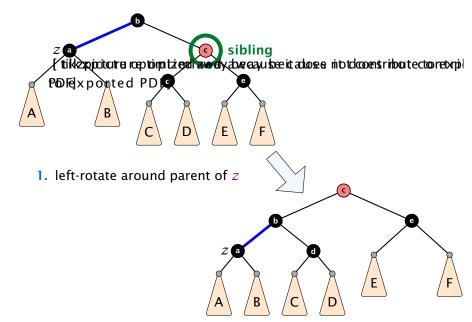
- ▶ the node *z* is black
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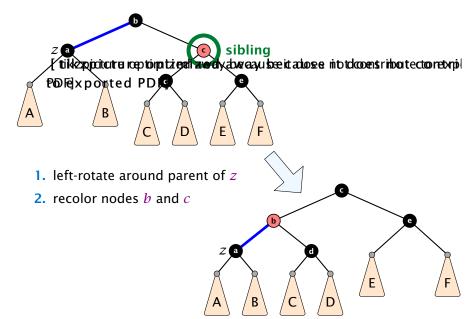
Goal: make rotations in such a way that you at some point can remove the fake black unit from the edge.

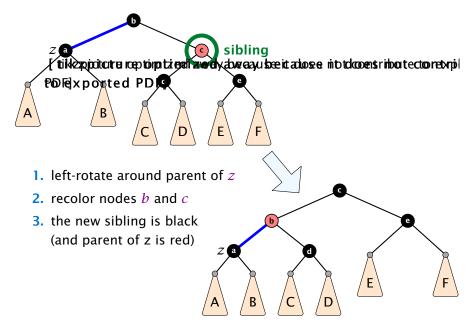
Case 1: Sibling of z is red

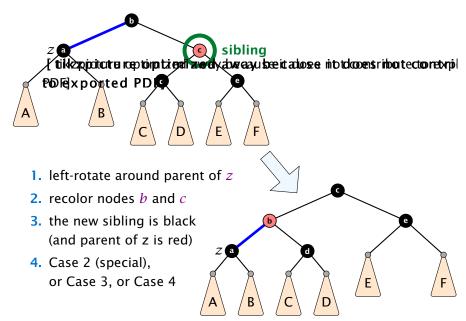


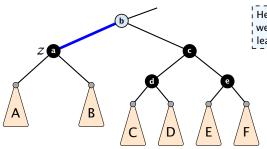




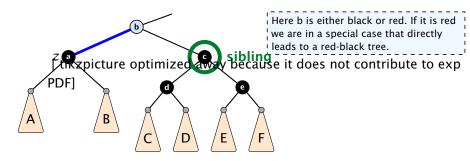


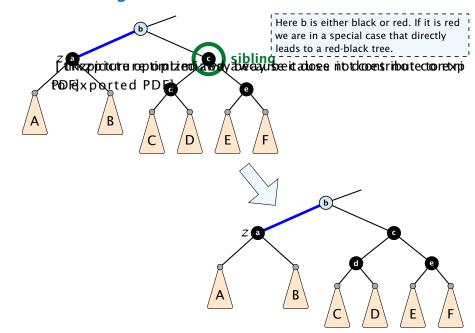


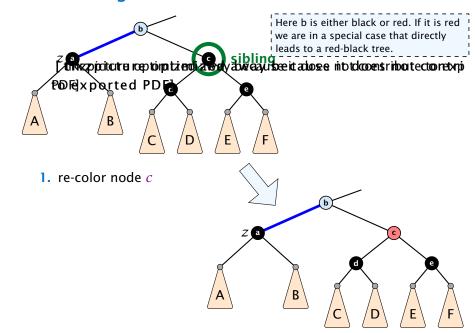


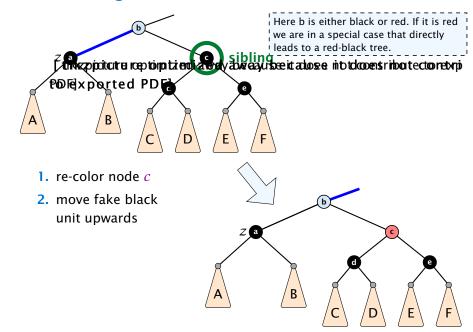


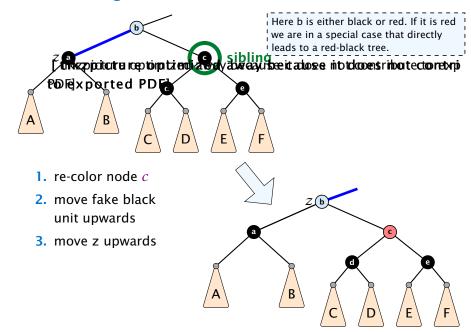
Here b is either black or red. If it is red we are in a special case that directly leads to a red-black tree.

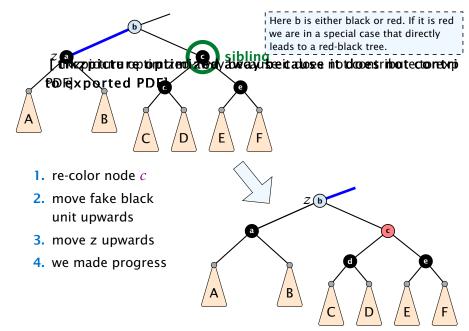


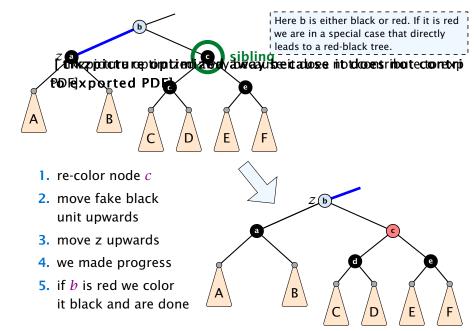


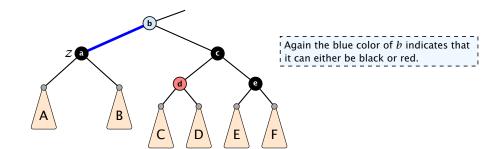




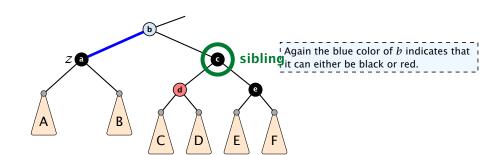


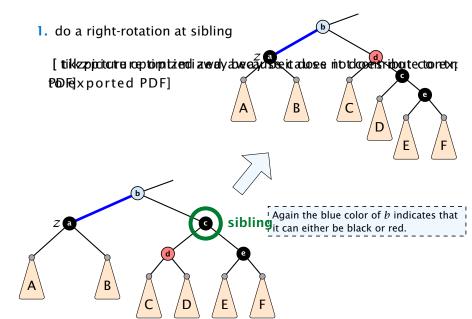


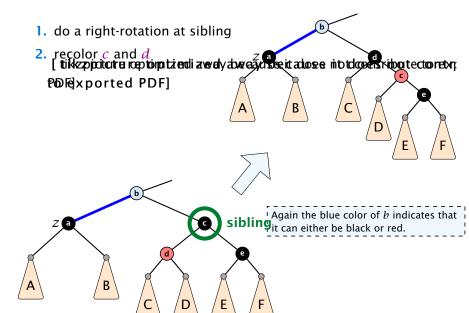


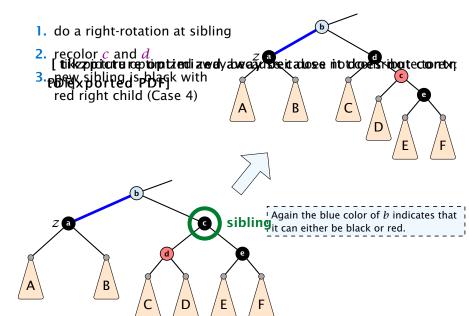


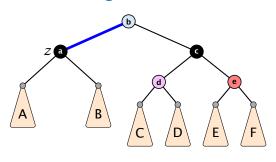
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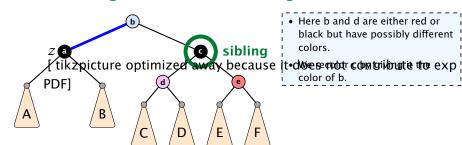


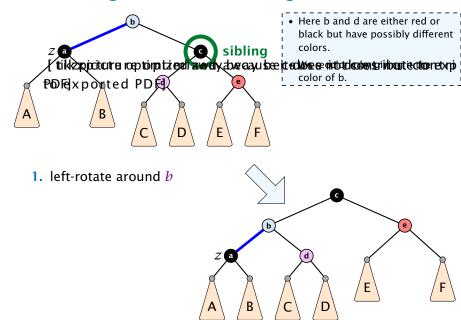


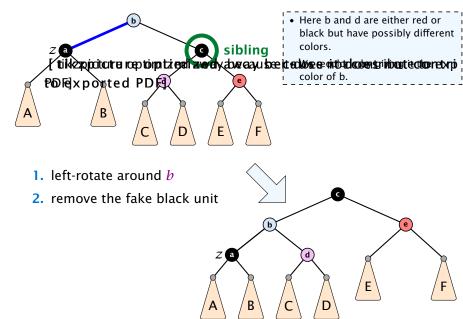


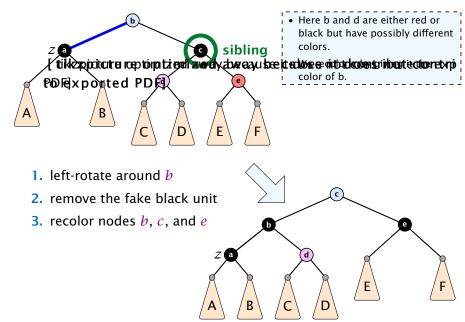


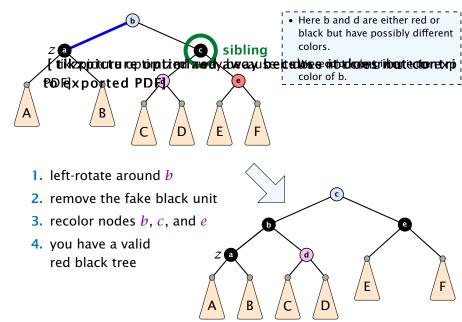
- Here b and d are either red or black but have possibly different colors.
- We recolor c by giving it the color of b.











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Performing Case 2 at most $\mathcal{O}(\log n)$ times and every other step at most once, we get a red black tree. Hence, $\mathcal{O}(\log n)$ re-colorings and at most 3 rotations.

Disadvantage of balanced search trees:

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- worst case; no advantage for easy inputs

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Splay Trees:

+ after access, an element is moved to the root; splay(x) repeated accesses are faster

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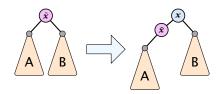
- + after access, an element is moved to the root; splay(x) repeated accesses are faster
- only amortized guarantee
- read-operations change the tree

find(x)

- ightharpoonup search for x according to a search tree
- let \bar{x} be last element on search-path
- ightharpoonup splay (\bar{x})

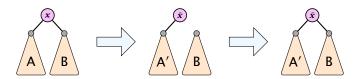
insert(x)

- search for x; \bar{x} is last visited element during search (successer or predecessor of x)
- splay(\bar{x}) moves \bar{x} to the root
- insert x as new root

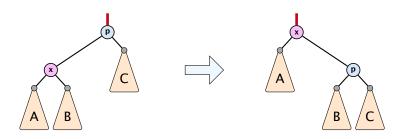


delete(x)

- search for x; splay(x); remove x
- **>** search largest element \bar{x} in A
- splay(\bar{x}) (on subtree A)
- connect root of B as right child of \bar{x}



Move to Root



How to bring element to root?

- one (bad) option: moveToRoot(x)
- iteratively do rotation around parent of x until x is root
- if x is left child do right rotation otw. left rotation

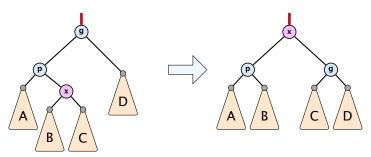
Splay: Zig Case



better option splay(x):

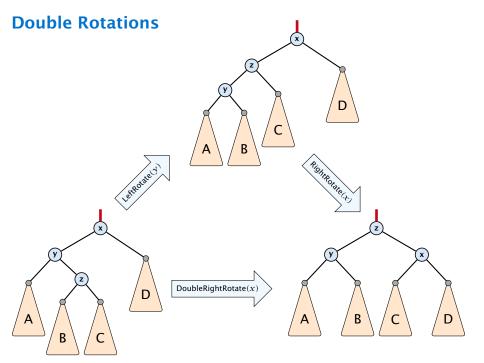
zig case: if x is child of root do left rotation or right rotation around parent

Splay: Zigzag Case

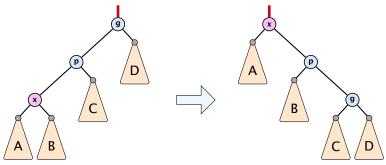


better option splay(x):

- zigzag case: if x is right child and parent of x is left child (or x left child parent of x right child)
- do double right rotation around grand-parent (resp. double left rotation)

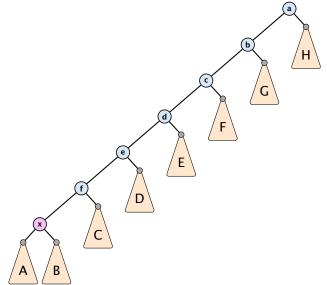


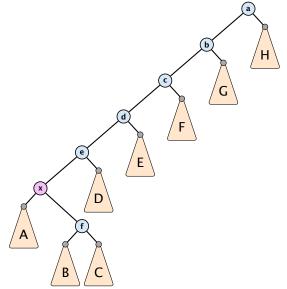
Splay: Zigzig Case

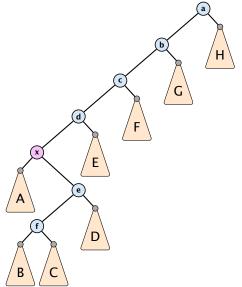


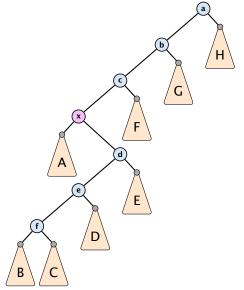
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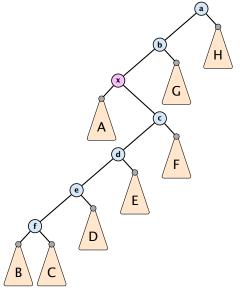
- zigzig case: if x is left child and parent of x is left child (or x right child, parent of x right child)
- do right roation around grand-parent followed by right rotation around parent (resp. left rotations)

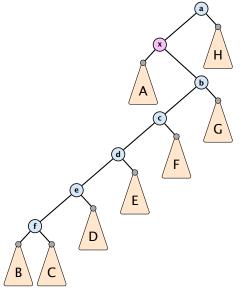


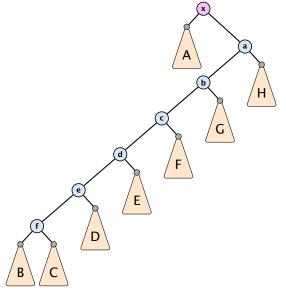


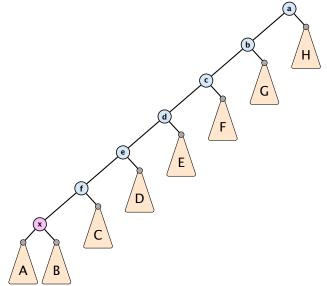




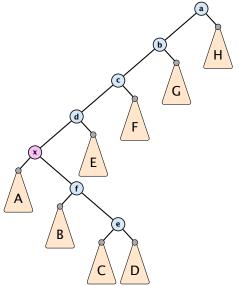


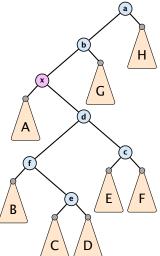


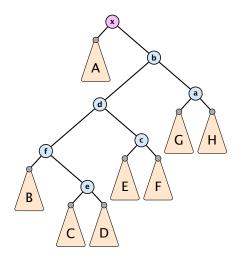




7.3 Splay Trees







Static Optimality

Suppose we have a sequence of m find-operations. find(x) appears h_x times in this sequence.

The cost of a static search tree *T* is:

$$cost(T) = m + \sum_{x} h_{x} \operatorname{depth}_{T}(x)$$

The total cost for processing the sequence on a splay-tree is $\mathcal{O}(\cos(T_{\min}))$, where T_{\min} is an optimal static search tree.

Dynamic Optimality

Let S be a sequence with m find-operations.

Let A be a data-structure based on a search tree:

- the cost for accessing element x is 1 + depth(x);
- after accessing x the tree may be re-arranged through rotations;

Conjecture:

A splay tree that only contains elements from S has cost $\mathcal{O}(\cos(A,S))$, for processing S.

Lemma 16

Splay Trees have an amortized running time of $O(\log n)$ for all operations.

Amortized Analysis

Definition 17

A data structure with operations $op_1(), \ldots, op_k()$ has amortized running times t_1, \ldots, t_k for these operations if the following holds.

Suppose you are given a sequence of operations (starting with an empty data-structure) that operate on at most n elements, and let k_i denote the number of occurences of $\operatorname{op}_i()$ within this sequence. Then the actual running time must be at most $\sum_i k_i \cdot t_i(n)$.

6. Feb. 2022

Introduce a potential for the data structure.

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• $\Phi(D_i)$ is the potential after the *i*-th operation.

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- $\Phi(D_i)$ is the potential after the *i*-th operation.
- ightharpoonup Amortized cost of the *i*-th operation is

$$\hat{c}_i = c_i + \Phi(D_i) - \Phi(D_{i-1}) \ . \label{eq:constraint}$$

7.3 Splay Trees

Introduce a potential for the data structure.

- $\Phi(D_i)$ is the potential after the *i*-th operation.
- ightharpoonup Amortized cost of the i-th operation is

$$\hat{c}_i = c_i + \Phi(D_i) - \Phi(D_{i-1}) .$$

▶ Show that $\Phi(D_i) \ge \Phi(D_0)$.

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► Show that $\Phi(D_i) \ge \Phi(D_0)$.

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$$\sum_{i=1}^{k} c_i \le \sum_{i=1}^{k} c_i + \Phi(D_k) - \Phi(D_0)$$

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▶ Show that $\Phi(D_i) \ge \Phi(D_0)$.

Then

$$\sum_{i=1}^k c_i \leq \sum_{i=1}^k c_i + \Phi(D_k) - \Phi(D_0) = \sum_{i=1}^k \hat{c}_i$$

This means the amortized costs can be used to derive a bound on the total cost.

Stack

- **►** *S.* push()
- **►** *S.* pop()
- ► *S.* multipop(*k*): removes *k* items from the stack. If the stack currently contains less than *k* items it empties the stack.
- ► The user has to ensure that pop and multipop do not generate an underflow.

Stack

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- ► The user has to ensure that pop and multipop do not generate an underflow.

Actual cost:

- **►** *S.* push(): cost 1.
- ► *S.* pop(): cost 1.
- *S.* multipop(k): cost min{size, k} = k.

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$$\hat{C}_{\text{pop}} = C_{\text{pop}} + \Delta \Phi = 1 - 1 \le 0 \ .$$

Example: Stack

Use potential function $\Phi(S) = \text{number of elements on the stack.}$

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$$\hat{C}_{\text{push}} = C_{\text{push}} + \Delta \Phi = 1 + 1 \le 2 .$$

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$$\hat{C}_{\mathrm{pop}} = C_{\mathrm{pop}} + \Delta \Phi = 1 - 1 \leq 0 \ .$$

 \triangleright S. multipop(k): cost

$$\hat{C}_{mn} = C_{mn} + \Delta \Phi = \min\{\text{size}, k\} - \min\{\text{size}, k\} \le 0$$
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Incrementing a binary counter:

Consider a computational model where each bit-operation costs one time-unit.

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Actual cost:

- ► Changing bit from 0 to 1: cost 1.
- Changing bit from 1 to 0: cost 1.
- ▶ Increment: cost is k + 1, where k is the number of consecutive ones in the least significant bit-positions (e.g, 001101 has k = 1).

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Choose potential function $\Phi(x)=k$, where k denotes the number of ones in the binary representation of x.

Amortized cost:

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$$\hat{C}_{0\to 1} = C_{0\to 1} + \Delta \Phi = 1 + 1 \le 2 .$$

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► Changing bit from 1 to 0:

$$\hat{C}_{1\to 0} = C_{1\to 0} + \Delta \Phi = 1 - 1 \le 0 .$$

▶ Increment: Let k denotes the number of consecutive ones in the least significant bit-positions. An increment involves k (1 \rightarrow 0)-operations, and one (0 \rightarrow 1)-operation.

Hence, the amortized cost is $k\hat{C}_{1\rightarrow 0} + \hat{C}_{0\rightarrow 1} \leq 2$.

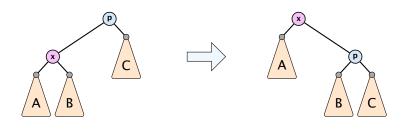
Splay Trees

potential function for splay trees:

- ightharpoonup size $s(x) = |T_x|$
- $rank r(x) = \log_2(s(x))$

amortized cost = real cost + potential change

The cost is essentially the cost of the splay-operation, which is 1 plus the number of rotations.



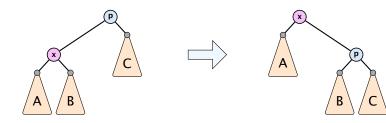
 $\Delta\Phi =$



$$\Delta\Phi = r'(x) + r'(p) - r(x) - r(p)$$



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$$= r'(p) - r(x)$$

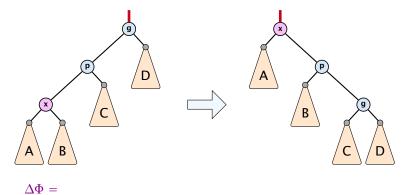


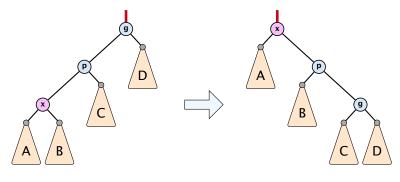
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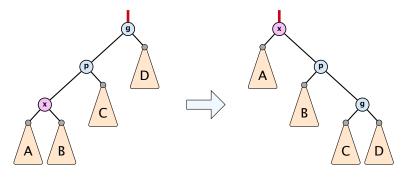
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$$\mathsf{cost}_{\mathsf{zig}} \leq 1 + 3(r'(x) - r(x))$$



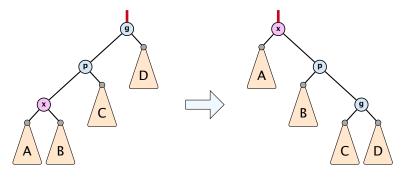


$$\Delta\Phi=r'(x)+r'(p)+r'(g)-r(x)-r(p)-r(g)$$



$$\Delta \Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

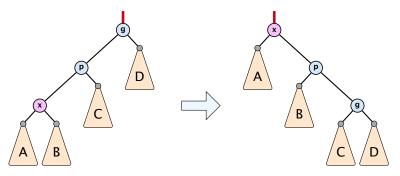
= $r'(p) + r'(g) - r(x) - r(p)$



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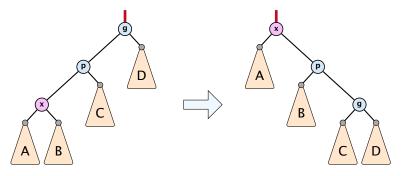


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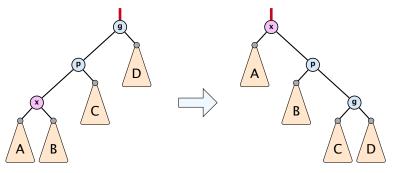
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$$= -2r'(x) + r'(g) + r(x) + 3(r'(x) - r(x))$$



$$\Delta\Phi = r'(x) + r'(p) + r'(g) - r(x) - r(p) - r(g)$$

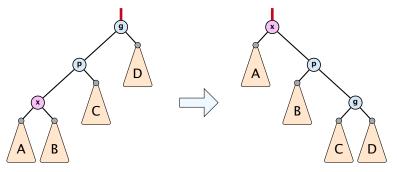
$$= r'(p) + r'(g) - r(x) - r(p)$$

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$$\leq -2 + 3(r'(x) - r(x))$$



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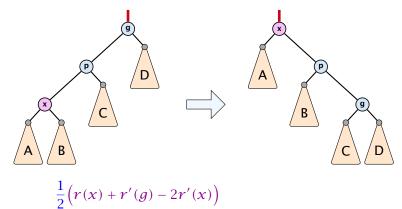
$$= r'(p) + r'(g) - r(x) - r(p)$$

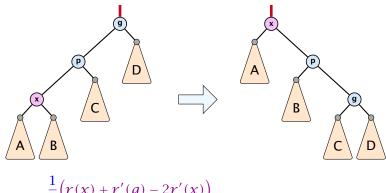
$$\leq r'(x) + r'(g) - r(x) - r(x)$$

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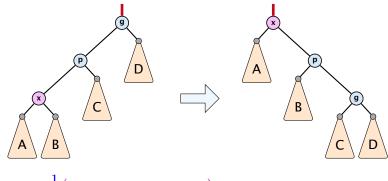
$$= -2r'(x) + r'(g) + r(x) + 3(r'(x) - r(x))$$

$$\leq -2 + 3(r'(x) - r(x)) \Rightarrow \cos t_{ziqziq} \leq 3(r'(x) - r(x))$$





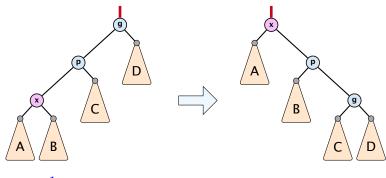
$$\frac{1}{2} \Big(r(x) + r'(g) - 2r'(x) \Big) \\
= \frac{1}{2} \Big(\log(s(x)) + \log(s'(g)) - 2\log(s'(x)) \Big)$$



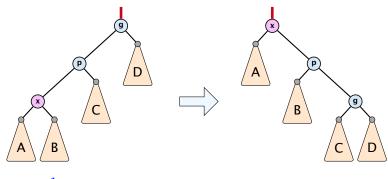
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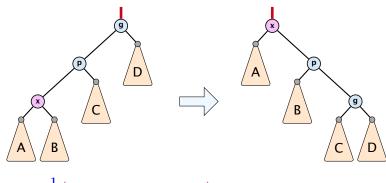
$$= \frac{1}{2} \log \left(\frac{s(x)}{s'(x)} \right) + \frac{1}{2} \log \left(\frac{s'(g)}{s'(x)} \right)$$



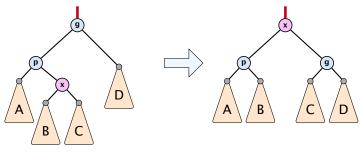
$$\begin{split} &\frac{1}{2}\Big(r(x) + r'(g) - 2r'(x)\Big) \\ &= \frac{1}{2}\Big(\log(s(x)) + \log(s'(g)) - 2\log(s'(x))\Big) \\ &= \frac{1}{2}\log\Big(\frac{s(x)}{s'(x)}\Big) + \frac{1}{2}\log\Big(\frac{s'(g)}{s'(x)}\Big) \\ &\leq \log\Big(\frac{1}{2}\frac{s(x)}{s'(x)} + \frac{1}{2}\frac{s'(g)}{s'(x)}\Big) \end{split}$$



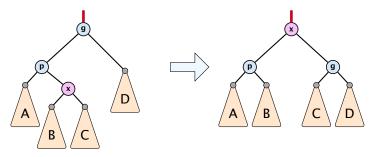
$$\begin{split} &\frac{1}{2}\Big(r(x)+r'(g)-2r'(x)\Big)\\ &=\frac{1}{2}\Big(\log(s(x))+\log(s'(g))-2\log(s'(x))\Big)\\ &=\frac{1}{2}\log\Big(\frac{s(x)}{s'(x)}\Big)+\frac{1}{2}\log\Big(\frac{s'(g)}{s'(x)}\Big)\\ &\leq\log\Big(\frac{1}{2}\frac{s(x)}{s'(x)}+\frac{1}{2}\frac{s'(g)}{s'(x)}\Big)\leq\log\Big(\frac{1}{2}\Big) \end{split}$$



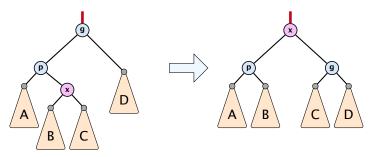
$$\begin{split} \frac{1}{2} \Big(r(x) + r'(g) - 2r'(x) \Big) \\ &= \frac{1}{2} \Big(\log(s(x)) + \log(s'(g)) - 2\log(s'(x)) \Big) \\ &= \frac{1}{2} \log \Big(\frac{s(x)}{s'(x)} \Big) + \frac{1}{2} \log \Big(\frac{s'(g)}{s'(x)} \Big) \\ &\leq \log \Big(\frac{1}{2} \frac{s(x)}{s'(x)} + \frac{1}{2} \frac{s'(g)}{s'(x)} \Big) \leq \log \Big(\frac{1}{2} \Big) = -1 \end{split}$$





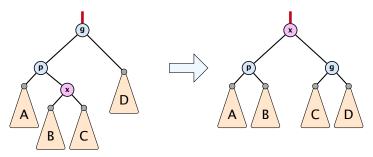


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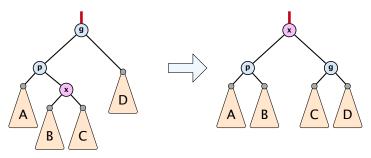
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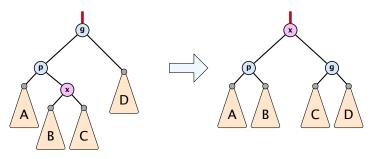


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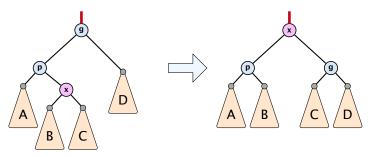
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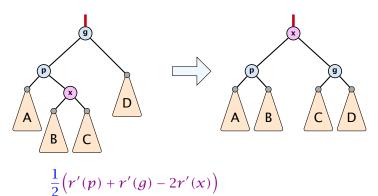
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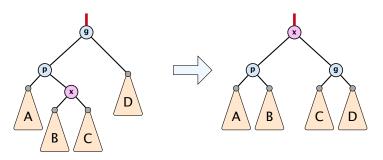
$$= r'(p) + r'(g) - r(x) - r(p)$$

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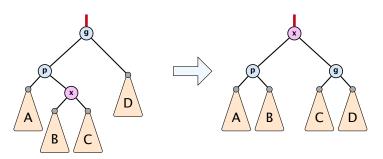
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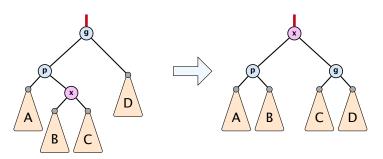




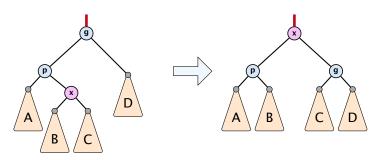
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Amortized cost of the whole splay operation:

$$\leq 1 + 1 + \sum_{\text{steps } t} 3(r_t(x) - r_{t-1}(x))$$

$$= 2 + 3(r(\text{root}) - r_0(x))$$

$$\leq \mathcal{O}(\log n)$$

Suppose you want to develop a data structure with:

- Insert(x): insert element x.
- **Search**(k): search for element with key k.
- **Delete**(x): delete element referenced by pointer x.
- ▶ find-by-rank(ℓ): return the ℓ -th element; return "error" if the data-structure contains less than ℓ elements.

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Augment an existing data-structure instead of developing a new one.

How to augment a data-structure

1. choose an underlying data-structure

- Of course, the above steps heavily depend on each other. For example it makes no sense to choose additional information to be stored (Step 2), and later realize that either the information cannot be maintained efficiently (Step 3) or is not sufficient to support the new operations (Step 4).
- However, the above outline is a good way to describe/document a new data-structure.

How to augment a data-structure

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- determine additional information to be stored in the underlying structure

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- verify/show how the additional information can be maintained for the basic modifying operations on the underlying structure.
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How to augment a data-structure

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- 3. verify/show how the additional information can be maintained for the basic modifying operations on the underlying structure.
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- However, the above outline is a good way to describe/document a new data-structure.

Goal: Design a data-structure that supports insert, delete, search, and find-by-rank in time $O(\log n)$.

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- 1. We choose a red-black tree as the underlying data-structure.
- 2. We store in each node v the size of the sub-tree rooted at v.
- 3. We need to be able to update the size-field in each node without asymptotically affecting the running time of insert, delete, and search. We come back to this step later...

Goal: Design a data-structure that supports insert, delete, search, and find-by-rank in time $O(\log n)$.

4. How does find-by-rank work? Find-by-rank(k) = Select(root,k) with

```
Algorithm 1 Select(x, i)
```

```
1: if x = \text{null} then return error
```

2: **if** $left[x] \neq null$ **then** $r \leftarrow left[x]$. size + 1 **else** $r \leftarrow 1$

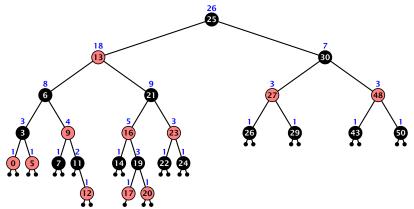
3: **if** i = r **then return** x

4: if i < r then

5: **return** Select(left[x], i)

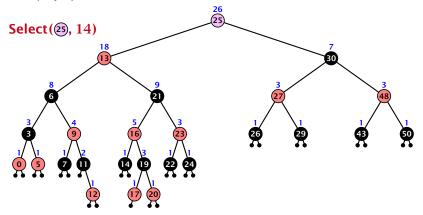
6: **else**

7: **return** Select(right[x], i - r)



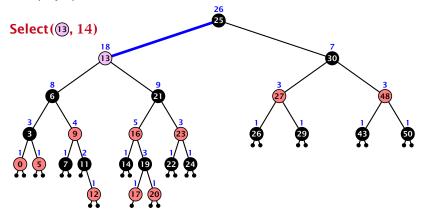
- decide whether you have to proceed into the left or right sub-tree
- adjust the rank that you are searching for if you go right





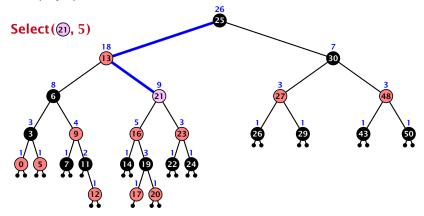
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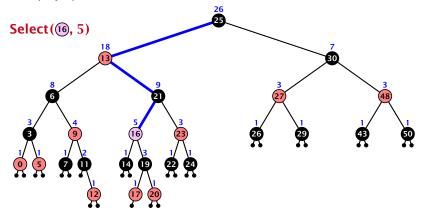
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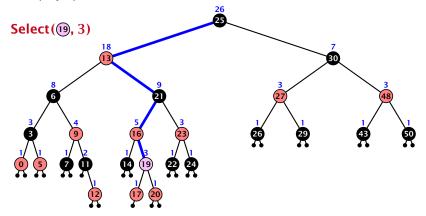
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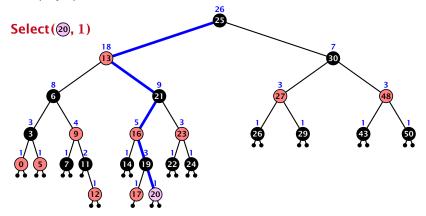
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Goal: Design a data-structure that supports insert, delete, search, and find-by-rank in time $\mathcal{O}(\log n)$.

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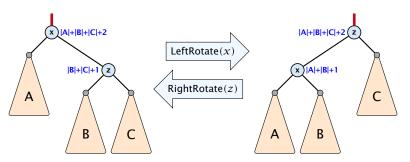
Search(k): Nothing to do.

Insert(x): When going down the search path increase the size field for each visited node. Maintain the size field during rotations.

Delete(*x*): Directly after splicing out a node traverse the path from the spliced out node upwards, and decrease the size counter on every node on this path. Maintain the size field during rotations.

Rotations

The only operation during the fix-up procedure that alters the tree and requires an update of the size-field:

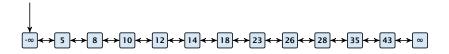


The nodes x and z are the only nodes changing their size-fields.

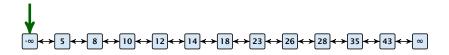
The new size-fields can be computed locally from the size-fields of the children.

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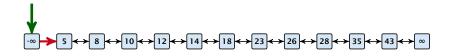
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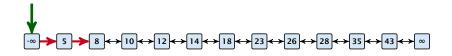
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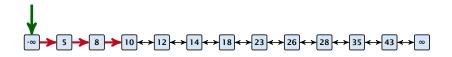
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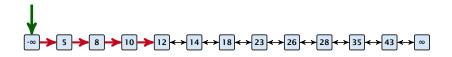
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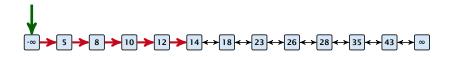


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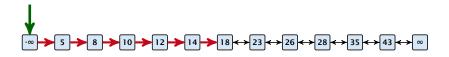
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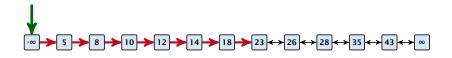
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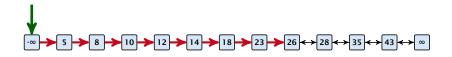
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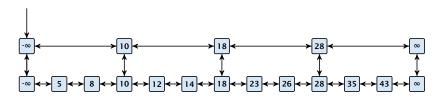
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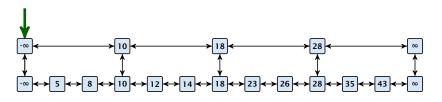
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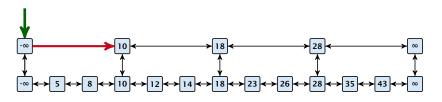
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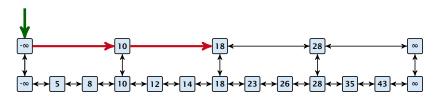
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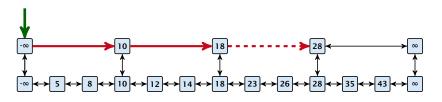
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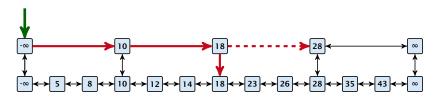
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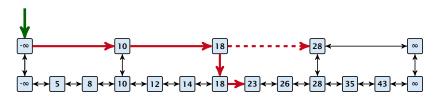
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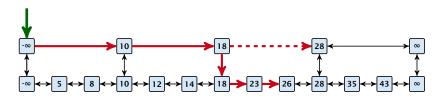
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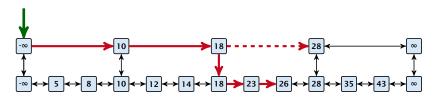


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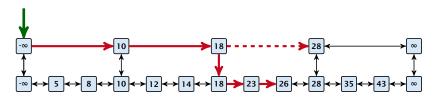
Add an express lane:



Let $|L_1|$ denote the number of elements in the "express lane", and $|L_0|=n$ the number of all elements (ignoring dummy elements).

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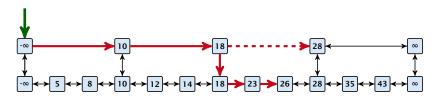


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Choose $|L_1| = \sqrt{n}$. Then search time $\Theta(\sqrt{n})$.

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6. Feb. 2022 205/616

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6. Feb. 2022

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- **.**..
- ► At most $|L_k| + \sum_{i=1}^k \frac{L_{i-1}}{L_i} + 3(k+1)$ steps.

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Choosing $k = \Theta(\log n)$ gives a logarithmic running time.

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Use randomization instead!

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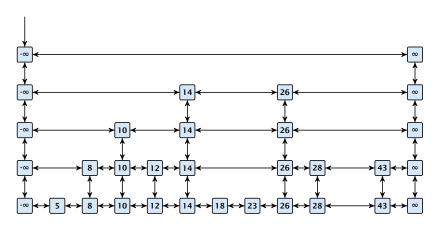
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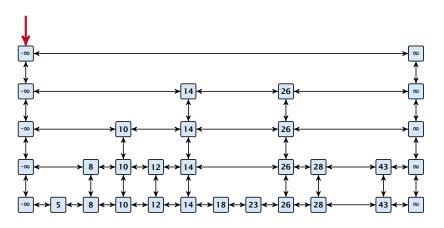
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The time for both operations is dominated by the search time.

Insert (35):

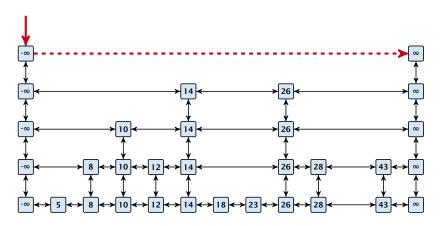


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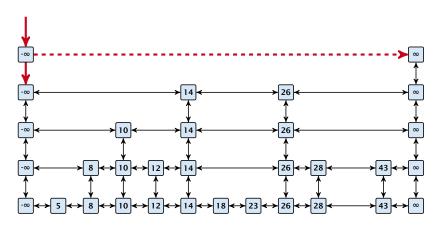




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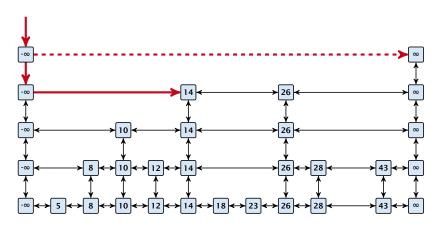


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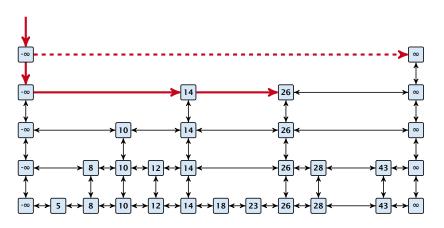


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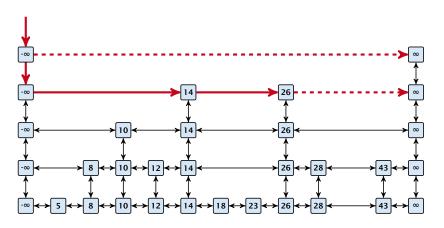
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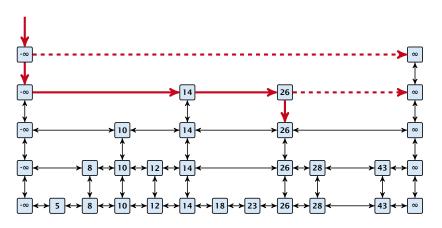
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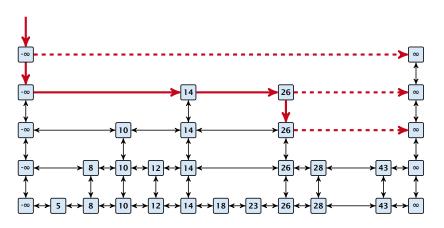
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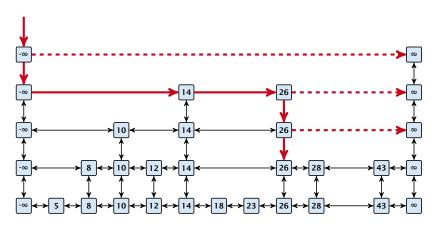
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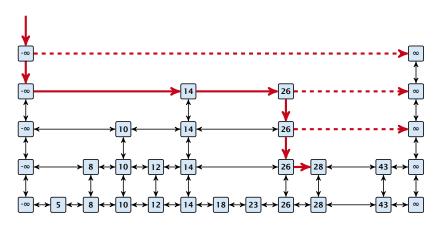


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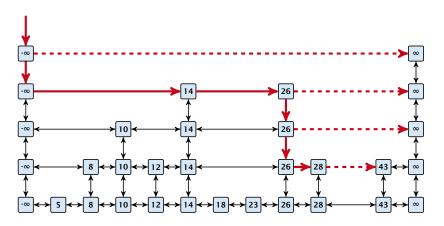
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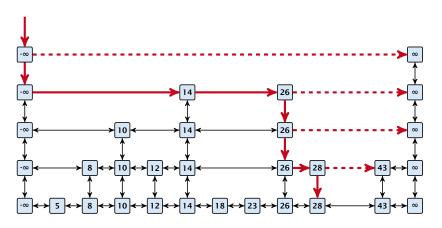
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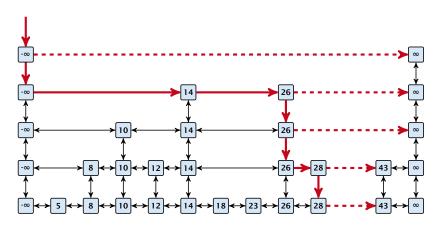




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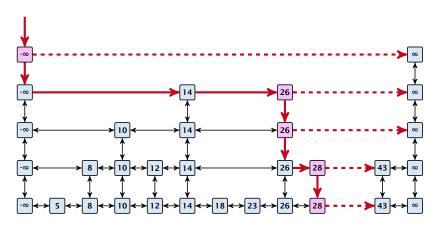


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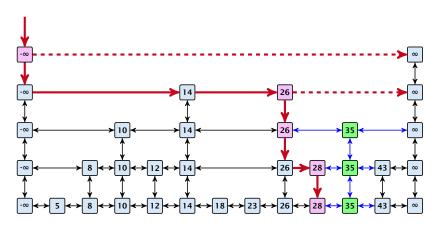


6. Feb. 2022 209/616

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Definition 18 (High Probability)

We say a **randomized** algorithm has running time $\mathcal{O}(\log n)$ with high probability if for any constant α the running time is at most $\mathcal{O}(\log n)$ with probability at least $1 - \frac{1}{n^{\alpha}}$.

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Here the \mathcal{O} -notation hides a constant that may depend on α .



Suppose there are polynomially many events E_1, E_2, \dots, E_ℓ , $\ell = n^c$ each holding with high probability (e.g. E_i may be the event that the i-th search in a skip list takes time at most $\mathcal{O}(\log n)$).

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$$\Pr[E_1 \wedge \cdots \wedge E_{\ell}] = 1 - \Pr[\bar{E}_1 \vee \cdots \vee \bar{E}_{\ell}]$$

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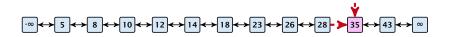
This means $Pr[E_1 \wedge \cdots \wedge E_{\ell}]$ holds with high probability.

Lemma 19

A search (and, hence, also insert and delete) in a skip list with n elements takes time O(logn) with high probability (w. h. p.).

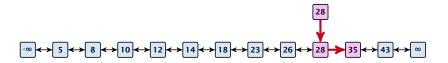
$$\begin{array}{c} -\infty \longleftrightarrow 5 \longleftrightarrow 8 \longleftrightarrow 10 \longleftrightarrow 12 \longleftrightarrow 14 \longleftrightarrow 18 \longleftrightarrow 23 \longleftrightarrow 26 \longleftrightarrow 28 \longleftrightarrow 35 \longleftrightarrow 43 \longleftrightarrow \infty \end{array}$$

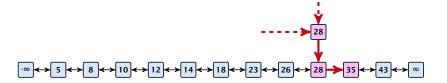
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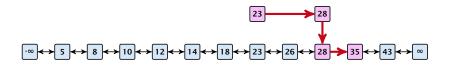


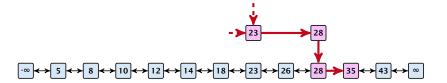
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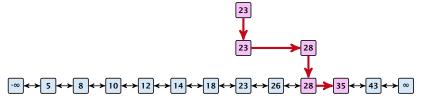


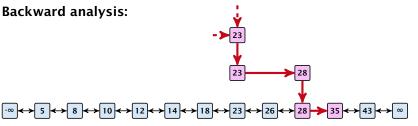




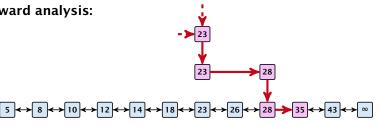






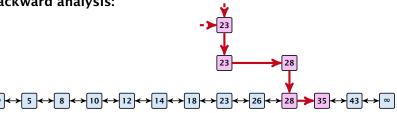


Backward analysis:



At each point the path goes up with probability 1/2 and left with probability 1/2.

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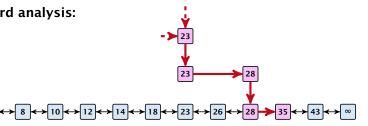


At each point the path goes up with probability 1/2 and left with probability 1/2.

We show that w.h.p:

A "long" search path must also go very high.

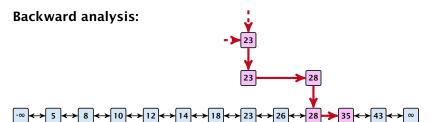
Backward analysis:



At each point the path goes up with probability 1/2 and left with probability 1/2.

We show that w.h.p:

- A "long" search path must also go very high.
- There are no elements in high lists.



At each point the path goes up with probability 1/2 and left with probability 1/2.

We show that w.h.p:

- A "long" search path must also go very high.
- There are no elements in high lists.

From this it follows that w.h.p. there are no long paths.



$$\left(\frac{n}{k}\right)^k \le \binom{n}{k} \le \left(\frac{en}{k}\right)^k$$

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Let $E_{z,k}$ denote the event that a search path is of length z (number of edges) but does not visit a list above L_k .

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In particular, this means that during the construction in the backward analysis we see at most k heads (i.e., coin flips that tell you to go up) in z trials.

 $\Pr[E_{z,k}]$

$$\leq {z \choose k} 2^{-(z-k)}$$

 $\Pr[E_{z,k}] \leq \Pr[\text{at most } k \text{ heads in } z \text{ trials}]$

$$\leq \binom{z}{k} 2^{-(z-k)} \leq \left(\frac{ez}{k}\right)^k 2^{-(z-k)}$$

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choosing $k = y \log n$ with $y \ge 1$ and $z = (\beta + \alpha)y \log n$

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for $\alpha \geq 1$.

So far we fixed $k = y \log n$, $y \ge 1$, and $z = 7\alpha y \log n$, $\alpha \ge 1$.

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Let A_{k+1} denote the event that the list L_{k+1} is non-empty. Then

$$\Pr[A_{k+1}] \le n2^{-(k+1)} \le n^{-(\gamma-1)}$$
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Pr[search requires z steps]

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 $\Pr[\text{search requires } z \text{ steps}] \leq \Pr[E_{z,k}] + \Pr[A_{k+1}]$

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For the search to take at least $z=7\alpha\gamma\log n$ steps either the event $E_{z,k}$ or the event A_{k+1} must hold. Hence,

$$\Pr[\text{search requires } z \text{ steps}] \le \Pr[E_{z,k}] + \Pr[A_{k+1}]$$

 $\le n^{-\alpha} + n^{-(\gamma-1)}$

This means, the search requires at most z steps, w.h.p.

7.6 van Emde Boas Trees

Dynamic Set Data Structure *S***:**

- \triangleright S.insert(x)
- \triangleright S. delete(x)
- \triangleright S. search(x)
- ► *S*. min()
- ► *S*. max()
- \triangleright S. succ(x)
- \triangleright S. pred(x)

7.6 van Emde Boas Trees

For this chapter we ignore the problem of storing satellite data:

- \triangleright S. insert(x): Inserts x into S.
- ▶ S. delete(x): Deletes x from S. Usually assumes that $x \in S$.
- **S.** member(x): Returns 1 if $x \in S$ and 0 otw.
- **S.** min(): Returns the value of the minimum element in S.
- **S.** $\max()$: Returns the value of the maximum element in S.
- ► *S.* succ(*x*): Returns successor of *x* in *S*. Returns null if *x* is maximum or larger than any element in *S*. Note that *x* needs not to be in *S*.
- ▶ S. pred(x): Returns the predecessor of x in S. Returns null if x is minimum or smaller than any element in S. Note that x needs not to be in S.

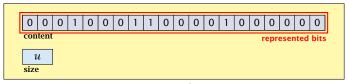


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7.6 van Emde Boas Trees

Can we improve the existing algorithms when the keys are from a restricted set?

In the following we assume that the keys are from $\{0, 1, \dots, u-1\}$, where u denotes the size of the universe.



one array of *u* bits

Use an array that encodes the indicator function of the dynamic set.

```
Algorithm 1 array.insert(x)
```

1: content[x] \leftarrow 1;

Algorithm 2 array.delete(x)

1: content[x] \leftarrow 0;

Algorithm 3 array.member(x)

1: return content[x];

- Note that we assume that x is valid, i.e., it falls within the array boundaries.
- ▶ Obviously(?) the running time is constant.

Algorithm 4 array.max()

1: for $(i = \text{size} -1; i \ge 0; i--)$ do 2: if content[i] = 1 then return i;

3: return null;

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Algorithm 5 array.min()

1: **for** (i = 0; i < size; i++) **do**

2: **if** content[i] = 1 **then return** i;

3: return null;

Algorithm 4 array.max()

```
1: for (i = \text{size} - 1; i \ge 0; i--) do

2: if content[i] = 1 then return i;

3: return null;
```

Algorithm 5 array.min()

```
1: for (i = 0; i < \text{size}; i++) do

2: if content[i] = 1 then return i;

3: return null;
```

▶ Running time is $\mathcal{O}(u)$ in the worst case.

Algorithm 6 array.succ(x)

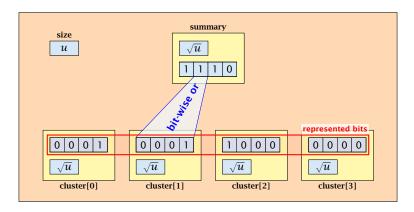
1: for (i = x + 1; i < size; i++) do 2: if content[i] = 1 then return i; 3: return null;

Algorithm 7 array.pred(x)

1: for $(i = x - 1; i \ge 0; i--)$ do 2: if content[i] = 1 then return i;

3: return null:

Running time is $\mathcal{O}(u)$ in the worst case.



- \sqrt{u} cluster-arrays of \sqrt{u} bits.
- One summary-array of \sqrt{u} bits. The *i*-th bit in the summary array stores the bit-wise or of the bits in the *i*-th cluster.

The bit for a key x is contained in cluster number $\left\lfloor \frac{x}{\sqrt{u}} \right\rfloor$.

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Within the cluster-array the bit is at position $x \mod \sqrt{u}$.

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Within the cluster-array the bit is at position $x \mod \sqrt{u}$.

For simplicity we assume that $u=2^{2k}$ for some $k\geq 1$. Then we can compute the cluster-number for an entry x as $\mathrm{high}(x)$ (the upper half of the dual representation of x) and the position of x within its cluster as $\mathrm{low}(x)$ (the lower half of the dual representation).

Algorithm 8 member(x)

1: **return** cluster[high(x)]. member(low(x));

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Algorithm 9 insert(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)].\operatorname{insert}(\operatorname{low}(x));$
- 2: summary.insert(high(x));

Algorithm 8 member(x)

1: **return** cluster[high(x)]. member(low(x));

Algorithm 9 insert(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)].\operatorname{insert}(\operatorname{low}(x));$
- 2: summary.insert(high(x));
- ► The running times are constant, because the corresponding array-functions have constant running times.

Algorithm 10 delete(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)]$. $\operatorname{delete}(\operatorname{low}(x))$;
- 2: **if** cluster[high(x)]. min() = null **then**
- 3: summary . delete(high(x));

Algorithm 10 delete(x)

```
1: \operatorname{cluster}[\operatorname{high}(x)]. \operatorname{delete}(\operatorname{low}(x));
```

2: **if** cluster[high(x)].min() = null **then**

3: summary . delete(high(x));

▶ The running time is dominated by the cost of a minimum computation on an array of size \sqrt{u} . Hence, $O(\sqrt{u})$.

Algorithm 11 max()

- 1: maxcluster ← summary.max(); 2: if maxcluster = null return null; 3: offs ← cluster[maxcluster].max() 4: return maxcluster ∘ offs;

Algorithm 11 max()

- 1: *maxcluster* ← summary.max();
- 2: **if** *maxcluster* = null **return** null;
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Algorithm 12 min()

- 1: *mincluster* ← summary.min();
- 2: **if** *mincluster* = null **return** null;
- 3: $offs \leftarrow cluster[mincluster].min();$
- 4: **return** *mincluster* ∘ *offs*;

Algorithm 11 max()

- 1: *maxcluster* ← summary.max();
- 2: **if** *maxcluster* = null **return** null;
- 3: $offs \leftarrow cluster[maxcluster].max()$
- 4: **return** *maxcluster* \circ *offs*;

Algorithm 12 min()

- 1: *mincluster* ← summary.min();
- 2: **if** *mincluster* = null **return** null;
- 3: $offs \leftarrow cluster[mincluster].min();$
- 4: **return** mincluster offs;

Running time is roughly $2\sqrt{u} = \mathcal{O}(\sqrt{u})$ in the worst case.

The operator \circ stands for the concatenation of two bitstrings.

This means if $x = 0111_2$ and $y = 0001_2$ then $x \circ y = 01110001_2$.

```
Algorithm 13 \operatorname{succ}(x)

1: m \leftarrow \operatorname{cluster}[\operatorname{high}(x)].\operatorname{succ}(\operatorname{low}(x))

2: if m \neq \operatorname{null} then return \operatorname{high}(x) \circ m;

3: \operatorname{succcluster} \leftarrow \operatorname{summary}.\operatorname{succ}(\operatorname{high}(x));

4: if \operatorname{succcluster} \neq \operatorname{null} then

5: \operatorname{offs} \leftarrow \operatorname{cluster}[\operatorname{succcluster}].\operatorname{min}();

6: \operatorname{return} \operatorname{succcluster} \circ \operatorname{offs};

7: \operatorname{return} \operatorname{null};
```

Implementation 2: Summary Array

```
Algorithm 13 \operatorname{succ}(x)

1: m \leftarrow \operatorname{cluster}[\operatorname{high}(x)].\operatorname{succ}(\operatorname{low}(x))

2: if m \neq \operatorname{null} then return \operatorname{high}(x) \circ m;

3: \operatorname{succcluster} \leftarrow \operatorname{summary}.\operatorname{succ}(\operatorname{high}(x));

4: if \operatorname{succcluster} \neq \operatorname{null} then

5: \operatorname{offs} \leftarrow \operatorname{cluster}[\operatorname{succcluster}].\operatorname{min}();

6: \operatorname{return} \operatorname{succcluster} \circ \operatorname{offs};

7: \operatorname{return} \operatorname{null};
```

▶ Running time is roughly $3\sqrt{u} = \mathcal{O}(\sqrt{u})$ in the worst case.

Implementation 2: Summary Array

```
Algorithm 14 \operatorname{pred}(x)

1: m \leftarrow \operatorname{cluster}[\operatorname{high}(x)].\operatorname{pred}(\operatorname{low}(x))

2: if m \neq \operatorname{null} then return \operatorname{high}(x) \circ m;

3: \operatorname{predcluster} \leftarrow \operatorname{summary.\operatorname{pred}}(\operatorname{high}(x));

4: if \operatorname{predcluster} \neq \operatorname{null} then

5: \operatorname{offs} \leftarrow \operatorname{cluster}[\operatorname{predcluster}].\operatorname{max}();

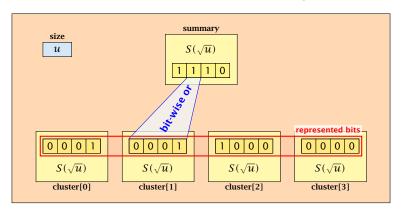
6: \operatorname{return} \operatorname{predcluster} \circ \operatorname{offs};

7: \operatorname{return} \operatorname{null};
```

• Running time is roughly $3\sqrt{u} = \mathcal{O}(\sqrt{u})$ in the worst case.

Instead of using sub-arrays, we build a recursive data-structure.

S(u) is a dynamic set data-structure representing u bits:



We assume that $u = 2^{2^k}$ for some k.

The data-structure S(2) is defined as an array of 2-bits (end of the recursion).

The code from Implementation 2 can be used unchanged. We only need to redo the analysis of the running time.

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Note that in the code we do not need to specifically address the non-recursive case. This is achieved by the fact that an S(4) will contain S(2)'s as sub-datastructures, which are arrays. Hence, a call like cluster[1]. min() from within the data-structure S(4) is not a recursive call as it will call the function array. min().

The code from Implementation 2 can be used unchanged. We only need to redo the analysis of the running time.

Note that in the code we do not need to specifically address the non-recursive case. This is achieved by the fact that an S(4) will contain S(2)'s as sub-datastructures, which are arrays. Hence, a call like cluster[1]. $\min()$ from within the data-structure S(4) is not a recursive call as it will call the function $\operatorname{array.min}()$.

This means that the non-recursive case is been dealt with while initializing the data-structure.

Algorithm 15 member(x)

1: **return** cluster[high(x)].member(low(x));

 $T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1.$

Algorithm 16 insert(x)

- 1: $\operatorname{cluster}[\operatorname{high}(x)]$. $\operatorname{insert}(\operatorname{low}(x))$;
- 2: summary.insert(high(x));
- $T_{ins}(u) = 2T_{ins}(\sqrt{u}) + 1.$

Algorithm 17 delete(x)

- 1: cluster[high(x)]. delete(low(x));
- 2: **if** cluster[high(x)].min() = null **then**
- 3: summary . delete(high(x));
- ► $T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1$.

Algorithm 18 min()

- 1: *mincluster* ← summary.min();
- 2: **if** *mincluster* = null **return** null;
- 3: *offs* ← cluster[*mincluster*].min();
- 4: **return** $mincluster \circ offs$;
- $T_{\min}(u) = 2T_{\min}(\sqrt{u}) + 1.$

```
Algorithm 19 succ(x)

1: m \leftarrow cluster[high(x)].succ(low(x))

2: if m \neq null then return high(x) \circ m;

3: succcluster \leftarrow summary.succ(high(x));

4: if succcluster \neq null then

5: offs \leftarrow cluster[succcluster].min();

6: return succcluster \circ offs;

7: return null;
```

 $T_{\text{succ}}(u) = 2T_{\text{succ}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1.$

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
:

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
:

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{mem}}(2^{\ell})$.

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
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Set $\ell := \log u$ and $X(\ell) := T_{\text{mem}}(2^{\ell})$. Then

$$T_{\mathrm{mem}}(u) = T_{\mathrm{mem}}(\sqrt{u}) + 1$$
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$$X(\ell) = T_{\text{mem}}(2^{\ell})$$

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$$\ell := \log u$$
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 and $X(\ell) := T_{\text{mem}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{mem}}(2^{\ell}) = T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$

= $T_{\text{mem}}(2^{\frac{\ell}{2}}) + 1$

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
:

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{mem}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{mem}}(2^{\ell}) = T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$

= $T_{\text{mem}}(2^{\frac{\ell}{2}}) + 1 = X(\frac{\ell}{2}) + 1$.

$$T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$
:

Set $\ell := \log u$ and $X(\ell) := T_{\text{mem}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{mem}}(2^{\ell}) = T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1$$

= $T_{\text{mem}}(2^{\frac{\ell}{2}}) + 1 = X(\frac{\ell}{2}) + 1$.

Using Master theorem gives $X(\ell) = \mathcal{O}(\log \ell)$, and hence $T_{\text{mem}}(u) = \mathcal{O}(\log \log u)$.

$$T_{\rm ins}(u) = 2T_{\rm ins}(\sqrt{u}) + 1.$$

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Set $\ell := \log u$ and $X(\ell) := T_{\text{ins}}(2^{\ell})$.

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m ins}(u)=2T_{
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 Set $\ell:=\log u$ and $X(\ell):=T_{
m ins}(2^\ell).$ Then $X(\ell)$

$$T_{\rm ins}(u) = 2T_{\rm ins}(\sqrt{u}) + 1.$$

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{ins}}(2^{\ell})$. Then

$$X(\ell) = T_{\rm ins}(2^{\ell})$$

$$T_{\rm ins}(u) = 2T_{\rm ins}(\sqrt{u}) + 1.$$

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{ins}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{ins}}(2^{\ell}) = T_{\text{ins}}(u)$$

$$T_{\rm ins}(u) = 2T_{\rm ins}(\sqrt{u}) + 1.$$

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{ins}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{ins}}(2^{\ell}) = T_{\text{ins}}(u) = 2T_{\text{ins}}(\sqrt{u}) + 1$$

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Set
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$$X(\ell) = T_{\text{ins}}(2^{\ell}) = T_{\text{ins}}(u) = 2T_{\text{ins}}(\sqrt{u}) + 1$$

= $2T_{\text{ins}}(2^{\frac{\ell}{2}}) + 1 = 2X(\frac{\ell}{2}) + 1$.

$$T_{\rm ins}(u) = 2T_{\rm ins}(\sqrt{u}) + 1.$$

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Using Master theorem gives $X(\ell) = \mathcal{O}(\ell)$, and hence $T_{\text{ins}}(u) = \mathcal{O}(\log u)$.

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Using Master theorem gives $X(\ell) = \mathcal{O}(\ell)$, and hence $T_{\rm ins}(u) = \mathcal{O}(\log u)$.

The same holds for $T_{\text{max}}(u)$ and $T_{\text{min}}(u)$.

$$T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1 \le 2T_{\text{del}}(\sqrt{u}) + \frac{c}{\log(u)}.$$

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{c}\log(u).$$

Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$.

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$T_{\rm del}(u)=2T_{\rm del}(\sqrt{u})+T_{\rm min}(\sqrt{u})+1\leq 2T_{\rm del}(\sqrt{u})+c\log(u).$$
 Set $\ell:=\log u$ and $X(\ell):=T_{\rm del}(2^\ell).$ Then
$$X(\ell)$$

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{del}}(2^{\ell})$$

$$T_{\rm del}(u) = 2T_{\rm del}(\sqrt{u}) + T_{\rm min}(\sqrt{u}) + 1 \le 2T_{\rm del}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set
$$\ell := \log u$$
 and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{del}}(2^{\ell}) = T_{\text{del}}(u)$$

$$T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1 \le 2T_{\text{del}}(\sqrt{u}) + \frac{c}{\log(u)}.$$

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 and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

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Set
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$$= 2T_{\text{del}}(2^{\frac{\ell}{2}}) + c\ell$$

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$$= 2T_{\text{del}}(2^{\frac{\ell}{2}}) + c\ell = 2X(\frac{\ell}{2}) + c\ell .$$

$$T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1 \le 2T_{\text{del}}(\sqrt{u}) + c \log(u).$$

Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{del}}(2^{\ell}) = T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + c \log u$$
$$= 2T_{\text{del}}(2^{\frac{\ell}{2}}) + c\ell = 2X(\frac{\ell}{2}) + c\ell .$$

Using Master theorem gives $X(\ell) = \Theta(\ell \log \ell)$, and hence $T_{\text{del}}(u) = \mathcal{O}(\log u \log \log u)$.

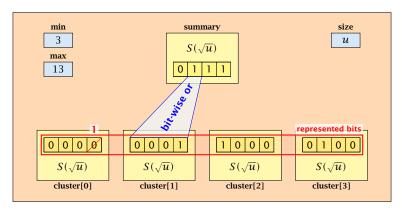
$$T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + T_{\min}(\sqrt{u}) + 1 \le 2T_{\text{del}}(\sqrt{u}) + \frac{c}{\log(u)}.$$

Set $\ell := \log u$ and $X(\ell) := T_{\text{del}}(2^{\ell})$. Then

$$X(\ell) = T_{\text{del}}(2^{\ell}) = T_{\text{del}}(u) = 2T_{\text{del}}(\sqrt{u}) + c \log u$$
$$= 2T_{\text{del}}(2^{\frac{\ell}{2}}) + c\ell = 2X(\frac{\ell}{2}) + c\ell .$$

Using Master theorem gives $X(\ell) = \Theta(\ell \log \ell)$, and hence $T_{\text{del}}(u) = \mathcal{O}(\log u \log \log u)$.

The same holds for $T_{\text{pred}}(u)$ and $T_{\text{succ}}(u)$.



- The bit referenced by min is not set within sub-datastructures.
- The bit referenced by max is set within sub-datastructures (if max ≠ min).

Advantages of having max/min pointers:

Recursive calls for min and max are constant time.

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- ightharpoonup min = null means that the data-structure is empty.
- min = max ≠ null means that the data-structure contains exactly one element.
- We can insert into an empty datastructure in constant time by only setting min = max = x.
- We can delete from a data-structure that just contains one element in constant time by setting min = max = null.

Algorithm 20 max()

1: **return** max;

Algorithm 21 min()

1: return min;

Constant time.

Algorithm 22 member(x)

- 1: if $x = \min$ then return 1; // TRUE
- 2: **return** cluster[high(x)]. member(low(x));
- $T_{\text{mem}}(u) = T_{\text{mem}}(\sqrt{u}) + 1 \Longrightarrow T(u) = \mathcal{O}(\log \log u).$

```
Algorithm 23 succ(x)
 1: if min \neq null \wedge x < min then return min:
 2: maxincluster \leftarrow cluster[high(x)].max();
 3: if maxincluster \neq null \land low(x) < maxincluster then
         offs \leftarrow cluster[high(x)]. succ(low(x));
4:
         return high(x) \circ offs;
 5:
6: else
         succeluster \leftarrow summary.succ(high(x));
7:
8:
         if succeluster = null then return null:
9:
         offs \leftarrow cluster[succeluster].min();
         return succeluster o offs:
10:
```

 $T_{\text{succ}}(u) = T_{\text{succ}}(\sqrt{u}) + 1 \Longrightarrow T_{\text{succ}}(u) = \mathcal{O}(\log \log u).$

```
Algorithm 35 insert(x)
 1: if min = null then
       \min = x; \max = x;
3: else
4:
       if x < \min then exchange x and \min;
     if x > \max then \max = x;
6:
     if cluster[high(x)]. min = null; then
7:
            summary insert(high(x));
8:
            cluster[high(x)].insert(low(x));
        else
9:
            cluster[high(x)].insert(low(x));
10:
```

 $T_{\text{ins}}(u) = T_{\text{ins}}(\sqrt{u}) + 1 \Longrightarrow T_{\text{ins}}(u) = \mathcal{O}(\log \log u).$

Note that the recusive call in Line 8 takes constant time as the if-condition in Line 6 ensures that we are inserting in an empty sub-tree.

The only non-constant recursive calls are the call in Line 7 and in Line 10. These are mutually exclusive, i.e., only one of these calls will actually occur.

From this we get that $T_{\text{ins}}(u) = T_{\text{ins}}(\sqrt{u}) + 1$.

Assumes that x is contained in the structure.

```
Algorithm 36 delete(x)
1: if min = max then
       min = max = null;
 3: else
4:
       if x = \min then
             firstcluster ← summary.min();
6:
             offs \leftarrow cluster[firstcluster].min();
       x \leftarrow firstcluster \circ offs;
 7:
         \min \leftarrow x;
        cluster[high(x)]. delete(low(x));
 9:
                         continued...
```

Assumes that x is contained in the structure.

```
Algorithm 36 delete(x)
 1: if min = max then
        min = max = null;
 3: else
4:
         if x = \min then
                                              find new minimum
              firstcluster \leftarrow summary.min();
 5:
              offs ← cluster[firstcluster]. min();
6:
              x \leftarrow firstcluster \circ offs;
 7:
 8:
          \min \leftarrow x:
         cluster[high(x)]. delete(low(x));
 9:
                          continued...
```

Assumes that x is contained in the structure.

```
Algorithm 36 delete(x)
 1: if min = max then
        min = max = null;
 3: else
4:
        if x = \min then
              firstcluster \leftarrow summary.min();
 5:
6:
               offs \leftarrow cluster[firstcluster].min();
              x \leftarrow firstcluster \circ offs;
 7:
 8:
              \min \leftarrow x:
         cluster[high(x)]. delete(low(x));
 9:
                                                           delete
                           continued...
```

```
Algorithm 36 delete(x)
                            ...continued
         if cluster[high(x)]. min() = null then
10:
              summary . delete(high(x));
11:
              if x = \max then
12:
13:
                   summax \leftarrow summary.max();
14:
                   if summax = null then max \leftarrow min;
                   else
15:
16:
                        offs \leftarrow cluster[summax]. max();
17:
                        \max \leftarrow summax \circ offs
         else
18:
              if x = \max then
19:
20:
                   offs \leftarrow cluster[high(x)]. max();
                   \max \leftarrow \text{high}(x) \circ \text{offs};
21:
```

```
Algorithm 36 delete(x)
                            ...continued
                                                      fix maximum
         if cluster[high(x)]. min() = null then
10:
              summary . delete(high(x));
11:
              if x = \max then
12:
13:
                   summax \leftarrow summary.max();
                   if summax = null then max \leftarrow min;
14:
                   else
15:
16:
                        offs \leftarrow cluster[summax]. max();
17:
                        \max \leftarrow summax \circ offs
         else
18:
              if x = \max then
19:
20:
                   offs \leftarrow cluster[high(x)]. max();
                   \max \leftarrow \text{high}(x) \circ \text{offs};
21:
```

Note that only one of the possible recusive calls in Line 9 and Line 11 in the deletion-algorithm may take non-constant time.

To see this observe that the call in Line 11 only occurs if the cluster where x was deleted is now empty. But this means that the call in Line 9 deleted the last element in cluster[high(x)]. Such a call only takes constant time.

Hence, we get a recurrence of the form

$$T_{\rm del}(u) = T_{\rm del}(\sqrt{u}) + c$$
.

This gives $T_{del}(u) = \mathcal{O}(\log \log u)$.

7.6 van Emde Boas Trees

Space requirements:

The space requirement fulfills the recurrence

$$S(u) = (\sqrt{u} + 1)S(\sqrt{u}) + \mathcal{O}(\sqrt{u}) .$$

- Note that we cannot solve this recurrence by the Master theorem as the branching factor is not constant.
- One can show by induction that the space requirement is $S(u) = \mathcal{O}(u)$. Exercise.

Let the "real" recurrence relation be

$$S(k^2) = (k+1)S(k) + c_1 \cdot k; S(4) = c_2$$

▶ Replacing S(k) by $R(k) := S(k)/c_2$ gives the recurrence

$$R(k^2) = (k+1)R(k) + ck; R(4) = 1$$

where $c = c_1/c_2 < 1$.

- Now, we show $R(k) \le k 2$ for squares $k \ge 4$.
 - Obviously, this holds for k = 4.
 - For $k = \ell^2 > 4$ with ℓ integral we have

$$R(k) = (1 + \ell)R(\ell) + c\ell$$

$$\leq (1 + \ell)(\ell - 2) + \ell \leq k - 2$$

▶ This shows that R(k) and, hence, S(k) grows linearly.

Dictionary:

- **S.** insert(x): Insert an element x.
- S. delete(x): Delete the element pointed to by x.
- ▶ *S.* search(k): Return a pointer to an element e with key[e] = k in S if it exists; otherwise return null.

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So far we have implemented the search for a key by carefully choosing split-elements.

Dictionary:

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So far we have implemented the search for a key by carefully choosing split-elements.

Then the memory location of an object x with key k is determined by successively comparing k to split-elements.

Dictionary:

- **S.** insert(x): Insert an element x.
- S. delete(x): Delete the element pointed to by x.
- ▶ *S.* search(k): Return a pointer to an element e with key[e] = k in S if it exists; otherwise return null.

So far we have implemented the search for a key by carefully choosing split-elements.

Then the memory location of an object x with key k is determined by successively comparing k to split-elements.

Hashing tries to directly compute the memory location from the given key. The goal is to have constant search time.

Definitions:

▶ Universe U of keys, e.g., $U \subseteq \mathbb{N}_0$. U very large.

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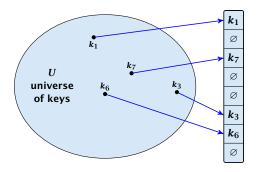
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The hash-function h should fulfill:

- Fast to evaluate.
- Small storage requirement.
- Good distribution of elements over the whole table.

Direct Addressing

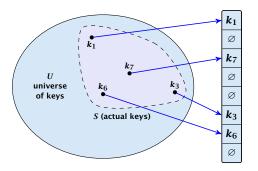
Ideally the hash function maps all keys to different memory locations.



This special case is known as Direct Addressing. It is usually very unrealistic as the universe of keys typically is quite large, and in particular larger than the available memory.

Perfect Hashing

Suppose that we know the set S of actual keys (no insert/no delete). Then we may want to design a simple hash-function that maps all these keys to different memory locations.



Such a hash function h is called a perfect hash function for set S.

If we do not know the keys in advance, the best we can hope for is that the hash function distributes keys evenly across the table.

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Hence, there may be two elements k_1, k_2 from the set S that map to the same memory location (i.e., $h(k_1) = h(k_2)$). This is called a collision.

Typically, collisions do not appear once the size of the set S of actual keys gets close to n, but already when $|S| \ge \omega(\sqrt{n})$.

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Lemma 20

The probability of having a collision when hashing m elements into a table of size n under uniform hashing is at least

$$1 - e^{-\frac{m(m-1)}{2n}} \approx 1 - e^{-\frac{m^2}{2n}}$$
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Uniform hashing:

Choose a hash function uniformly at random from all functions $f: U \to [0, \dots, n-1]$.



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Proof.

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Let $A_{m,n}$ denote the event that inserting m keys into a table of size n does not generate a collision. Then

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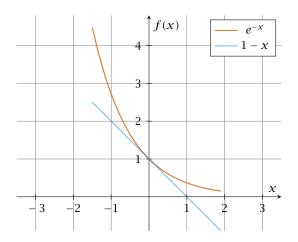
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Here the first equality follows since the ℓ -th element that is hashed has a probability of $\frac{n-\ell+1}{n}$ to not generate a collision under the condition that the previous elements did not induce collisions.

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The inequality $1-x \le e^{-x}$ is derived by stopping the Taylor-expansion of e^{-x} after the second term.



Resolving Collisions

The methods for dealing with collisions can be classified into the two main types

- open addressing, aka. closed hashing
- hashing with chaining, aka. closed addressing, open hashing.

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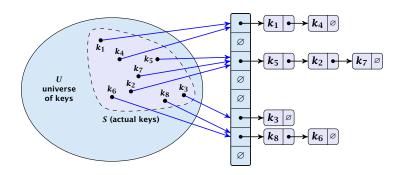
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- open addressing, aka. closed hashing
- hashing with chaining, aka. closed addressing, open hashing.

There are applications e.g. computer chess where you do not resolve collisions at all.

Arrange elements that map to the same position in a linear list.

- Access: compute h(x) and search list for key[x].
- Insert: insert at the front of the list.



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$$A^- = 1 + \alpha .$$

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$$= 1+\frac{m-1}{2n}=1+\frac{\alpha}{2}-\frac{\alpha}{2m}.$$

$$\begin{split} \mathbf{E} \left[\frac{1}{m} \sum_{i=1}^{m} \left(1 + \sum_{j=i+1}^{m} X_{ij} \right) \right] &= \frac{1}{m} \sum_{i=1}^{m} \left(1 + \sum_{j=i+1}^{m} \mathbf{E} \left[X_{ij} \right] \right) \\ &= \frac{1}{m} \sum_{i=1}^{m} \left(1 + \sum_{j=i+1}^{m} \frac{1}{n} \right) \\ &= 1 + \frac{1}{mn} \sum_{i=1}^{m} (m-i) \\ &= 1 + \frac{1}{mn} \left(m^2 - \frac{m(m+1)}{2} \right) \\ &= 1 + \frac{m-1}{2n} = 1 + \frac{\alpha}{2} - \frac{\alpha}{2m} \end{split} .$$

Hence, the expected cost for a successful search is $A^+ \leq 1 + \frac{\alpha}{2}$.

Disadvantages:

- pointers increase memory requirements
- pointers may lead to bad cache efficiency

Advantages:

- no à priori limit on the number of elements
- deletion can be implemented efficiently
- by using balanced trees instead of linked list one can also obtain worst-case guarantees.

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Define a function h(k, j) that determines the table-position to be examined in the j-th step. The values $h(k,0),\ldots,h(k,n-1)$ must form a permutation of $0, \ldots, n-1$.

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Insert(x): Search until you find an empty slot; insert your element there. If your search reaches h(k, n-1), and this slot is non-empty then your table is full.

Choices for h(k, j):

Linear probing:

```
h(k,i) = h(k) + i \mod n
(sometimes: h(k,i) = h(k) + ci \mod n).
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Choices for h(k, j):

Linear probing: $h(k, i) = h(k) + i \mod n$ (sometimes: $h(k, i) = h(k) + ci \mod n$).

- Quadratic probing: $h(k, i) = h(k) + c_1 i + c_2 i^2 \mod n$.
- Double hashing: $h(k, i) = h_1(k) + ih_2(k) \mod n$.

For quadratic probing and double hashing one has to ensure that the search covers all positions in the table (i.e., for double hashing $h_2(k)$ must be relatively prime to n (teilerfremd); for quadratic probing c_1 and c_2 have to be chosen carefully).

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Linear Probing

Advantage: Cache-efficiency. The new probe position is very likely to be in the cache.

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Lemma 21

Let L be the method of linear probing for resolving collisions:

$$L^+ \approx \frac{1}{2} \left(1 + \frac{1}{1 - \alpha} \right)$$

$$L^- \approx \frac{1}{2} \left(1 + \frac{1}{(1 - \alpha)^2} \right)$$



Quadratic Probing

- Not as cache-efficient as Linear Probing.
- Secondary clustering: caused by the fact that all keys mapped to the same position have the same probe sequence.

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Lemma 22

Let Q be the method of quadratic probing for resolving collisions:

$$Q^+ \approx 1 + \ln\left(\frac{1}{1-\alpha}\right) - \frac{\alpha}{2}$$

$$Q^- \approx \frac{1}{1-\alpha} + \ln\left(\frac{1}{1-\alpha}\right) - \alpha$$

Double Hashing

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Lemma 23

Let D be the method of double hashing for resolving collisions:

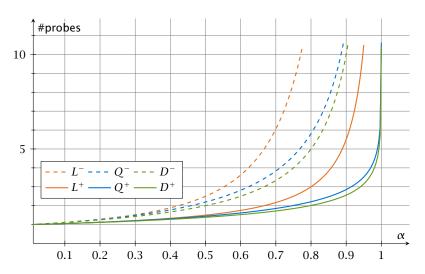
$$D^+ \approx \frac{1}{\alpha} \ln \left(\frac{1}{1 - \alpha} \right)$$

$$D^- \approx \frac{1}{1-\alpha}$$

Some values:

α	Linear Probing		Quadratic Probing		Double Hashing	
	L^+	L^{-}	Q^+	Q^-	D^+	D^-
0.5	1.5	2.5	1.44	2.19	1.39	2
0.9	5.5	50.5	2.85	11.40	2.55	10
0.95	10.5	200.5	3.52	22.05	3.15	20

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We analyze the time for a search in a very idealized Open Addressing scheme.

► The probe sequence h(k,0), h(k,1), h(k,2),... is equally likely to be any permutation of (0,1,...,n-1).

Let X denote a random variable describing the number of probes in an unsuccessful search.

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$$Pr[A_1 \cap A_2 \cap \cdots \cap A_{i-1}]$$

$$= Pr[A_1] \cdot Pr[A_2 \mid A_1] \cdot Pr[A_3 \mid A_1 \cap A_2] \cdot \cdots \cdot Pr[A_{i-1} \mid A_1 \cap \cdots \cap A_{i-2}]$$

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$$\le \left(\frac{m}{n}\right)^{i-1} = \alpha^{i-1} .$$

E[X]

$$E[X] = \sum_{i=1}^{\infty} \Pr[X \ge i]$$

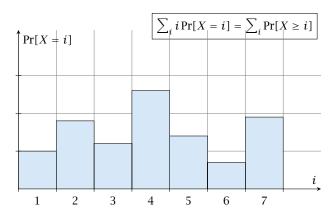
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$$E[X] = \sum_{i=1}^{\infty} \Pr[X \ge i] \le \sum_{i=1}^{\infty} \alpha^{i-1} = \sum_{i=0}^{\infty} \alpha^{i}$$

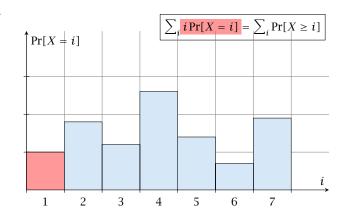
$$E[X] = \sum_{i=1}^{\infty} \Pr[X \ge i] \le \sum_{i=1}^{\infty} \alpha^{i-1} = \sum_{i=0}^{\infty} \alpha^{i} = \frac{1}{1-\alpha}.$$

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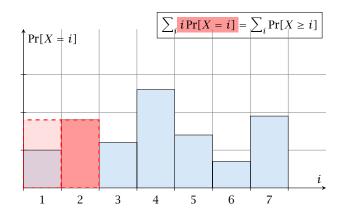
$$\frac{1}{1-\alpha}=1+\alpha+\alpha^2+\alpha^3+\dots$$



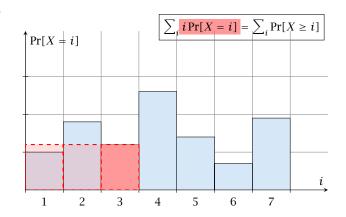
i = 1



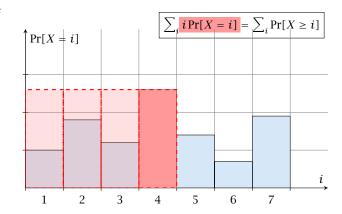
$$i = 2$$



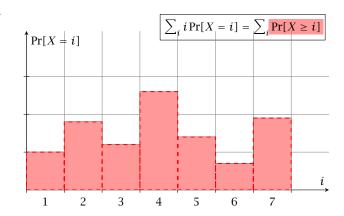
$$i = 3$$



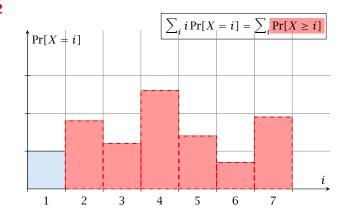
$$i = 4$$



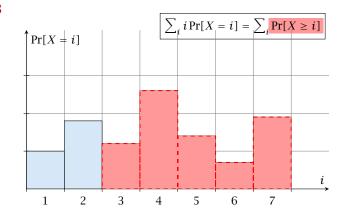




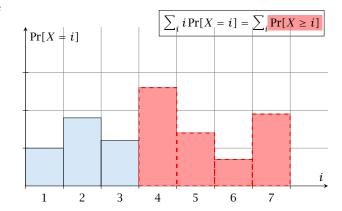
$$i = 2$$

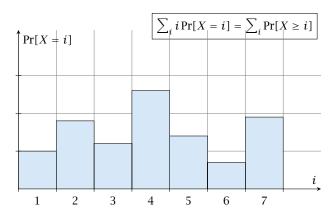


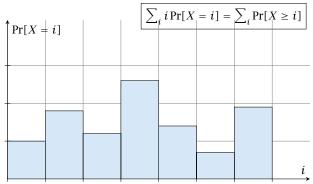
$$i = 3$$



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The j-th rectangle appears in both sums j^6 times. (j times in the first due to multiplication with j; and j times in the second for summands $i = 1, 2, \dots, j$

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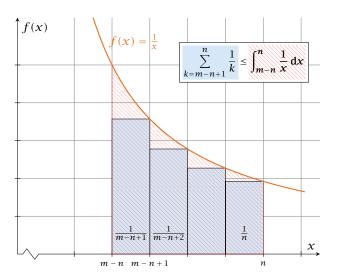
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$$\leq \frac{1}{\alpha} \int_{n-m}^{n} \frac{1}{x} dx = \frac{1}{\alpha} \ln \frac{n}{n-m} = \frac{1}{\alpha} \ln \frac{1}{1-\alpha} .$$



How do we delete in a hash-table?

For hashing with chaining this is not a problem. Simply search for the key, and delete the item in the corresponding list.

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- For hashing with chaining this is not a problem. Simply search for the key, and delete the item in the corresponding list.
- For open addressing this is difficult.

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Deletions in Hashtables

- Simply removing a key might interrupt the probe sequence of other keys which then cannot be found anymore.
- One can delete an element by replacing it with a deleted-marker.
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 - During a search a deleted-marker must not be used to terminate the probe sequence.

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Deletions in Hashtables

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 - During a search a deleted-marker must not be used to terminate the probe sequence.
- The table could fill up with deleted-markers leading to bad performance.
- ▶ If a table contains many deleted-markers (linear fraction of the keys) one can rehash the whole table and amortize the cost for this rehash against the cost for the deletions.



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For Linear Probing one can delete elements without using deletion-markers.

- For Linear Probing one can delete elements without using deletion-markers.
- Upon a deletion elements that are further down in the probe-sequence may be moved to guarantee that they are still found during a search.

Algorithm 37 delete(p) 1: $T[p] \leftarrow \text{null}$ 2: $p \leftarrow \text{succ}(p)$ 3: while $T[p] \neq \text{null do}$ 4: $y \leftarrow T[p]$ 5: $T[p] \leftarrow \text{null}$ 6: $p \leftarrow \text{succ}(p)$ 7: insert(y)

p is the index into the table-cell that contains the object to be deleted.

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Pointers into the hash-table become invalid.



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However, the assumption of uniform hashing that h is chosen randomly from all functions $f:U\to [0,\dots,n-1]$ is clearly unrealistic as there are $n^{|U|}$ such functions. Even writing down such a function would take $|U|\log n$ bits.

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Universal hashing tries to define a set $\mathcal H$ of functions that is much smaller but still leads to good average case behaviour when selecting a hash-function uniformly at random from $\mathcal H$.

Definition 24

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\dots,n-1\}$ is called universal if for all $u_1,u_2\in U$ with $u_1\neq u_2$

$$\Pr[h(u_1) = h(u_2)] \le \frac{1}{n}$$
,

where the probability is w.r.t. the choice of a random hash-function from set \mathcal{H} .

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Note that this means that the probability of a collision between two arbitrary elements is at most $\frac{1}{n}$.

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Definition 25

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\dots,n-1\}$ is called 2-independent (pairwise independent) if the following two conditions hold

- For any key $u \in U$, and $t \in \{0, ..., n-1\}$ $\Pr[h(u) = t] = \frac{1}{n}$, i.e., a key is distributed uniformly within the hash-table.
- For all $u_1, u_2 \in U$ with $u_1 \neq u_2$, and for any two hash-positions t_1, t_2 :

$$\Pr[h(u_1) = t_1 \land h(u_2) = t_2] \le \frac{1}{n^2}$$
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.

This requirement clearly implies a universal hash-function.



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Definition 26

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\dots,n-1\}$ is called k-independent if for any choice of $\ell \le k$ distinct keys $u_1,\dots,u_\ell \in U$, and for any set of ℓ not necessarily distinct hash-positions t_1,\dots,t_ℓ :

$$\Pr[h(u_1) = t_1 \wedge \cdots \wedge h(u_\ell) = t_\ell] \le \frac{1}{n^\ell} ,$$

where the probability is w.r.t. the choice of a random hash-function from set \mathcal{H} .



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Definition 27

A class $\mathcal H$ of hash-functions from the universe U into the set $\{0,\ldots,n-1\}$ is called (μ,k) -independent if for any choice of $\ell \leq k$ distinct keys $u_1,\ldots,u_\ell \in U$, and for any set of ℓ not necessarily distinct hash-positions t_1,\ldots,t_ℓ :

$$\Pr[h(u_1) = t_1 \wedge \cdots \wedge h(u_\ell) = t_\ell] \le \frac{\mu}{n^\ell} ,$$

where the probability is w.r.t. the choice of a random hash-function from set \mathcal{H} .



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Let $U := \{0, \dots, p-1\}$ for a prime p. Let $\mathbb{Z}_p := \{0, \dots, p-1\}$, and let $\mathbb{Z}_p^* := \{1, \dots, p-1\}$ denote the set of invertible elements in \mathbb{Z}_p .

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Define

$$h_{a,b}(x) := (ax + b \bmod p) \bmod n$$

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Define

$$h_{a,b}(x) := (ax + b \bmod p) \bmod n$$

Lemma 28

The class

$$\mathcal{H} = \{ h_{a,b} \mid a \in \mathbb{Z}_p^*, b \in \mathbb{Z}_p \}$$

is a universal class of hash-functions from U to $\{0, \ldots, n-1\}$.

Proof.

Let $x, y \in U$ be two distinct keys. We have to show that the probability of a collision is only 1/n.

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where we use that \mathbb{Z}_p is a field (Körper) and, hence, has no zero divisors (nullteilerfrei).

The hash-function does not generate collisions before the \pmod{n} -operation. Furthermore, every choice (a,b) is mapped to a different pair (t_x,t_y) with $t_x:=ax+b$ and $t_y:=ay+b$.

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This holds because we can compute a and b when given t_x and t_y :

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$$t_{x} \equiv ax + b \qquad (\text{mod } p)$$

$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

$$t_{x} - t_{y} \equiv a(x - y) \qquad (\text{mod } p)$$

$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

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$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

$$t_{x} - t_{y} \equiv a(x - y) \qquad (\text{mod } p)$$

$$t_{y} \equiv ay + b \qquad (\text{mod } p)$$

$$a \equiv (t_{x} - t_{y})(x - y)^{-1} \qquad (\text{mod } p)$$

$$b \equiv t_{y} - ay \qquad (\text{mod } p)$$

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What happens when we do the mod n operation?

Fix a value t_x . There are p-1 possible values for choosing t_y .

From the range $0, \ldots, p-1$ the values $t_x, t_x + n, t_x + 2n, \ldots$ map to t_x after the modulo-operation. These are at most $\lceil p/n \rceil$ values.

As $t_y \neq t_x$ there are

$$\left\lceil \frac{p}{n} \right\rceil - 1$$

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possibilities for choosing $t_{\mathcal{Y}}$ such that the final hash-value creates a collision.

This happens with probability at most $\frac{1}{n}$.

It is also possible to show that $\mathcal H$ is an (almost) pairwise independent class of hash-functions.

$$\Pr_{t_{\mathcal{X}} \neq t_{\mathcal{Y}} \in \mathbb{Z}_p^2} \left[\begin{array}{c} t_{\mathcal{X}} \bmod n = h_1 \\ t_{\mathcal{Y}} \bmod n = h_2 \end{array} \right]$$

7.7 Hashing

It is also possible to show that $\mathcal H$ is an (almost) pairwise independent class of hash-functions.

$$\frac{\left\lfloor \frac{p}{n} \right\rfloor^2}{p(p-1)} \le \Pr_{t_X \neq t_Y \in \mathbb{Z}_p^2} \left[\begin{array}{c} t_X \bmod n = h_1 \\ t_Y \bmod n = h_2 \end{array} \right] \le \frac{\left\lceil \frac{p}{n} \right\rceil^2}{p(p-1)}$$

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$$\frac{\left\lfloor \frac{p}{n} \right\rfloor^2}{p(p-1)} \le \Pr_{t_x \neq t_y \in \mathbb{Z}_p^2} \left[\begin{array}{c} t_x \bmod n = h_1 \\ t_y \bmod n = h_2 \end{array} \right] \le \frac{\left\lceil \frac{p}{n} \right\rceil^2}{p(p-1)}$$

Note that the middle is the probability that $h(x) = h_1$ and $h(y) = h_2$. The total number of choices for (t_x, t_y) is p(p-1). The number of choices for t_x (t_y) such that $t_x \mod n = h_1$ $(t_y \mod n = h_2)$ lies between $\lfloor \frac{p}{n} \rfloor$ and $\lceil \frac{p}{n} \rceil$.

Definition 29

Let $d \in \mathbb{N}$; $q \ge (d+1)n$ be a prime; and let $\bar{a} \in \{0, \dots, q-1\}^{d+1}$. Define for $x \in \{0, \dots, q-1\}$

$$h_{\bar{a}}(x) := \left(\sum_{i=0}^d a_i x^i \bmod q\right) \bmod n$$
.

Let $\mathcal{H}_n^d := \{h_{\bar{a}} \mid \bar{a} \in \{0, \dots, q-1\}^{d+1}\}$. The class \mathcal{H}_n^d is (e, d+1)-independent.

Note that in the previous case we had d = 1 and chose $a_d \neq 0$.

For the coefficients $\bar{a} \in \{0,\ldots,q-1\}^{d+1}$ let $f_{\bar{a}}$ denote the polynomial

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The polynomial is defined by d+1 distinct points.

Fix $\ell \leq d+1$; let $x_1, \ldots, x_\ell \in \{0, \ldots, q-1\}$ be keys, and let t_1, \ldots, t_ℓ denote the corresponding hash-function values.

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$$A^{\ell} = \{h_{\bar{a}} \in \mathcal{H} \mid h_{\bar{a}}(x_i) = t_i \text{ for all } i \in \{1, \dots, \ell\}\}$$

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$$A^{\ell}=\{h_{\bar{a}}\in\mathcal{H}\mid h_{\bar{a}}(x_i)=t_i \text{ for all } i\in\{1,\ldots,\ell\}\}$$

Then

$$h_{\bar{a}} \in A^{\ell} \Leftrightarrow h_{\bar{a}} = f_{\bar{a}} \bmod n$$
 and

$$f_{\bar{a}}(x_i) \in \underbrace{\{t_i + \alpha \cdot n \mid \alpha \in \{0, \dots, \lceil \frac{q}{n} \rceil - 1\}\}}_{=:B_i}$$

Fix $\ell \leq d+1$; let $x_1,\ldots,x_\ell \in \{0,\ldots,q-1\}$ be keys, and let t_1,\ldots,t_ℓ denote the corresponding hash-function values.

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Then
$$h_{\bar{a}} \in A^{\ell} \Leftrightarrow h_{\bar{a}} = f_{\bar{a}} \bmod n \text{ and}$$

$$f_{\bar{a}}(x_i) \in \underbrace{\{t_i + \alpha \cdot n \mid \alpha \in \{0, \dots, \lceil \frac{q}{n} \rceil - 1\} \}}_{=:B_i}$$

In order to obtain the cardinality of A^{ℓ} we choose our polynomial by fixing d+1 points.

Fix $\ell \leq d+1$; let $x_1, \ldots, x_\ell \in \{0, \ldots, q-1\}$ be keys, and let t_1, \ldots, t_ℓ denote the corresponding hash-function values.

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Then

 $h_{\bar{a}} \in A^{\ell} \Leftrightarrow h_{\bar{a}} = f_{\bar{a}} \bmod n$ and

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In order to obtain the cardinality of A^ℓ we choose our polynomial by fixing d+1 points.

We first fix the values for inputs x_1, \ldots, x_ℓ . We have

$$|B_1| \cdot \ldots \cdot |B_{\ell}|$$

possibilities to do this (so that $h_{\bar{a}}(x_i) = t_i$).

Now, we choose $d-\ell+1$ other inputs and choose their value arbitrarily. We have $q^{d-\ell+1}$ possibilities to do this.

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Therefore we have

$$|B_1| \cdot \ldots \cdot |B_{\ell}| \cdot q^{d-\ell+1} \le \lceil \frac{q}{n} \rceil^{\ell} \cdot q^{d-\ell+1}$$

possibilities to choose \bar{a} such that $h_{\bar{a}} \in A_{\ell}$.

$$\frac{\lceil \frac{q}{n} \rceil^{\ell} \cdot q^{d-\ell+1}}{q^{d+1}}$$

$$\frac{\lceil \frac{q}{n} \rceil^{\ell} \cdot q^{d-\ell+1}}{q^{d+1}} \leq \frac{(\frac{q+n}{n})^{\ell}}{q^{\ell}}$$

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$$\begin{split} & \frac{\lceil \frac{q}{n} \rceil^{\ell} \cdot q^{d-\ell+1}}{q^{d+1}} \leq \frac{(\frac{q+n}{n})^{\ell}}{q^{\ell}} \leq \left(\frac{q+n}{q}\right)^{\ell} \cdot \frac{1}{n^{\ell}} \\ & \leq \left(1 + \frac{1}{\ell}\right)^{\ell} \cdot \frac{1}{n^{\ell}} \end{split}$$

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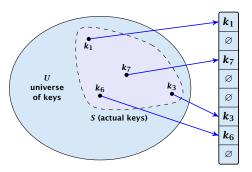
Therefore the probability of choosing $h_{ ilde{a}}$ from A_{ℓ} is only

$$\begin{split} & \frac{\lceil \frac{q}{n} \rceil^{\ell} \cdot q^{d-\ell+1}}{q^{d+1}} \leq \frac{(\frac{q+n}{n})^{\ell}}{q^{\ell}} \leq \left(\frac{q+n}{q}\right)^{\ell} \cdot \frac{1}{n^{\ell}} \\ & \leq \left(1 + \frac{1}{\ell}\right)^{\ell} \cdot \frac{1}{n^{\ell}} \leq \frac{e}{n^{\ell}} \ . \end{split}$$

This shows that the \mathcal{H} is (e, d+1)-universal.

The last step followed from $q \ge (d+1)n$, and $\ell \le d+1$.

Suppose that we **know** the set S of actual keys (no insert/no delete). Then we may want to design a **simple** hash-function that maps all these keys to different memory locations.



6. Feb. 2022

Let m = |S|. We could simply choose the hash-table size very large so that we don't get any collisions.

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6. Feb. 2022

304/616

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Can we get an upper bound on the probability of having collisions?

The probability of having 1 or more collisions can be at most $\frac{1}{2}$ as otherwise the expectation would be larger than $\frac{1}{2}$.



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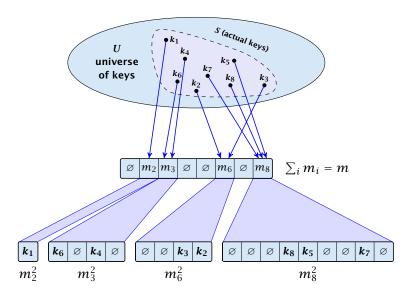
We construct a two-level scheme. We first use a hash-function that maps elements from ${\cal S}$ to ${\cal m}$ buckets.

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However, a hash-table size of $n = m^2$ is very very high.

We construct a two-level scheme. We first use a hash-function that maps elements from ${\it S}$ to ${\it m}$ buckets.

Let m_j denote the number of items that are hashed to the j-th bucket. For each bucket we choose a second hash-function that maps the elements of the bucket into a table of size m_j^2 . The second function can be chosen such that all elements are mapped to different locations.





6. Feb. 2022

The total memory that is required by all hash-tables is $\mathcal{O}(\sum_j m_j^2)$. Note that m_j is a random variable.

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The first expectation is simply the expected number of collisions, for the first level. Since we use universal hashing we have

$$= 2\binom{m}{2} \frac{1}{m} + m = 2m - 1 .$$

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We need only $\mathcal{O}(m)$ time to construct a hash-function h with $\sum_j m_j^2 = \mathcal{O}(4m)$, because with probability at least 1/2 a random function from a universal family will have this property.

Then we construct a hash-table h_j for every bucket. This takes expected time $\mathcal{O}(m_j)$ for every bucket. A random function h_j is collision-free with probability at least 1/2. We need $\mathcal{O}(m_j)$ to test this.

We only need that the hash-functions are chosen from a universal family!!!

6. Feb. 2022

Goal:

Try to generate a hash-table with constant worst-case search time in a dynamic scenario.

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► Two hash-tables $T_1[0,...,n-1]$ and $T_2[0,...,n-1]$, with hash-functions h_1 , and h_2 .

309/616

Goal:

Try to generate a hash-table with constant worst-case search time in a dynamic scenario.

- ▶ Two hash-tables $T_1[0,...,n-1]$ and $T_2[0,...,n-1]$, with hash-functions h_1 , and h_2 .
- ▶ An object x is either stored at location $T_1[h_1(x)]$ or $T_2[h_2(x)]$.

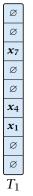
Goal:

Try to generate a hash-table with constant worst-case search time in a dynamic scenario.

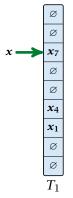
- ▶ Two hash-tables $T_1[0,...,n-1]$ and $T_2[0,...,n-1]$, with hash-functions h_1 , and h_2 .
- ▶ An object x is either stored at location $T_1[h_1(x)]$ or $T_2[h_2(x)]$.
- A search clearly takes constant time if the above constraint is met.

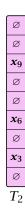
6. Feb. 2022 **309/616**

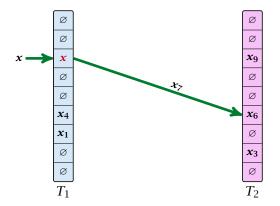
Insert:

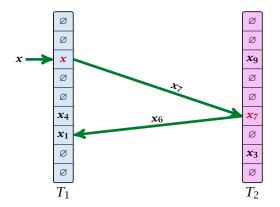


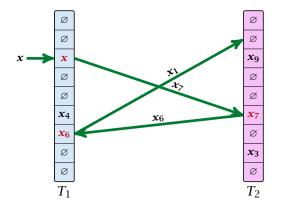
Ø Ø x_9 Ø Ø x_6 Ø \boldsymbol{x}_3 T_2











Algorithm 38 Cuckoo-Insert(x)

```
1: if T_1[h_1(x)] = x \vee T_2[h_2(x)] = x then return

2: steps \leftarrow 1

3: while steps \leq maxsteps do

4: exchange x and T_1[h_1(x)]

5: if x = \text{null} then return

6: exchange x and T_2[h_2(x)]

7: if x = \text{null} then return

8: steps \leftarrow steps +1

9: rehash() // change hash-functions; rehash everything

10: Cuckoo-Insert(x)
```

► We call one iteration through the while-loop a step of the algorithm.

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- We call a sequence of iterations through the while-loop without the termination condition becoming true a phase of the algorithm.

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- We call a sequence of iterations through the while-loop without the termination condition becoming true a phase of the algorithm.
- We say a phase is successful if it is not terminated by the maxstep-condition, but the while loop is left because x = null.

7.7 Hashing 312/616

What is the expected time for an insert-operation?

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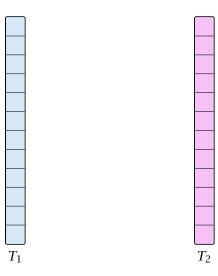
We first analyze the probability that we end-up in an infinite loop (that is then terminated after maxsteps steps).

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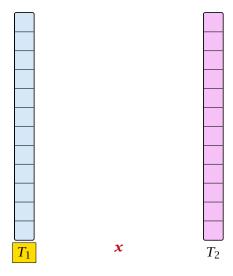
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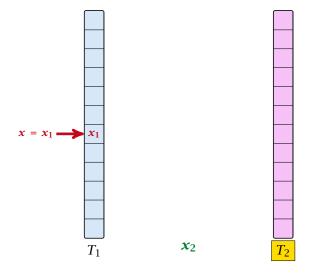
Formally what is the probability to enter an infinite loop that touches s different keys?

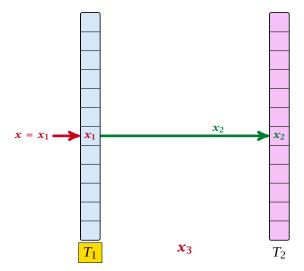
Cuckoo Hashing: Insert

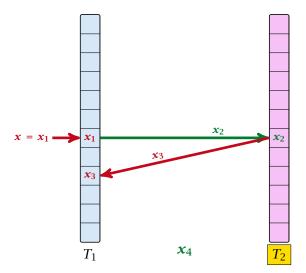


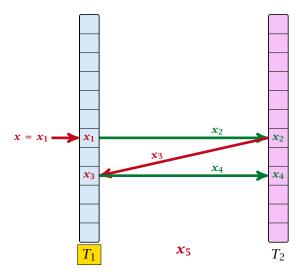
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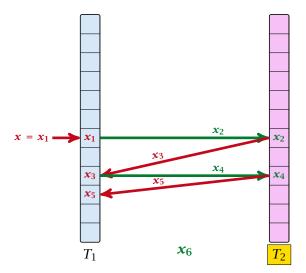


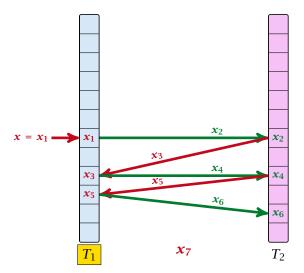




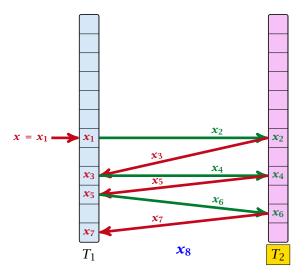




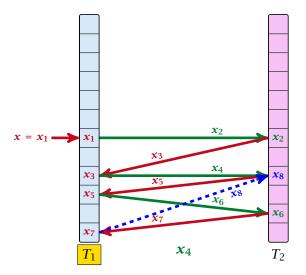


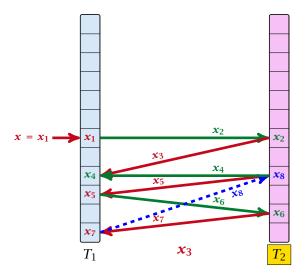




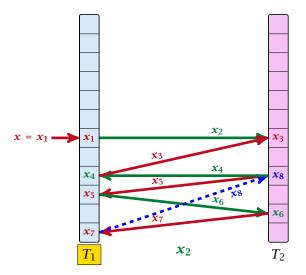


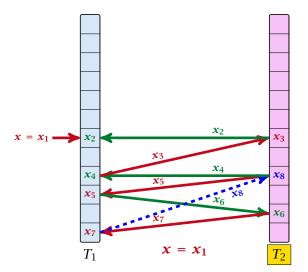


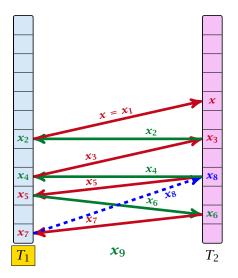


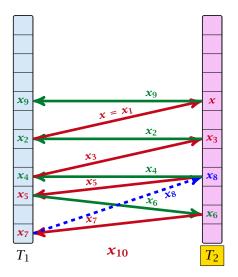


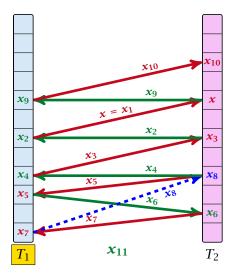


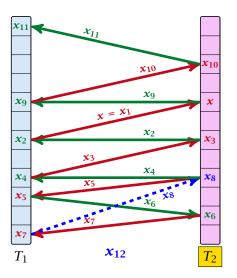


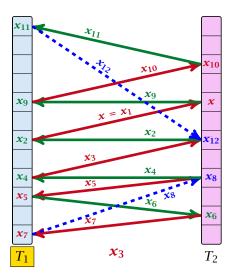


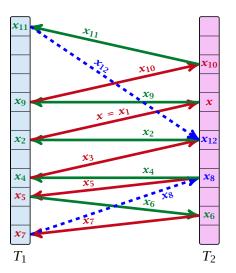


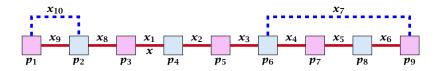




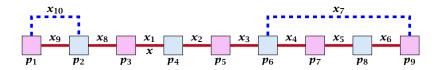






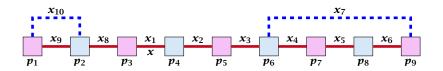


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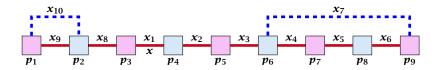
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A cycle-structure of size s is defined by

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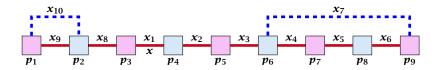
7.7 Hashing



A cycle-structure of size s is defined by

- ▶ s-1 different cells (alternating btw. cells from T_1 and T_2).
- *s* distinct keys $x = x_1, x_2, ..., x_s$, linking the cells.
- The leftmost cell is "linked forward" to some cell on the right.

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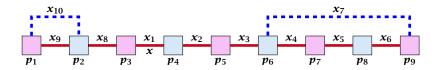


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- ▶ s-1 different cells (alternating btw. cells from T_1 and T_2).
- *s* distinct keys $x = x_1, x_2, ..., x_s$, linking the cells.
- The leftmost cell is "linked forward" to some cell on the right.
- The rightmost cell is "linked backward" to a cell on the left.
- ▶ One link represents key x; this is where the counting starts.



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A cycle-structure is active if for every key x_{ℓ} (linking a cell p_i from T_1 and a cell p_j from T_2) we have

$$h_1(x_\ell) = p_i$$
 and $h_2(x_\ell) = p_i$

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$$h_1(x_\ell) = p_i$$
 and $h_2(x_\ell) = p_j$

Observation:

If during a phase the insert-procedure runs into a cycle there must exist an active cycle structure of size $s \ge 3$.

What is the probability that all keys in a cycle-structure of size s correctly map into their T_1 -cell?

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This probability is at most $\frac{\mu}{n^s}$ since h_1 is a (μ,s) -independent hash-function.

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What is the probability that all keys in the cycle-structure of size s correctly map into their T_2 -cell?

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These events are independent.

The probability that a given cycle-structure of size s is active is at most $\frac{\mu^2}{n^{2s}}$.

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What is the probability that there exists an active cycle structure of size *s*?

The number of cycle-structures of size s is at most

$$s^3 \cdot n^{s-1} \cdot m^{s-1}$$
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- ▶ There are m^{s-1} possibilities to choose the keys apart from x.
- ▶ There are n^{s-1} possibilities to choose the cells.

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The probability that there exists an active cycle-structure is therefore at most

$$\sum_{s=3}^{\infty} s^3 \cdot n^{s-1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}}$$

The probability that there exists an active cycle-structure is therefore at most

$$\sum_{s=3}^{\infty} s^3 \cdot n^{s-1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}} = \frac{\mu^2}{nm} \sum_{s=3}^{\infty} s^3 \left(\frac{m}{n}\right)^s$$

The probability that there exists an active cycle-structure is therefore at most

$$\sum_{s=3}^{\infty} s^3 \cdot n^{s-1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}} = \frac{\mu^2}{nm} \sum_{s=3}^{\infty} s^3 \left(\frac{m}{n}\right)^s$$

$$\leq \frac{\mu^2}{m^2} \sum_{s=3}^{\infty} s^3 \left(\frac{1}{1+\epsilon}\right)^s$$

The probability that there exists an active cycle-structure is therefore at most

$$\begin{split} \sum_{s=3}^{\infty} s^3 \cdot n^{s-1} \cdot m^{s-1} \cdot \frac{\mu^2}{n^{2s}} &= \frac{\mu^2}{nm} \sum_{s=3}^{\infty} s^3 \left(\frac{m}{n}\right)^s \\ &\leq \frac{\mu^2}{m^2} \sum_{s=3}^{\infty} s^3 \left(\frac{1}{1+\epsilon}\right)^s \leq \mathcal{O}\left(\frac{1}{m^2}\right) \; . \end{split}$$

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Here we used the fact that $(1 + \epsilon)m \le n$.

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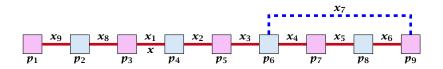
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Here we used the fact that $(1 + \epsilon)m \le n$.

Hence,

$$\Pr[\mathsf{cycle}] = \mathcal{O}\left(\frac{1}{m^2}\right)$$
.

Now, we analyze the probability that a phase is not successful without running into a closed cycle.



Sequence of visited keys:

$$x = x_1, x_2, x_3, x_4, x_5, x_6, x_7, x_3, x_2, x_1 = x, x_8, x_9, \dots$$

Consider the sequence of not necessarily distinct keys starting with \boldsymbol{x} in the order that they are visited during the phase.

Consider the sequence of not necessarily distinct keys starting with \boldsymbol{x} in the order that they are visited during the phase.

Lemma 30

If the sequence is of length p then there exists a sub-sequence of at least $\frac{p+2}{3}$ keys starting with x of distinct keys.

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Proof.

Let i be the number of keys (including x) that we see before the first repeated key. Let j denote the total number of distinct keys.

The sequence is of the form:

$$x = x_1 \rightarrow x_2 \rightarrow \cdots \rightarrow x_i \rightarrow x_r \rightarrow x_{r-1} \rightarrow \cdots \rightarrow x_1 \rightarrow x_{i+1} \rightarrow \cdots \rightarrow x_j$$

As $r \le i - 1$ the length p of the sequence is

$$p = i + r + (j - i) \le i + j - 1$$
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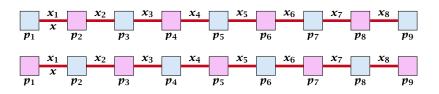
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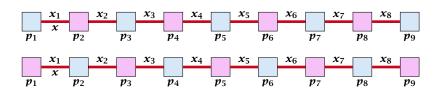
Either sub-sequence $x_1 \rightarrow x_2 \rightarrow \cdots \rightarrow x_i$ or sub-sequence $x_1 \rightarrow x_{i+1} \rightarrow \cdots \rightarrow x_j$ has at least $\frac{p+2}{3}$ elements.





A path-structure of size s is defined by

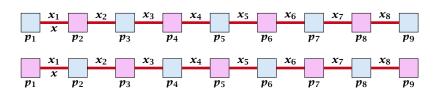
7.7 Hashing



A path-structure of size s is defined by

ightharpoonup s+1 different cells (alternating btw. cells from T_1 and T_2).

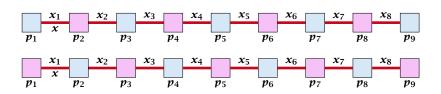
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- ightharpoonup s + 1 different cells (alternating btw. cells from T_1 and T_2).
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- ightharpoonup s + 1 different cells (alternating btw. cells from T_1 and T_2).
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- ▶ The leftmost cell is either from T_1 or T_2 .

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A path-structure is active if for every key x_ℓ (linking a cell p_i from T_1 and a cell p_j from T_2) we have

$$h_1(x_\ell) = p_i$$
 and $h_2(x_\ell) = p_j$

Observation:

If a phase takes at least t steps without running into a cycle there must exist an active path-structure of size (2t+2)/3.

The probability that a given path-structure of size s is active is at most $\frac{\mu^2}{n^{2s}}$.

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The probability that there exists an active path-structure of size s is at most

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$$\leq 2\mu^2 \left(\frac{m}{n}\right)^{s-1}$$

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Plugging in s = (2t + 2)/3 gives

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$$\leq 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2t+2)/3-1}$$

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$$\leq 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2t+2)/3-1} = 2\mu^2 \left(\frac{1}{1+\epsilon}\right)^{(2t-1)/3} \ .$$

We choose maxsteps $\geq 3\ell/2 + 1/2$.

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\begin{split} & Pr[\text{unsuccessful} \mid \text{no cycle}] \\ & \leq Pr[\exists \text{ active path-structure of size at least } \tfrac{2\text{maxsteps}+2}{3}] \end{split}
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$$\ell \geq \log{(\frac{1}{2\mu^2m^2})}/\log{(\frac{1}{1+\epsilon})} = \log{(2\mu^2m^2)}/\log{(1+\epsilon)}$$

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This gives maxsteps = $\Theta(\log m)$.

So far we estimated

$$\Pr[\mathsf{cycle}] \leq \mathcal{O}\left(\frac{1}{m^2}\right)$$

and

$$\Pr[\mathsf{unsuccessful} \mid \mathsf{no} \; \mathsf{cycle}] \leq \mathcal{O}\left(\frac{1}{m^2}\right)$$

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Pr[successful] = Pr[no cycle] - Pr[unsuccessful | no cycle]

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Observe that

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for a suitable constant c > 0.

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Pr[search at least t steps | successful]

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We have

Pr[search at least *t* steps | successful]

= $Pr[search at least t steps \land successful] / Pr[successful]$

The expected number of complete steps in the successful phase of an insert operation is:

$$\begin{split} & \text{E}[\text{number of steps} \mid \text{phase successful}] \\ &= \sum_{t \geq 1} \Pr[\text{search takes at least } t \text{ steps} \mid \text{phase successful}] \end{split}$$

We have

```
\begin{split} \Pr[\mathsf{search} \ \mathsf{at} \ \mathsf{least} \ t \ \mathsf{steps} \ | \ \mathsf{successful}] \\ &= \Pr[\mathsf{search} \ \mathsf{at} \ \mathsf{least} \ t \ \mathsf{steps} \ \land \ \mathsf{successful}] / \Pr[\mathsf{successful}] \\ &\leq \frac{1}{c} \Pr[\mathsf{search} \ \mathsf{at} \ \mathsf{least} \ t \ \mathsf{steps} \ \land \ \mathsf{successful}] / \Pr[\mathsf{no} \ \mathsf{cycle}] \end{split}
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Hence,

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Hence,

E[number of steps | phase successful]

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This means the expected cost for a successful phase is constant (even after accounting for the cost of the incomplete step that finishes the phase).

A phase that is not successful induces cost for doing a complete rehash (this dominates the cost for the steps in the phase).

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The probability that a phase is not successful is $q = O(1/m^2)$ (probability $\mathcal{O}(1/m^2)$ of running into a cycle and probability $\mathcal{O}(1/m^2)$ of reaching maxsteps without running into a cycle).

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A rehash try requires m insertions and takes expected constant time per insertion. It fails with probability p := O(1/m).

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The expected number of unsuccessful rehashes is $\sum_{i\geq 1} p^i = \frac{1}{1-p} - 1 = \frac{p}{1-p} = \mathcal{O}(p)$.



7.7 Hashing

A phase that is not successful induces cost for doing a complete rehash (this dominates the cost for the steps in the phase).

The probability that a phase is not successful is $q = \mathcal{O}(1/m^2)$ (probability $\mathcal{O}(1/m^2)$ of running into a cycle and probability $\mathcal{O}(1/m^2)$ of reaching maxsteps without running into a cycle).

A rehash try requires m insertions and takes expected constant time per insertion. It fails with probability $p := \mathcal{O}(1/m)$.

The expected number of unsuccessful rehashes is $\sum_{i\geq 1} p^i = \frac{1}{1-p} - 1 = \frac{p}{1-p} = \mathcal{O}(p)$.

Therefore the expected cost for re-hashes is $\mathcal{O}(m) \cdot \mathcal{O}(p) = \mathcal{O}(1)$.



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Let Y_i denote the event that the i-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

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Let Z_i denote the event that the i-th rehash occurs:

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Let Z_i denote the event that the *i*-th rehash occurs:

The 0-th (re)hash is the initial configuration when doing the $\Pr[Z_i] \leq \Pr[\wedge_{j=0}^{i-1} Y_j] \leq p^i$ insert.

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Let X_i^s , $s \in \{1, ..., m+1\}$ denote the cost for inserting the s-th element during the *i*-th rehash (assuming *i*-th rehash occurs):

$$E[X_i^s]$$

Let Y_i denote the event that the *i*-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

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$$\begin{split} \mathbf{E}[X_i^s] &= \mathbf{E}[\mathsf{steps} \mid \mathsf{phase} \; \mathsf{successful}] \cdot \Pr[\mathsf{phase} \; \mathsf{sucessful}] \\ &+ \mathsf{maxsteps} \cdot \Pr[\mathsf{not} \; \mathsf{sucessful}] \end{split}$$

Let Y_i denote the event that the *i*-th rehash occurs and does not lead to a valid configuration (i.e., one of the m+1 insertions fails):

$$\Pr[Y_i|Z_i] \le (m+1) \cdot \mathcal{O}(1/m^2) \le \mathcal{O}(1/m) =: p.$$

Let Z_i denote the event that the *i*-th rehash occurs:

The 0-th (re)hash is the initial

configuration when doing the insert.
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Therefore, it is sufficient to have $(\mu, \Theta(\log m))$ -independent hash-functions.

How do we make sure that $n \ge (1 + \epsilon)m$?

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7.7 Hashing 6. Feb. 2022

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6. Feb. 2022 336/616

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6. Feb. 2022

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- Therefore we can amortize the rehash cost after a change in table-size against the cost for insertions and deletions.

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Lemma 31

Cuckoo Hashing has an expected constant insert-time and a worst-case constant search-time.

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Note that the above lemma only holds if the fill-factor (number of keys/total number of hash-table slots) is at most $\frac{1}{2(1+\epsilon)}$.

The $1/(2(1+\epsilon))$ fill-factor comes from the fact that the total hash-table is of size 2n (because we have two tables of size n); moreover $m \le (1+\epsilon)n$.

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Sometimes we also have

▶ *S.* merge(S'): $S := S \cup S'$; $S' := \emptyset$.



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- handle S. insert(x): Adds element x to the data-structure, and returns a handle to the object for future reference.
- **S.** delete(h): Deletes element specified through handle h.
- S. decrease-key(h, k): Decreases the key of the element specified by handle h to k. Assumes that the key is at least k before the operation.

Dijkstra's Shortest Path Algorithm

```
Algorithm 1 Shortest-Path(G = (V, E, d), s \in V)
 1: Input: weighted graph G = (V, E, d); start vertex s;
 2: Output: key-field of every node contains distance from s;
 3: S.build(); // build empty priority queue
 4: for all v \in V \setminus \{s\} do
 5: v \cdot \text{key} \leftarrow \infty;
 6: h_v \leftarrow S.insert(v);
 7: s. \text{key} \leftarrow 0; S. \text{insert}(s);
 8: while S.is-empty() = false do
 9:
     v \leftarrow S. delete-min():
10: for all x \in V s.t. (v, x) \in E do
11:
                if x. key > v. key +d(v,x) then
12:
                      S.decrease-key(h_x, v. key + d(v, x));
13:
                      x. \text{key} \leftarrow v. \text{key} + d(v, x);
```

Prim's Minimum Spanning Tree Algorithm

```
Algorithm 2 Prim-MST(G = (V, E, d), s \in V)
 1: Input: weighted graph G = (V, E, d); start vertex s;
 2: Output: pred-fields encode MST;
 3: S.build(); // build empty priority queue
 4: for all v \in V \setminus \{s\} do
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 6: h_v \leftarrow S.insert(v);
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12:
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                      x. key \leftarrow d(v, x);
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14:
                      x. pred \leftarrow v:
```

Analysis of Dijkstra and Prim

Both algorithms require:

- 1 build() operation
- ightharpoonup |V| insert() operations
- ▶ |V| delete-min() operations
- ightharpoonup |V| is-empty() operations
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Analysis of Dijkstra and Prim

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How good a running time can we obtain?

Operation	Binary Heap	BST	Binomial Heap	Fibonacci Heap*
build	n	$n \log n$	$n \log n$	n
minimum	1	$\log n$	$\log n$	1
is-empty	1	1	1	1
insert	$\log n$	$\log n$	$\log n$	1
delete	$\log n^{**}$	$\log n$	$\log n$	$\log n$
delete-min	$\log n$	$\log n$	$\log n$	$\log n$
decrease-key	$\log n$	$\log n$	$\log n$	1
merge	n	$n \log n$	$\log n$	1

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Note that most applications use $\mathbf{build}()$ only to create an empty heap which then costs time 1.

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The standard version of binary heaps is not addressable, and hence does not support a delete operation.

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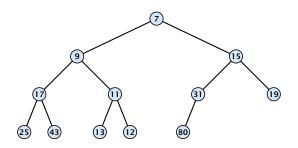
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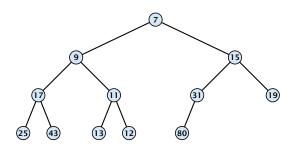
Fibonacci heaps only give an amortized guarantee.

Using Binary Heaps, Prim and Dijkstra run in time $\mathcal{O}((|V|+|E|)\log |V|)$.

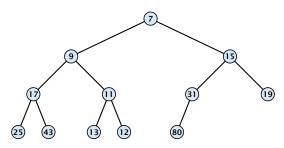
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Nearly complete binary tree; only the last level is not full, and this one is filled from left to right.



- Nearly complete binary tree; only the last level is not full, and this one is filled from left to right.
- Heap property: A node's key is not larger than the key of one of its children.



Binary Heaps

Operations:

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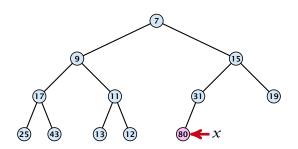
minimum(): return the root-element. Time O(1).

Binary Heaps

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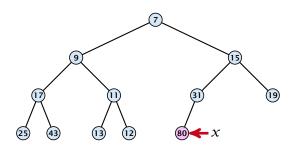
- **minimum():** return the root-element. Time $\mathcal{O}(1)$.
- **is-empty():** check whether root-pointer is null. Time $\mathcal{O}(1)$.

Maintain a pointer to the last element x.



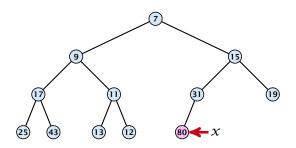
Maintain a pointer to the last element x.

We can compute the predecessor of x (last element when x is deleted) in time $\mathcal{O}(\log n)$.



Maintain a pointer to the last element x.

We can compute the predecessor of x (last element when x is deleted) in time O(log n). go up until the last edge used was a right edge. go left; go right until you reach a leaf

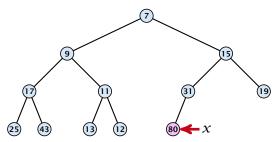


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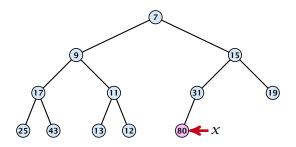
We can compute the predecessor of x (last element when x is deleted) in time $\mathcal{O}(\log n)$.

go up until the last edge used was a right edge. go left; go right until you reach a leaf

if you hit the root on the way up, go to the rightmost element

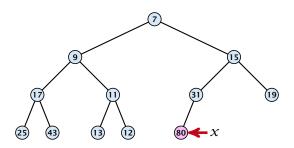


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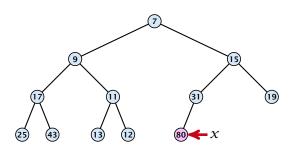
• We can compute the successor of x (last element when an element is inserted) in time $\mathcal{O}(\log n)$.



Maintain a pointer to the last element x.

We can compute the successor of *x* (last element when an element is inserted) in time *O*(log *n*).

 go up until the last edge used was a left edge.
 go right; go left until you reach a null-pointer.

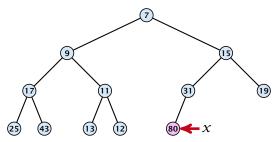


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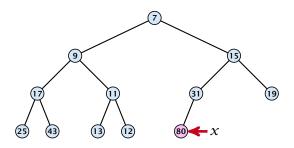
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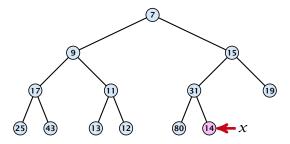
if you hit the root on the way up, go to the leftmost element; insert a new element as a left child;



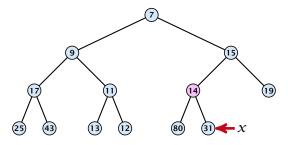
1. Insert element at successor of x.



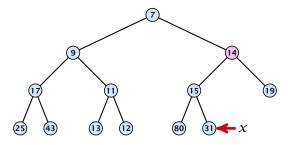
- 1. Insert element at successor of x.
- 2. Exchange with parent until heap property is fulfilled.



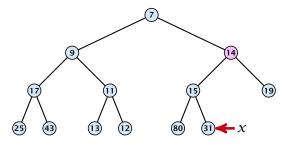
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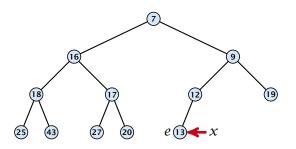


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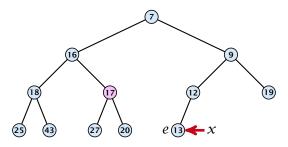


Note that an exchange can either be done by moving the data or by changing pointers. The latter method leads to an addressable priority queue.

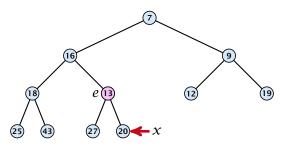
1. Exchange the element to be deleted with the element e pointed to by x.



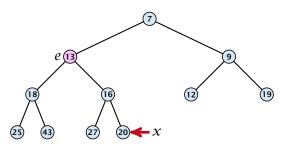
- 1. Exchange the element to be deleted with the element e pointed to by x.
- **2.** Restore the heap-property for the element e.



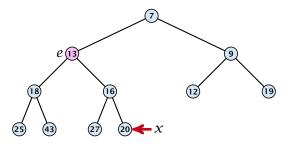
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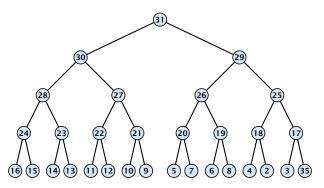
At its new position e may either travel up or down in the tree (but not both directions).

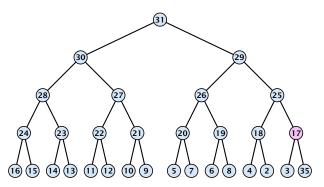
Binary Heaps

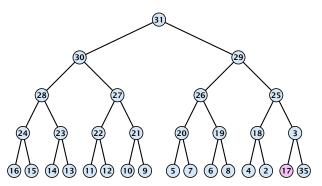
Operations:

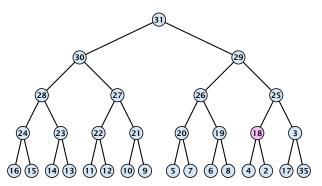
- **minimum()**: return the root-element. Time O(1).
- **is-empty():** check whether root-pointer is null. Time O(1).
- insert(k): insert at successor of x and bubble up. Time $O(\log n)$.
- **delete**(h): swap with x and bubble up or sift-down. Time $O(\log n)$.

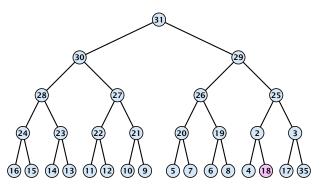
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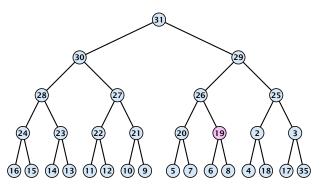


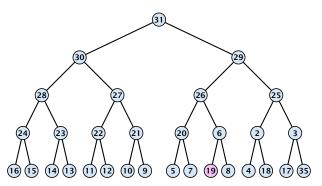


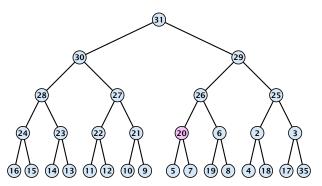


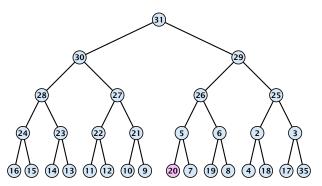


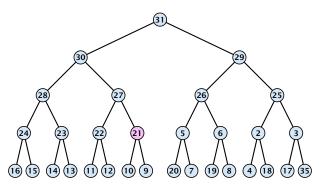


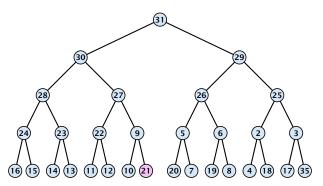


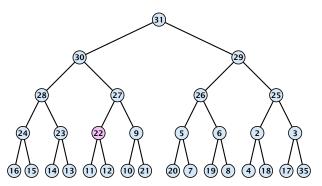


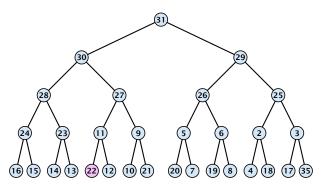


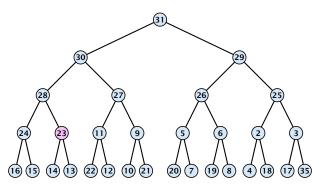


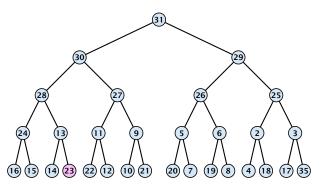


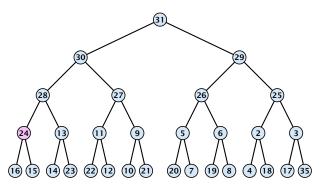


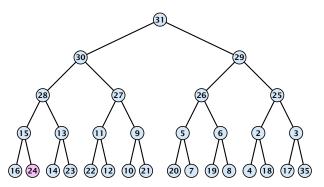


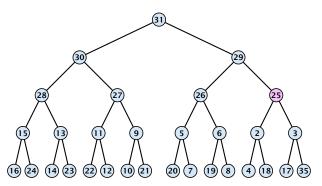


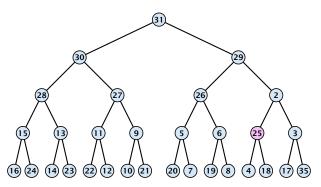


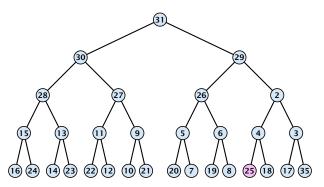


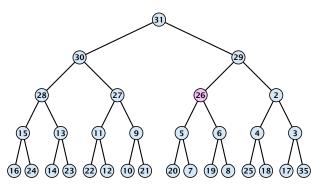


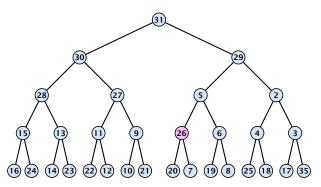


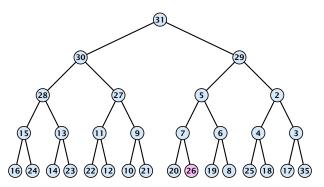


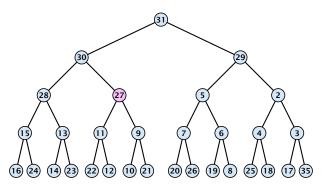


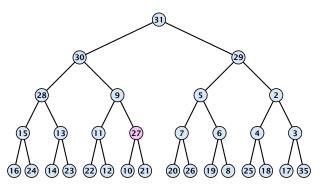


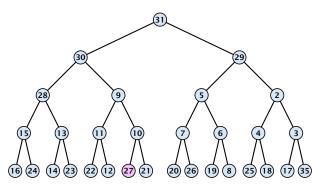


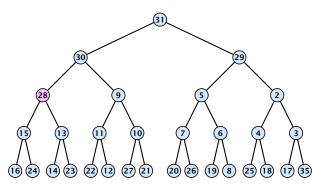


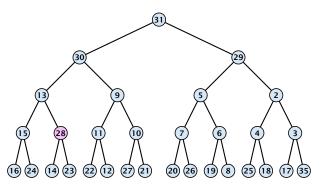


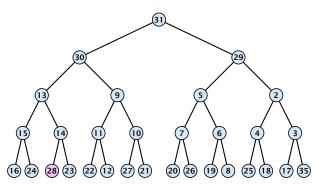


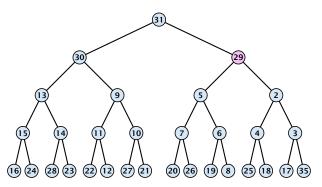


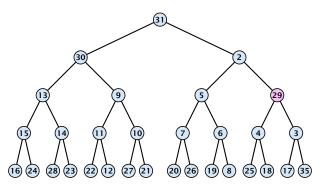


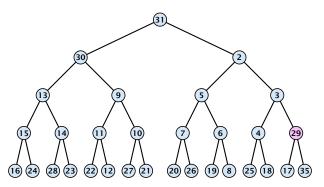


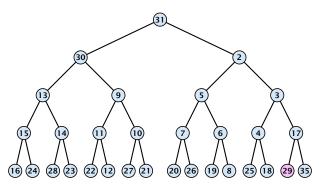


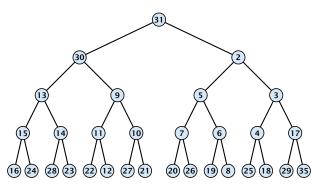


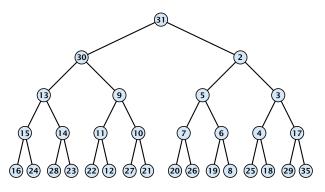












$$\sum_{\text{levels } \ell} 2^{\ell} \cdot (h - \ell) = \sum_{i} i 2^{h - i} = \mathcal{O}(2^h) = \mathcal{O}(n)$$

Operations:

- **minimum():** Return the root-element. Time $\mathcal{O}(1)$.
- **is-empty():** Check whether root-pointer is null. Time O(1).
- ▶ insert(k): Insert at x and bubble up. Time O(log n).
- **delete**(h): Swap with x and bubble up or sift-down. Time $O(\log n)$.
- **build** (x_1, \ldots, x_n) : Insert elements arbitrarily; then do sift-down operations starting with the lowest layer in the tree. Time $\mathcal{O}(n)$.



The standard implementation of binary heaps is via arrays. Let A[0,...,n-1] be an array

- ▶ The parent of i-th element is at position $\lfloor \frac{i-1}{2} \rfloor$.
- ▶ The left child of i-th element is at position 2i + 1.
- ► The right child of i-th element is at position 2i + 2.

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The resulting binary heap is not addressable. The elements don't maintain their positions and therefore there are no stable handles.

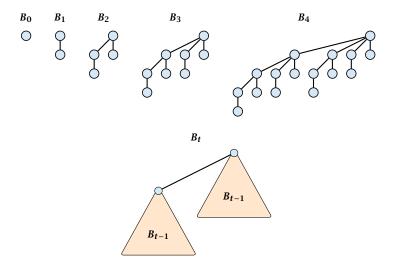
6. Feb. 2022

8.2 Binomial Heaps

Operation	Binary Heap	BST	Binomial Heap	Fibonacci Heap*
build	n	$n \log n$	$n \log n$	n
minimum	1	$\log n$	$\log n$	1
is-empty	1	1	1	1
insert	$\log n$	$\log n$	$\log n$	1
delete	$\log n^{**}$	$\log n$	$\log n$	$\log n$
delete-min	$\log n$	$\log n$	$\log n$	$\log n$
decrease-key	$\log n$	$\log n$	$\log n$	1
merge	n	$n \log n$	$\log n$	1



6. Feb. 2022



Properties of Binomial Trees

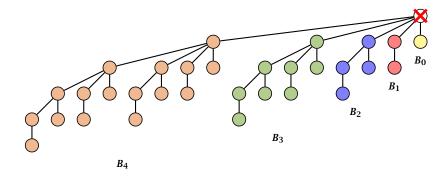
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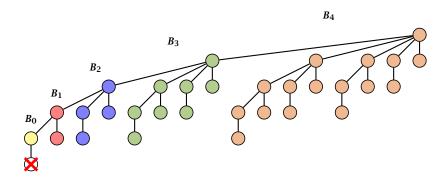
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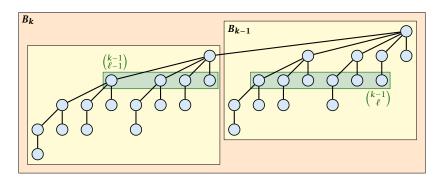
- \triangleright B_k has 2^k nodes.
- \triangleright B_k has height k.
- ▶ The root of B_k has degree k.
- ▶ B_k has $\binom{k}{\ell}$ nodes on level ℓ .
- Deleting the root of B_k gives trees $B_0, B_1, \ldots, B_{k-1}$.



Deleting the root of B_5 leaves sub-trees B_4 , B_3 , B_2 , B_1 , and B_0 .

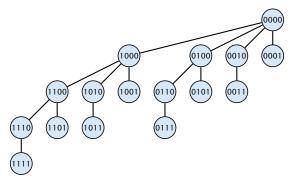


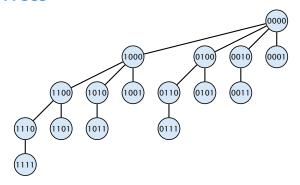
Deleting the leaf furthest from the root (in B_5) leaves a path that connects the roots of sub-trees B_4 , B_3 , B_2 , B_1 , and B_0 .



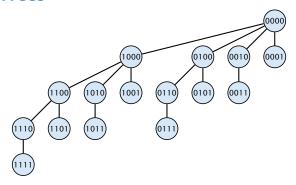
The number of nodes on level ℓ in tree B_k is therefore

$$\binom{k-1}{\ell-1}+\binom{k-1}{\ell}=\binom{k}{\ell}$$



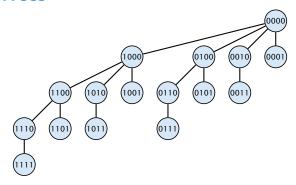


The binomial tree B_k is a sub-graph of the hypercube H_k .



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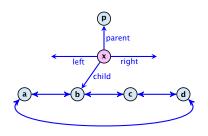
The parent of a node with label $b_k, ..., b_1$ is obtained by setting the least significant 1-bit to 0.

The ℓ -th level contains nodes that have ℓ 1's in their label.



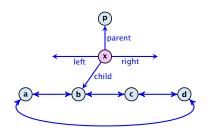
How do we implement trees with non-constant degree?

The children of a node are arranged in a circular linked list.



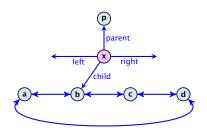
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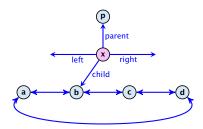
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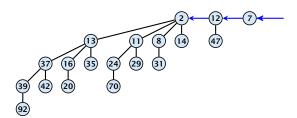


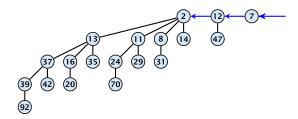
How do we implement trees with non-constant degree?

- The children of a node are arranged in a circular linked list.
- A child-pointer points to an arbitrary node within the list.
- A parent-pointer points to the parent node.
- Pointers x. left and x. right point to the left and right sibling of x (if x does not have siblings then x. left = x. right = x).

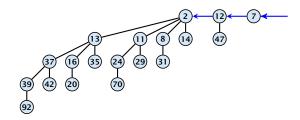


- \blacktriangleright Given a pointer to a node x we can splice out the sub-tree rooted at x in constant time.
- We can add a child-tree T to a node x in constant time if we are given a pointer to x and a pointer to the root of T.



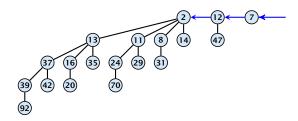


In a binomial heap the keys are arranged in a collection of binomial trees.



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Every tree fulfills the heap-property



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Every tree fulfills the heap-property

There is at most one tree for every dimension/order. For example the above heap contains trees B_0 , B_1 , and B_4 .

Given the number n of keys to be stored in a binomial heap we can deduce the binomial trees that will be contained in the collection.

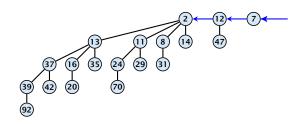
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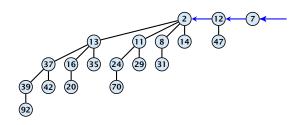
Let B_{k_1} , B_{k_2} , B_{k_3} , $k_i < k_{i+1}$ denote the binomial trees in the collection and recall that every tree may be contained at most once.

Then $n=\sum_i 2^{k_i}$ must hold. But since the k_i are all distinct this means that the k_i define the non-zero bit-positions in the binary representation of n.

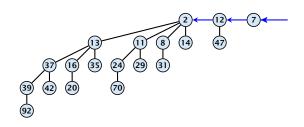


Properties of a heap with n keys:

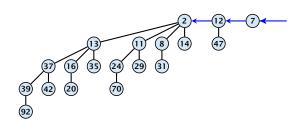
Let $n = b_d b_{d-1}, \dots, b_0$ denote binary representation of n.



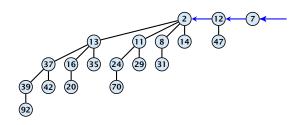
- Let $n = b_d b_{d-1}, \dots, b_0$ denote binary representation of n.
- ▶ The heap contains tree B_i iff $b_i = 1$.



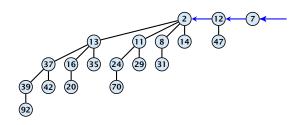
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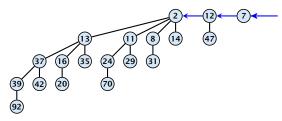
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- The minimum must be contained in one of the roots.
- ▶ The height of the largest tree is at most $\lfloor \log n \rfloor$.
- The trees are stored in a single-linked list; ordered by dimension/size.



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A merge is easy if we have two heaps with different binomial trees. We can simply merge the tree-lists.

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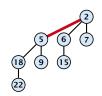
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Merging two trees of the same size: Add the tree with larger root-value as a child to the other tree.



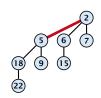
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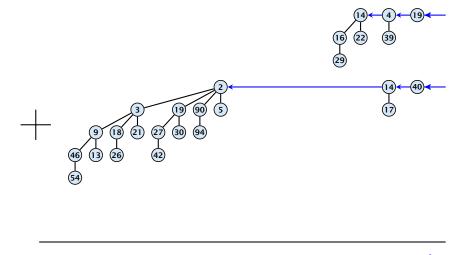
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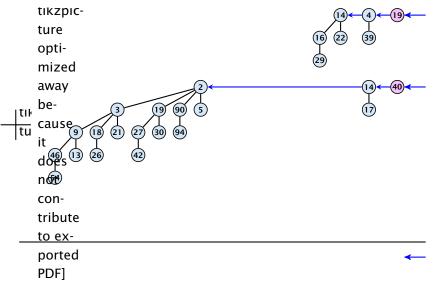
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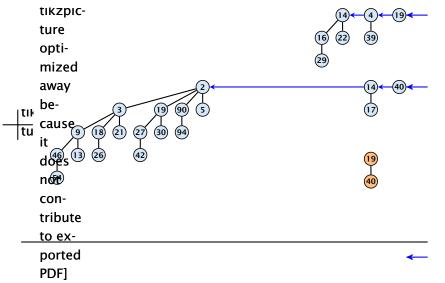
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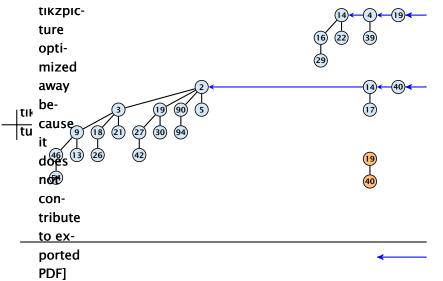
For more trees the technique is analogous to binary addition.

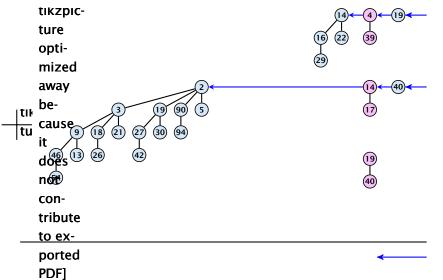


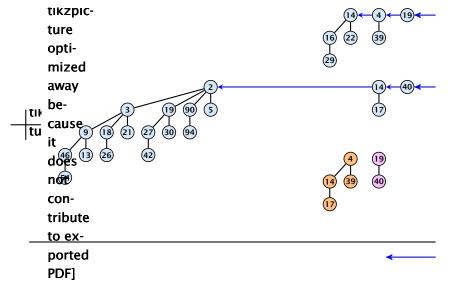


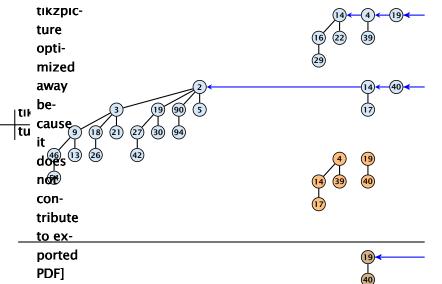


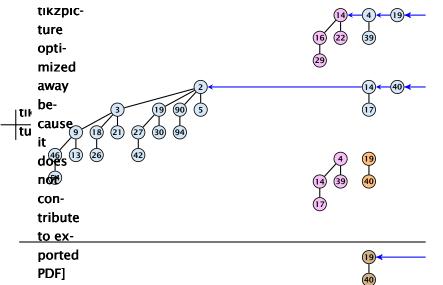


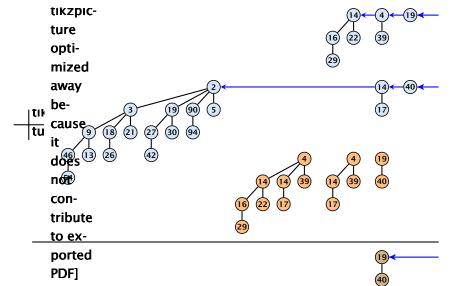


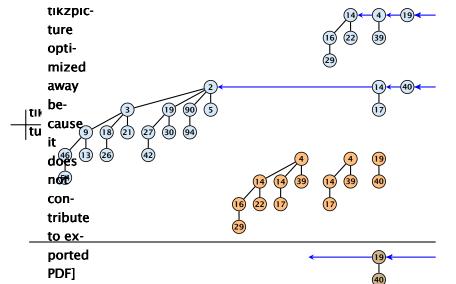


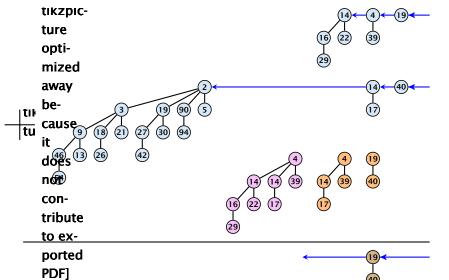


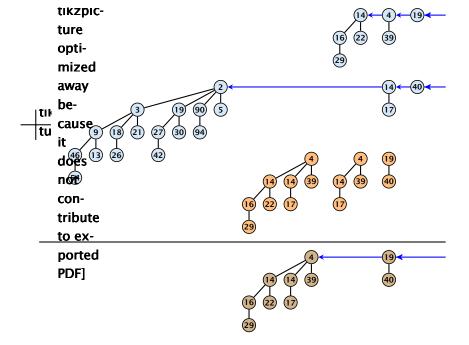


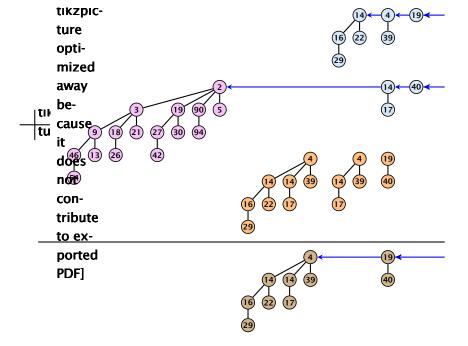


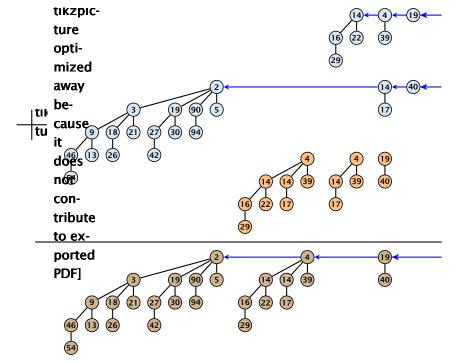


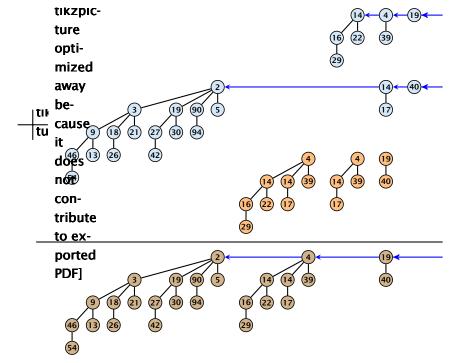












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• Create a new heap S' that contains just the element x.

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- Find the minimum key-value among all roots.
- ▶ Time: $O(\log n)$.

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- ▶ Time: $O(\log n)$ since the trees have height $O(\log n)$.

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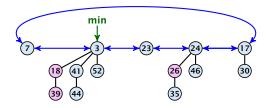
- ► Execute *S*. decrease-key(h, $-\infty$).
- ► Execute *S*. delete-min().

S. delete(handle *h*):

- **Execute** *S*. decrease-key $(h, -\infty)$.
- ► Execute *S*. delete-min().
- ▶ Time: $\mathcal{O}(\log n)$.

Collection of trees that fulfill the heap property.

Structure is much more relaxed than binomial heaps.

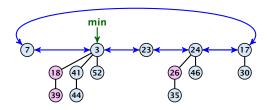


Additional implementation details:

- Every node x stores its degree in a field x. degree. Note that this can be updated in constant time when adding a child to x.
- Every node stores a boolean value x. marked that specifies whether x is marked or not.

The potential function:

- ightharpoonup t(S) denotes the number of trees in the heap.
- \blacktriangleright m(S) denotes the number of marked nodes.
- We use the potential function $\Phi(S) = t(S) + 2m(S)$.



The potential is $\Phi(S) = 5 + 2 \cdot 3 = 11$.

We assume that one unit of potential can pay for a constant amount of work, where the constant is chosen "big enough" (to take care of the constants that occur).

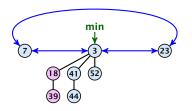
To make this more explicit we use \boldsymbol{c} to denote the amount of work that a unit of potential can pay for.

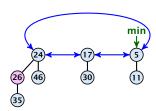
S. minimum()

- Access through the min-pointer.
- Actual cost $\mathcal{O}(1)$.
- No change in potential.
- ▶ Amortized cost $\mathcal{O}(1)$.

S. merge(S')

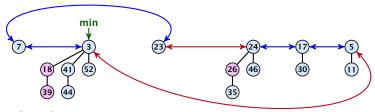
- Merge the root lists.
- Adjust the min-pointer





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- Adjust the min-pointer

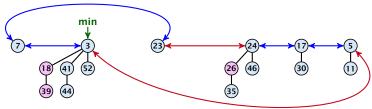


Running time:

Actual cost $\mathcal{O}(1)$.

S. merge(S')

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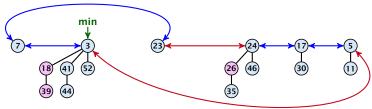


Running time:

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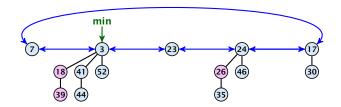
Running time:

- Actual cost $\mathcal{O}(1)$.
- No change in potential.
- \blacktriangleright Hence, amortized cost is $\mathcal{O}(1)$.



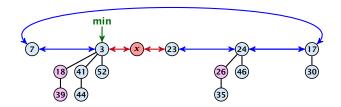
S.insert(x)

- ightharpoonup Create a new tree containing x.
- ► Insert *x* into the root-list.
- Update min-pointer, if necessary.



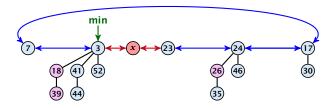
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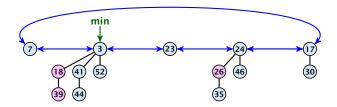


Running time:

- ightharpoonup Actual cost $\mathcal{O}(1)$.
- \triangleright Change in potential is +1.
- ▶ Amortized cost is c + O(1) = O(1).

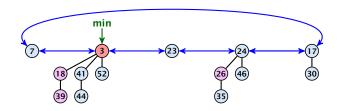


S. delete-min(x)



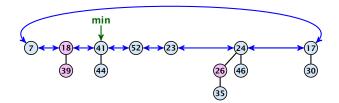
S. delete-min(x)

▶ Delete minimum; add child-trees to heap; time: $D(\min) \cdot \mathcal{O}(1)$.



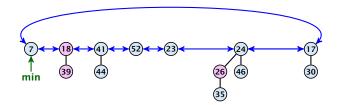
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- Update min-pointer; time: $(t + D(\min)) \cdot \mathcal{O}(1)$.



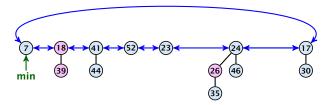
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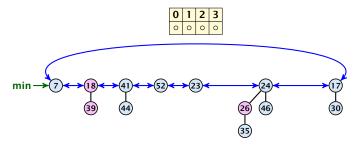


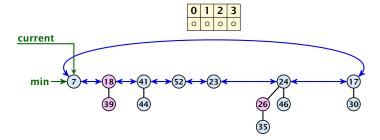
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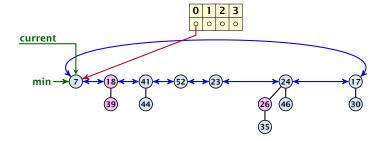
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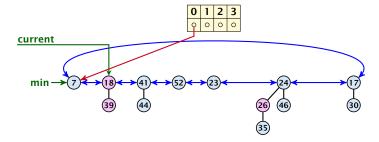


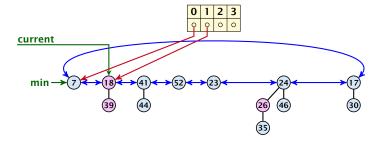
Consolidate root-list so that no roots have the same degree. Time $t\cdot\mathcal{O}(1)$ (see next slide).

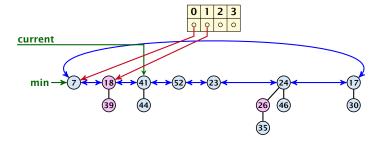


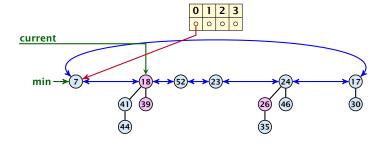


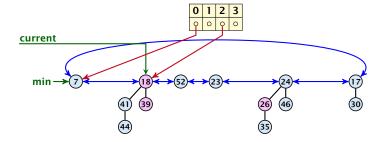


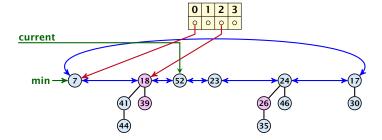


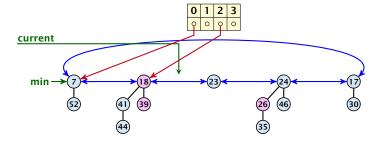


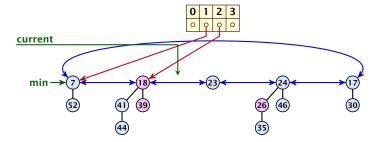


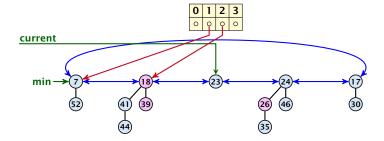


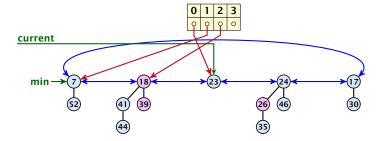


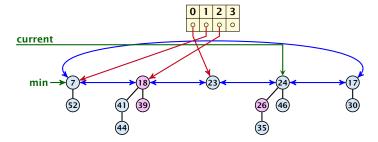


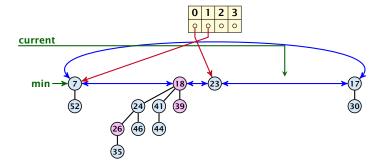


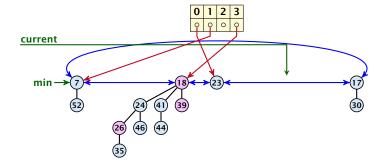


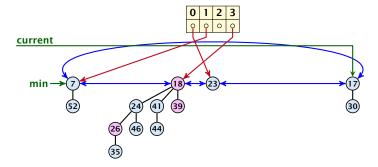


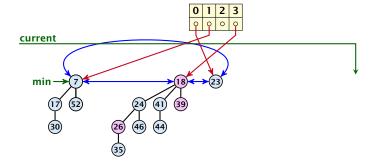


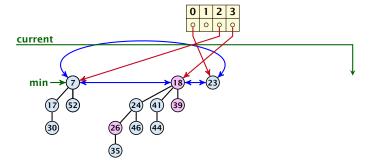


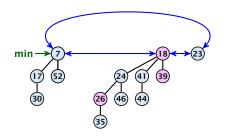












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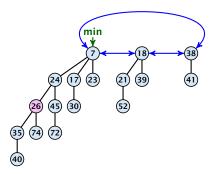
$$\leq (c_1 + c)D_n + (c_1 - c)t + c \leq 2c(D_n + 1) \leq \mathcal{O}(D_n)$$

for $c \ge c_1$.

If the input trees of the consolidation procedure are binomial trees (for example only singleton vertices) then the output will be a set of distinct binomial trees, and, hence, the Fibonacci heap will be (more or less) a Binomial heap right after the consolidation.

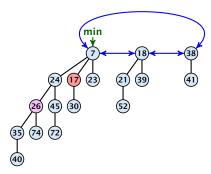
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If we do not have delete or decrease-key operations then $D_n \leq \log n$.



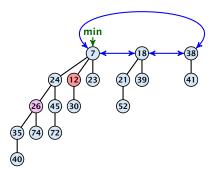
Case 1: decrease-key does not violate heap-property

Just decrease the key-value of element referenced by h. Nothing else to do.



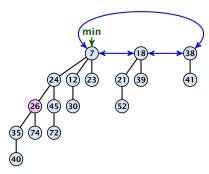
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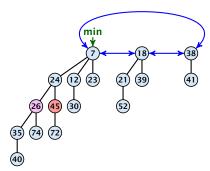
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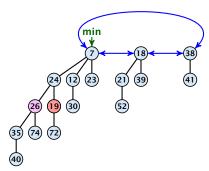
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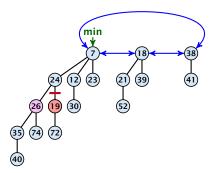
- Decrease key-value of element x reference by h.
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- Adjust min-pointers, if necessary.
- Mark the (previous) parent of x (unless it's a root).





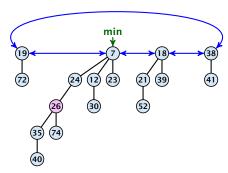
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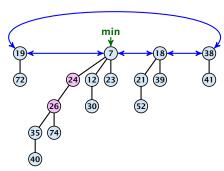
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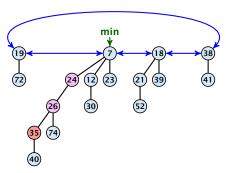
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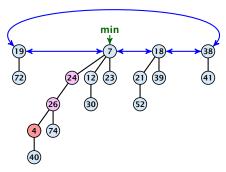
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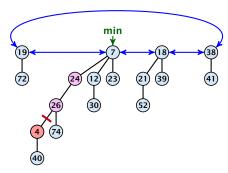
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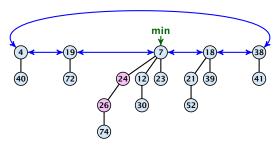
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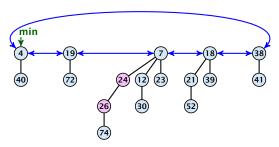
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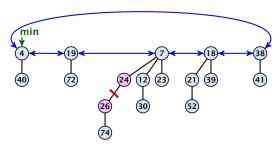
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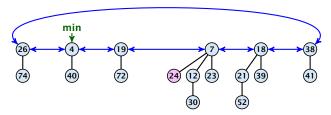
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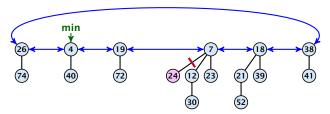
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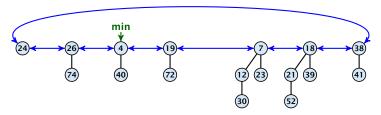
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Constant cost for decreasing the value.

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- ightharpoonup Constant cost for each of ℓ cuts.

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- Constant cost for decreasing the value.
- ▶ Constant cost for each of ℓ cuts.
- ▶ Hence, cost is at most $c_2 \cdot (\ell + 1)$, for some constant c_2 .

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$$c_2(\ell+1) + c(4-\ell) \le (c_2-c)\ell + 4c + c_2 = \mathcal{O}(1)$$
, if $c \ge c_2$.

Delete node

H. delete(x):

- ▶ decrease value of x to $-\infty$.
- delete-min.

Amortized cost: $\mathcal{O}(D_n)$

- $ightharpoonup \mathcal{O}(1)$ for decrease-key.
- \triangleright $\mathcal{O}(D_n)$ for delete-min.

Lemma 32

Let x be a node with degree k and let y_1, \ldots, y_k denote the children of x in the order that they were linked to x. Then

$$\operatorname{degree}(y_i) \geq \left\{ \begin{array}{ll} 0 & \textit{if } i = 1 \\ i - 2 & \textit{if } i > 1 \end{array} \right.$$

Proof

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- ▶ Hence, at this time $degree(x) \ge i 1$, and therefore also $degree(y_i) \ge i 1$ as the algorithm links nodes of equal degree only.
- Since, then y_i has lost at most one child.
- ► Therefore, degree(y_i) $\geq i 2$.

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$$= 2 + \sum_{i=2}^{k-2} s_i$$

 $\phi=rac{1}{2}(1+\sqrt{5})$ denotes the *golden ratio*. Note that $\phi^2=1+\phi$.

Definition 33

Consider the following non-standard Fibonacci type sequence:

$$F_k = \begin{cases} 1 & \text{if } k = 0 \\ 2 & \text{if } k = 1 \\ F_{k-1} + F_{k-2} & \text{if } k \ge 2 \end{cases}$$

Facts:

- 1. $F_k \geq \phi^k$.
- **2.** For $k \ge 2$: $F_k = 2 + \sum_{i=0}^{k-2} F_i$.

The above facts can be easily proved by induction. From this it follows that $s_k \ge F_k \ge \phi^k$, which gives that the maximum degree in a Fibonacci heap is logarithmic.

k=0:
$$1 = F_0 \ge \Phi^0 = 1$$

k=1: $2 = F_1 \ge \Phi^1 \approx 1.61$
k-2,k-1 \rightarrow k: $F_k = F_{k-1} + F_{k-2} \ge \Phi^{k-1} + \Phi^{k-2} = \Phi^{k-2}(\Phi^{+1}) = \Phi^k$

k=2:
$$3 = F_2 = 2 + 1 = 2 + F_0$$

k-1 \rightarrow k: $F_k = F_{k-1} + F_{k-2} = 2 + \sum_{i=0}^{k-3} F_i + F_{k-2} = 2 + \sum_{i=0}^{k-2} F_i$



Union Find Data Structure \mathcal{P} : Maintains a partition of disjoint sets over elements.

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- ▶ **P.** makeset(x): Given an element x, adds x to the data-structure and creates a singleton set that contains only this element. Returns a locator/handle for x in the data-structure.
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Union Find Data Structure \mathcal{P} : Maintains a partition of disjoint sets over elements.

- ▶ **P.** makeset(x): Given an element x, adds x to the data-structure and creates a singleton set that contains only this element. Returns a locator/handle for x in the data-structure.
- ▶ \mathcal{P} . find(x): Given a handle for an element x; find the set that contains x. Returns a representative/identifier for this set.
- ▶ \mathcal{P} . union(x, y): Given two elements x, and y that are currently in sets S_x and S_y , respectively, the function replaces S_x and S_y by $S_x \cup S_y$ and returns an identifier for the new set.

Applications:

► Keep track of the connected components of a dynamic graph that changes due to insertion of nodes and edges.

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- Keep track of the connected components of a dynamic graph that changes due to insertion of nodes and edges.
- Kruskals Minimum Spanning Tree Algorithm

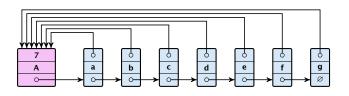
Algorithm 1 Kruskal-MST(G = (V, E), w)

- 1: $A \leftarrow \emptyset$;
- 2: for all $v \in V$ do
- 3: $v. set \leftarrow P. makeset(v. label)$
- 4: sort edges in non-decreasing order of weight w
- 5: **for all** $(u, v) \in E$ in non-decreasing order **do**
- 6: **if** \mathcal{P} . find(u. set) $\neq \mathcal{P}$. find(v. set) **then**
- 7: $A \leftarrow A \cup \{(u, v)\}$
- 8: $\mathcal{P}.union(u.set, v.set)$

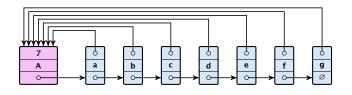
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► The elements of a set are stored in a list; each node has a backward pointer to the head.

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- ► The head of the list contains the identifier for the set and a field that stores the size of the set.

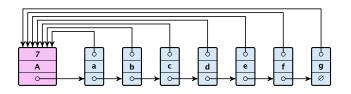


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- find(x) can be performed in constant time.

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union(x, y)

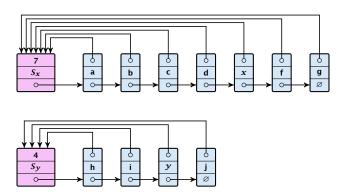
▶ Determine sets S_X and S_Y .

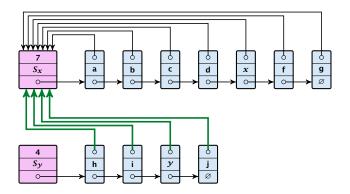
- ▶ Determine sets S_X and S_Y .
- ► Traverse the smaller list (say S_y), and change all backward pointers to the head of list S_x .

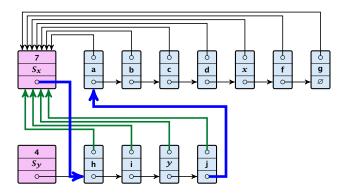
- ▶ Determine sets S_X and S_Y .
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- Adjust the size-field of list S_x .
- ► Time: $\min\{|S_x|, |S_y|\}$.

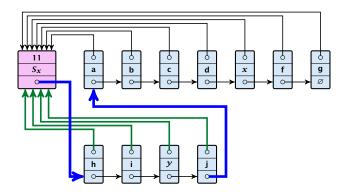








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Running times:

- ightharpoonup find(x): constant
- makeset(x): constant
- union(x, y): O(n), where n denotes the number of elements contained in the set system.

Lemma 34

The list implementation for the ADT union find fulfills the following amortized time bounds:

- find(x): $\mathcal{O}(1)$.
- ightharpoonup makeset(x): $O(\log n)$.
- ightharpoonup union(x, y): O(1).

The Accounting Method for Amortized Time Bounds

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- Whenever for an operation the actual cost exceeds the amortized time bound, the difference is charged to bank accounts of some of the elements involved.
- If we can find a charging scheme that guarantees that balances always stay positive the amortized time bounds are proven.

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- Later operations charge the account but the balance never drops below zero.

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- ▶ Charge c to every element in set S_x .



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An element is charged at most $\lfloor \log_2 n \rfloor$ times, where n is the total number of elements in the set system.

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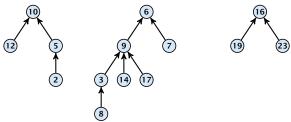
Proof.

Whenever an element x is charged the number of elements in x's set doubles. This can happen at most $|\log n|$ times.



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- Only pointer to parent exists; we cannot list all elements of a given set.

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- The root of the tree is the label of the set.
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- Example:



Set system {2,5,10,12}, {3,6,7,8,9,14,17}, {16,19,23}.

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Create a singleton tree. Return pointer to the root.

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find(x)

- Start at element x in the tree. Go upwards until you reach the root.
- ► Time: $\mathcal{O}(\text{level}(x))$, where level(x) is the distance of element x to the root in its tree. Not constant.

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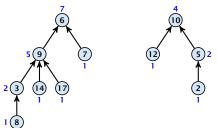
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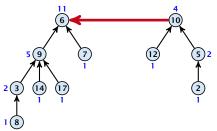


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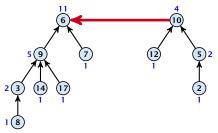


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▶ Time: constant for link(a, b) plus two find-operations.

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Path Compression

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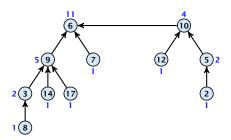
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- Re-attach all visited nodes as children of the root.

Path Compression

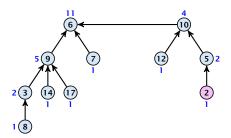
find(x):

- Go upward until you find the root.
- Re-attach all visited nodes as children of the root.
- Speeds up successive find-operations.

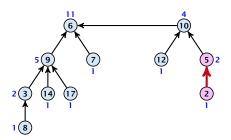
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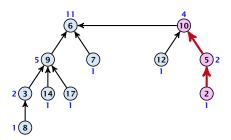
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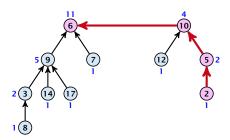
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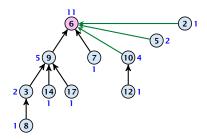
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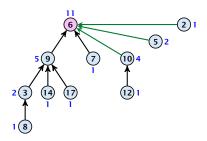


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Note that the size-fields now only give an upper bound on the size of a sub-tree.

9 Union Find

Asymptotically the cost for a find-operation does not increase due to the path compression heuristic.

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However, for a worst-case analysis there is no improvement on the running time. It can still happen that a find-operation takes time $\mathcal{O}(\log n)$.

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size(v) = the number of nodes that were in the sub-tree rooted at v when v became the child of another node (or the number of nodes if v is the root).

Note that this is the same as the size of ν 's subtree in the case that there are no find-operations.

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Lemma 37

The rank of a parent must be strictly larger than the rank of a child



Lemma 38

There are at most $n/2^s$ nodes of rank s.

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- A node v sees at most one node of rank s during the running time of the algorithm.

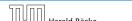


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- Let's say a node v sees node x if v is in x's sub-tree at the time that x becomes a child.
- A node v sees at most one node of rank s during the running time of the algorithm.
- This holds because the rank-sequence of the roots of the different trees that contain v during the running time of the algorithm is a strictly increasing sequence.
- Hence, every node *sees* at most one rank s node, but every rank s node is seen by at least 2^s different nodes.

We define

$$tow(i) := \left\{ \begin{array}{ll} 1 & \text{if } i = 0 \\ 2^{tow(i-1)} & \text{otw.} \end{array} \right.$$

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$$\log^*(n) := \min\{i \mid \text{tow}(i) \ge n\} .$$

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Theorem 39

Union find with path compression fulfills the following amortized running times:

- ightharpoonup makeset(x) : $\mathcal{O}(\log^*(n))$
- $ightharpoonup find(x) : \mathcal{O}(\log^*(n))$
- ightharpoonup union(x, y): $\mathcal{O}(\log^*(n))$

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416/616

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- ► The maximum non-empty rank group is $\log^*(\lfloor \log n \rfloor) \leq \log^*(n) 1$ (which holds for $n \geq 2$).
- ▶ Hence, the total number of rank-groups is at most $\log^* n$.

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Accounting Scheme:

create an account for every find-operation

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- If parent[v] is the root we charge the cost to the find-account.
- If the group-number of $\operatorname{rank}(v)$ is the same as that of $\operatorname{rank}(\operatorname{parent}[v])$ (before starting path compression) we charge the cost to the node-account of v.

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- create an account for every find-operation
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The cost for a find-operation is equal to the length of the path traversed. We charge the cost for going from v to parent[v] as follows:

- If parent[v] is the root we charge the cost to the find-account.
- If the group-number of rank(v) is the same as that of rank(parent[v]) (before starting path compression) we charge the cost to the node-account of v.
- Otherwise we charge the cost to the find-account.

Observations:

▶ A find-account is charged at most $\log^*(n)$ times (once for the root and at most $\log^*(n) - 1$ times when increasing the rank-group).

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- After a node v is charged its parent-edge is re-assigned. The rank of the parent strictly increases.
- After some charges to v the parent will be in a larger rank-group. $\Rightarrow v$ will never be charged again.
- ► The total charge made to a node in rank-group g is at most $tow(g) tow(g-1) 1 \le tow(g)$.

What is the total charge made to nodes?

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The total charge is at most

$$\sum_{g} n(g) \cdot \text{tow}(g) ,$$

where n(g) is the number of nodes in group g.

For $g \ge 1$ we have

n(g)

$$n(g) \le \sum_{s=\text{tow}(g-1)+1}^{\text{tow}(g)} \frac{n}{2^s}$$

$$n(g) \leq \sum_{s=\mathsf{tow}(g-1)+1}^{\mathsf{tow}(g)} \frac{n}{2^s} \leq \sum_{s=\mathsf{tow}(g-1)+1}^{\infty} \frac{n}{2^s}$$

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Hence,

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Hence,

$$\sum_{g} n(g) \operatorname{tow}(g) \le n(0) \operatorname{tow}(0) + \sum_{g>1} n(g) \operatorname{tow}(g) \le n \log^*(n)$$

Without loss of generality we can assume that all makeset-operations occur at the start.

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This means if we inflate the cost of makeset to $\log^* n$ and add this to the node account of v then the balances of all node accounts will sum up to a positive value (this is sufficient to obtain an amortized bound).



The analysis is not tight. In fact it has been shown that the amortized time for the union-find data structure with path compression is $\mathcal{O}(\alpha(m,n))$, where $\alpha(m,n)$ is the inverse Ackermann function which grows a lot lot slower than $\log^* n$. (Here, we consider the average running time of m operations on at most *n* elements).

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There is also a lower bound of $\Omega(\alpha(m, n))$.

$$A(x,y) = \begin{cases} y+1 & \text{if } x = 0 \\ A(x-1,1) & \text{if } y = 0 \\ A(x-1,A(x,y-1)) & \text{otw.} \end{cases}$$

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- A(0, v) = v + 1
- A(1, v) = v + 2
- $A(2, \nu) = 2\nu + 3$
- ► $A(3, y) = 2^{y+3} 3$ ► $A(4, y) = 2^{2^{2^2}} 3$

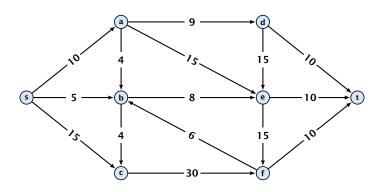
Part IV

Flows and Cuts

The following slides are partially based on slides by Kevin Wayne.

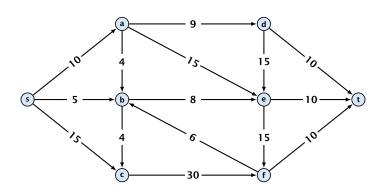
Flow Network

• directed graph G = (V, E); edge capacities c(e)



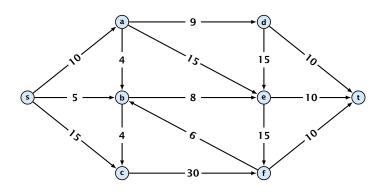
Flow Network

- directed graph G = (V, E); edge capacities c(e)
- two special nodes: source s; target t;



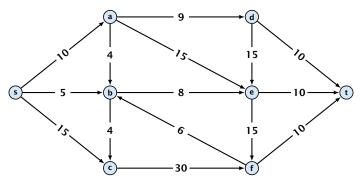
Flow Network

- directed graph G = (V, E); edge capacities c(e)
- two special nodes: source s; target t;
- ightharpoonup no edges entering s or leaving t;



Flow Network

- directed graph G = (V, E); edge capacities c(e)
- two special nodes: source s; target t;
- no edges entering s or leaving t;
- at least for now: no parallel edges;



Definition 40

An (s, t)-cut in the graph G is given by a set $A \subset V$ with $s \in A$ and $t \in V \setminus A$.

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Definition 41

The capacity of a cut A is defined as

$$cap(A, V \setminus A) := \sum_{e \in out(A)} c(e) ,$$

where $\operatorname{out}(A)$ denotes the set of edges of the form $A \times V \setminus A$ (i.e. edges leaving A).

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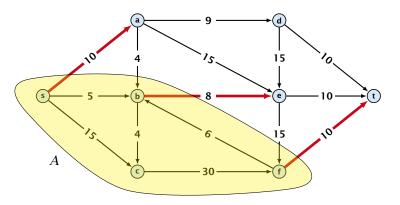
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Minimum Cut Problem: Find an (s, t)-cut with minimum capacity.

Example 42



The capacity of the cut is $cap(A, V \setminus A) = 28$.



Flows

Definition 43

An (s,t)-flow is a function $f: E \rightarrow \mathbb{R}^+$ that satisfies

1. For each edge e

$$0 \le f(e) \le c(e)$$
.

(capacity constraints)

Flows

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1. For each edge e

$$0 \le f(e) \le c(e)$$
.

(capacity constraints)

2. For each $v \in V \setminus \{s, t\}$

$$\sum_{e \in \text{out}(v)} f(e) = \sum_{e \in \text{into}(v)} f(e) .$$

(flow conservation constraints)

Definition 44

The value of an (s, t)-flow f is defined as

$$val(f) = \sum_{e \in out(s)} f(e)$$
.

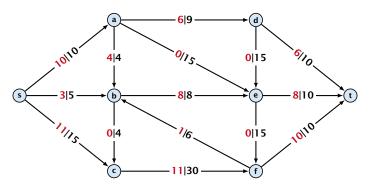
Definition 44

The value of an (s, t)-flow f is defined as

$$\operatorname{val}(f) = \sum_{e \in \operatorname{out}(s)} f(e)$$
.

Maximum Flow Problem: Find an (s, t)-flow with maximum value.

Example 45



The value of the flow is val(f) = 24.

Lemma 46 (Flow value lemma)

Let f be a flow, and let $A \subseteq V$ be an (s,t)-cut. Then the net-flow across the cut is equal to the amount of flow leaving s, i.e.,

$$\operatorname{val}(f) = \sum_{e \in \operatorname{out}(A)} f(e) - \sum_{e \in \operatorname{into}(A)} f(e)$$
.

val(f)

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$$= \sum_{e \in out(A)} f(e) - \sum_{e \in into(A)} f(e)$$

$$val(f) = \sum_{e \in out(s)} f(e)$$

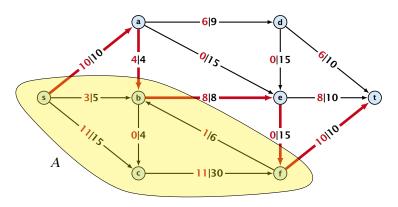
$$= \sum_{e \in out(s)} f(e) + \sum_{v \in A \setminus \{s\}} \left(\sum_{e \in out(v)} f(e) - \sum_{e \in in(v)} f(e) \right)$$

$$= \sum_{e \in out(A)} f(e) - \sum_{e \in into(A)} f(e)$$

The last equality holds since every edge with both end-points in A contributes negatively as well as positively to the sum in Line 2. The only edges whose contribution doesn't cancel out are edges leaving or entering A.

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Example 47



The net-flow across the cut is val(f) = 24.

10 Introduction

Let f be an (s,t)-flow and let A be an (s,t)-cut, such that

$$\operatorname{val}(f) = \operatorname{cap}(A, V \setminus A).$$

Then f is a maximum flow.

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Proof.

Let f be an (s,t)-flow and let A be an (s,t)-cut, such that

$$\operatorname{val}(f) = \operatorname{cap}(A, V \setminus A).$$

Then f is a maximum flow.

Proof.

$$cap(A, V \setminus A) < val(f')$$

Let f be an (s,t)-flow and let A be an (s,t)-cut, such that

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Proof.

$$cap(A, V \setminus A) < val(f')$$

$$= \sum_{e \in out(A)} f'(e) - \sum_{e \in into(A)} f'(e)$$

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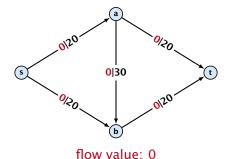
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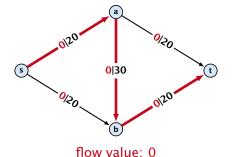
$$\leq \sum_{e \in out(A)} f'(e)$$

$$\leq cap(A, V \setminus A)$$

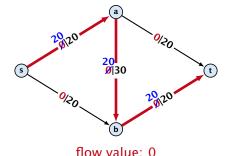
- start with f(e) = 0 everywhere
- find an s-t path with f(e) < c(e) on every edge
- augment flow along the path
- repeat as long as possible



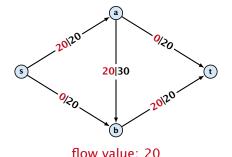
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From the graph G = (V, E, c) and the current flow f we construct an auxiliary graph $G_f = (V, E_f, c_f)$ (the residual graph):

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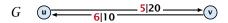
▶ Suppose the original graph has edges $e_1 = (u, v)$, and $e_2 = (v, u)$ between u and v.

From the graph G = (V, E, c) and the current flow f we construct an auxiliary graph $G_f = (V, E_f, c_f)$ (the residual graph):

- Suppose the original graph has edges $e_1 = (u, v)$, and $e_2 = (v, u)$ between u and v.
- ▶ G_f has edge e_1' with capacity $\max\{0, c(e_1) f(e_1) + f(e_2)\}$ and e_2' with with capacity $\max\{0, c(e_2) f(e_2) + f(e_1)\}$.

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Definition 49

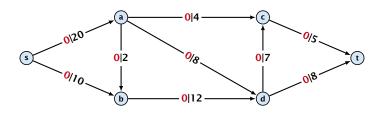
An augmenting path with respect to flow f, is a path from s to t in the auxiliary graph G_f that contains only edges with non-zero capacity.

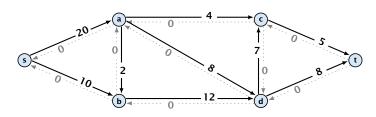
Definition 49

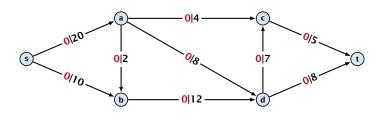
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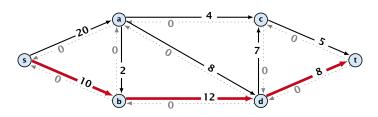
Algorithm 1 FordFulkerson(G = (V, E, c))

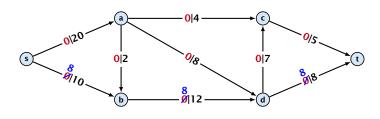
- 1: Initialize $f(e) \leftarrow 0$ for all edges.
- 2: **while** \exists augmenting path p in G_f **do**
- 3: augment as much flow along p as possible.

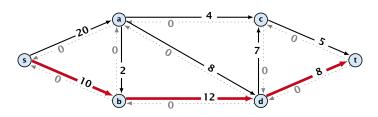


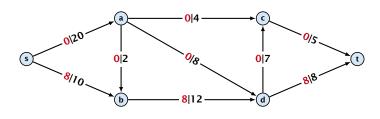


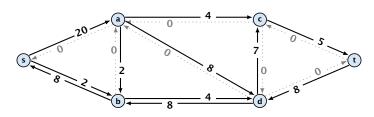


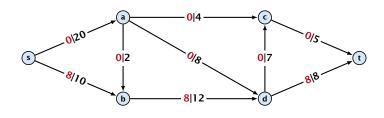


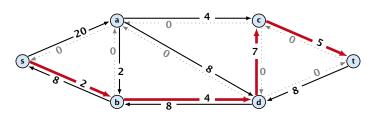


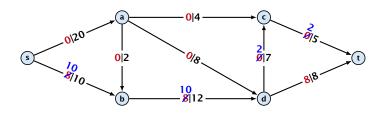


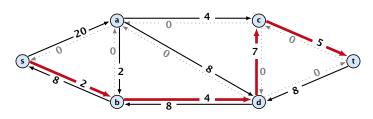


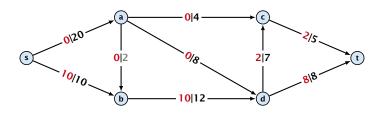


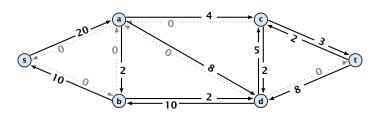


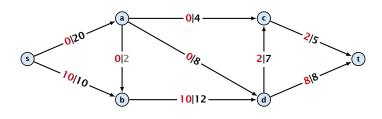


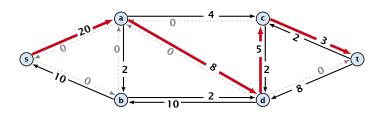




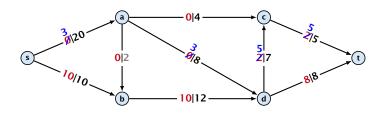




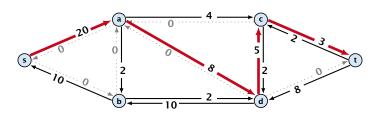




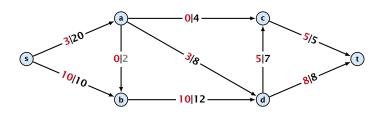
Augmenting Paths



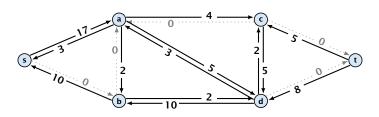
flow value: 10



Augmenting Paths



flow value: 13



Theorem 50

A flow f is a maximum flow **iff** there are no augmenting paths.

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Theorem 51

The value of a maximum flow is equal to the value of a minimum cut.

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Proof.

Let f be a flow. The following are equivalent:

1. There exists a cut A such that $val(f) = cap(A, V \setminus A)$.



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Proof.

Let f be a flow. The following are equivalent:

- **1.** There exists a cut A such that $val(f) = cap(A, V \setminus A)$.
- 2. Flow f is a maximum flow.



Theorem 50

A flow f is a maximum flow **iff** there are no augmenting paths.

Theorem 51

The value of a maximum flow is equal to the value of a minimum cut.

Proof.

Let f be a flow. The following are equivalent:

- **1.** There exists a cut A such that $val(f) = cap(A, V \setminus A)$.
- 2. Flow f is a maximum flow.
- 3. There is no augmenting path w.r.t. f.



 $1. \Rightarrow 2.$

This we already showed.

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 $2. \Rightarrow 3.$

If there were an augmenting path, we could improve the flow. Contradiction.

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If there were an augmenting path, we could improve the flow. Contradiction.

- $3. \Rightarrow 1.$
 - Let f be a flow with no augmenting paths.

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This we already showed.

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If there were an augmenting path, we could improve the flow. Contradiction.

- $3. \Rightarrow 1.$
 - Let f be a flow with no augmenting paths.
 - Let A be the set of vertices reachable from s in the residual graph along non-zero capacity edges.

$$1. \Rightarrow 2.$$

This we already showed.

$$2. \Rightarrow 3.$$

If there were an augmenting path, we could improve the flow. Contradiction.

- $3. \Rightarrow 1.$
 - Let f be a flow with no augmenting paths.
 - Let A be the set of vertices reachable from s in the residual graph along non-zero capacity edges.
 - ▶ Since there is no augmenting path we have $s \in A$ and $t \notin A$.

val(f)

$$\operatorname{val}(f) = \sum_{e \in \operatorname{out}(A)} f(e) - \sum_{e \in \operatorname{into}(A)} f(e)$$

$$val(f) = \sum_{e \in out(A)} f(e) - \sum_{e \in into(A)} f(e)$$
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$$= cap(A, V \setminus A)$$

This finishes the proof.

Here the first equality uses the flow value lemma, and the second exploits the fact that the flow along incoming edges must be 0 as the residual graph does not have edges leaving A.

Analysis

Assumption:

All capacities are integers between 1 and ${\cal C}$.

Analysis

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All capacities are integers between 1 and C.

Invariant:

Every flow value f(e) and every residual capacity $c_f(e)$ remains integral troughout the algorithm.

Lemma 52

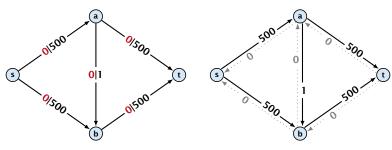
The algorithm terminates in at most $val(f^*) \le nC$ iterations, where f^* denotes the maximum flow. Each iteration can be implemented in time $\mathcal{O}(m)$. This gives a total running time of $\mathcal{O}(nmC)$.

Lemma 52

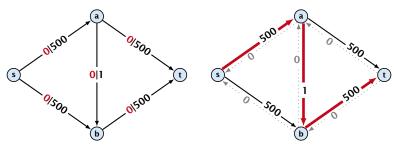
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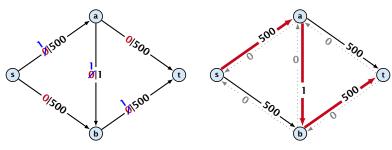
Theorem 53

If all capacities are integers, then there exists a maximum flow for which every flow value f(e) is integral.

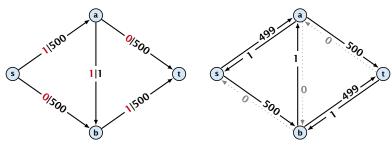


flow value: 0

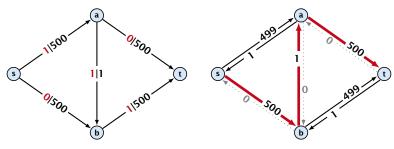


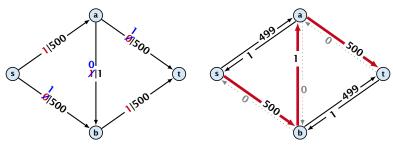


flow value: 0

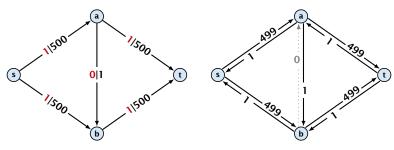


flow value: 1

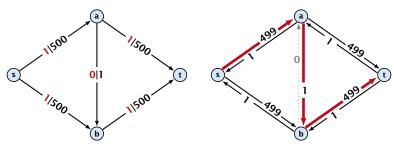




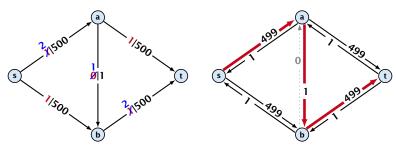
flow value: 1



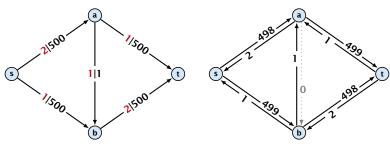
flow value: 2



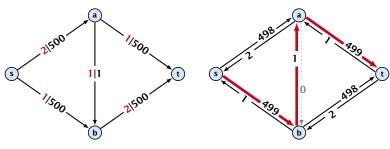
flow value: 2



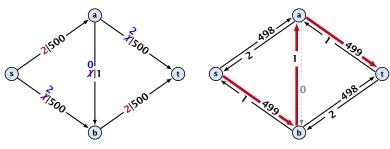
flow value: 2



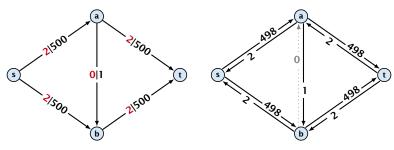
flow value: 3



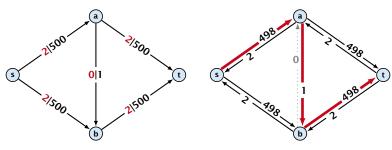
flow value: 3



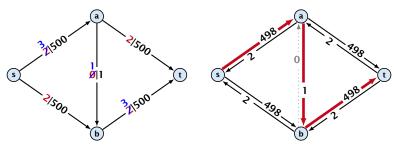
flow value: 3



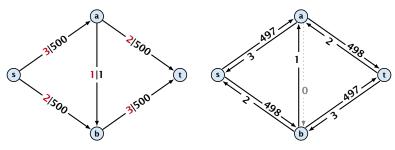
flow value: 4



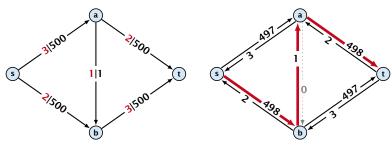
flow value: 4



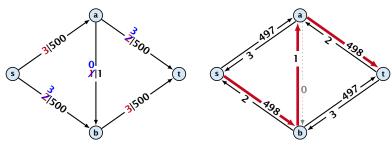
flow value: 4



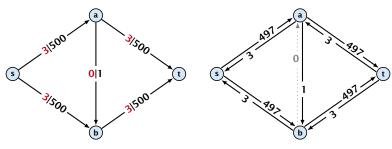
flow value: 5



flow value: 5

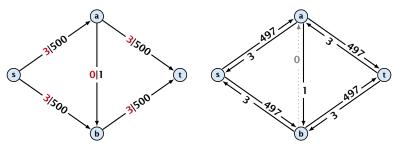


flow value: 5



flow value: 6

Problem: The running time may not be polynomial

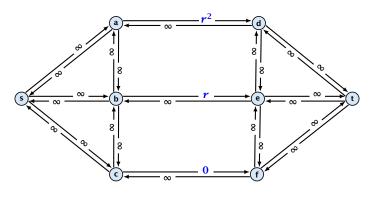


flow value: 6

Question:

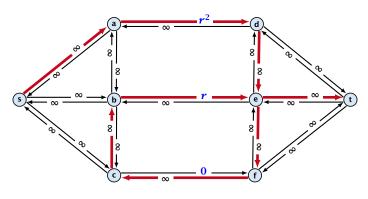
Can we tweak the algorithm so that the running time is polynomial in the input length?

Let
$$r = \frac{1}{2}(\sqrt{5} - 1)$$
. Then $r^{n+2} = r^n - r^{n+1}$.



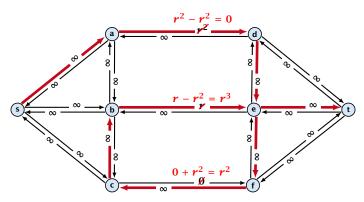
flow value: 0

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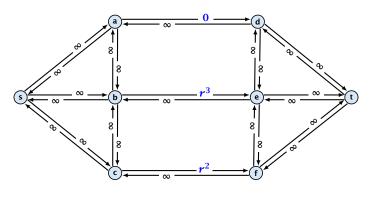
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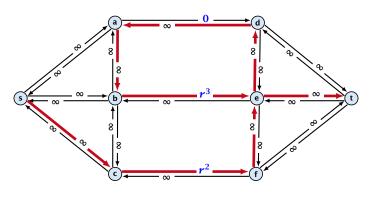
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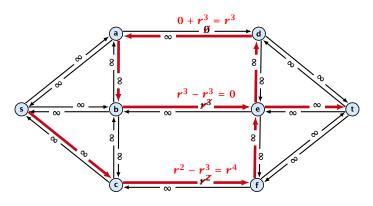
flow value: r^2

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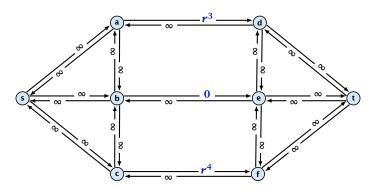
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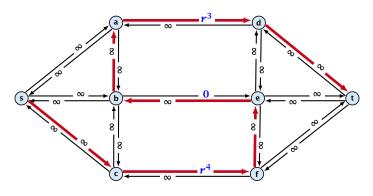
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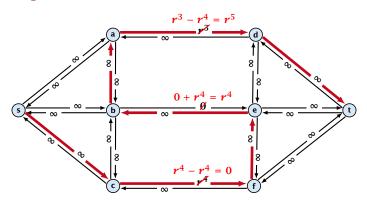
flow value: $r^2 + r^3$

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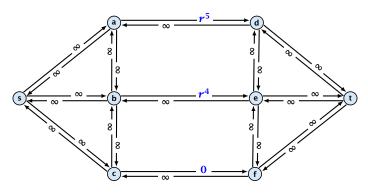
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flow value: $r^2 + r^3 + r^4$

Running time may be infinite!!!

We need to find paths efficiently.

- We need to find paths efficiently.
- We want to guarantee a small number of iterations.

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Several possibilities:

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Choose path with maximum bottleneck capacity.

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Several possibilities:

- Choose path with maximum bottleneck capacity.
- Choose path with sufficiently large bottleneck capacity.
- Choose the shortest augmenting path.

Lemma 54

The length of the shortest augmenting path never decreases.

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Lemma 55

After at most $\mathcal{O}(m)$ augmentations, the length of the shortest augmenting path strictly increases.

These two lemmas give the following theorem:

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Theorem 56

The shortest augmenting path algorithm performs at most $\mathcal{O}(mn)$ augmentations. This gives a running time of $\mathcal{O}(m^2n)$.

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Proof.

We can find the shortest augmenting paths in time $\mathcal{O}(m)$ via BFS.



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Theorem 56

The shortest augmenting path algorithm performs at most $\mathcal{O}(mn)$ augmentations. This gives a running time of $\mathcal{O}(m^2n)$.

Proof.

- We can find the shortest augmenting paths in time $\mathcal{O}(m)$ via BFS.
- O(m) augmentations for paths of exactly k < n edges.



Define the level $\ell(v)$ of a node as the length of the shortest s-v path in G_f (along non-zero edges).

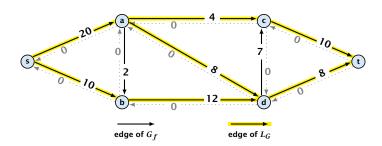
Define the level $\ell(v)$ of a node as the length of the shortest s-v path in G_f (along non-zero edges).

Let L_G denote the subgraph of the residual graph G_f that contains only those edges (u, v) with $\ell(v) = \ell(u) + 1$.

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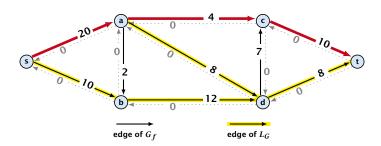
A path P is a shortest s-u path in G_f iff it is an s-u path in L_G .



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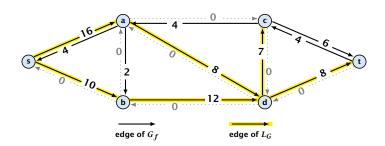
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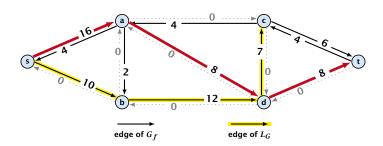
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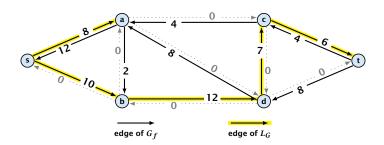
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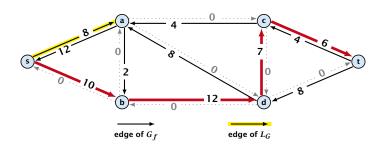
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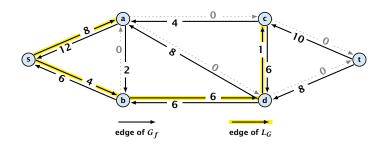
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In the following we assume that the residual graph \mathcal{G}_f does not contain zero capacity edges.

This means, we construct it in the usual sense and then delete edges of zero capacity.

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The length of the shortest augmenting path never decreases.

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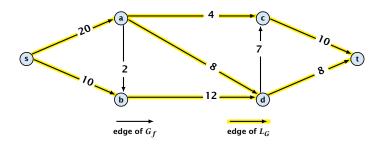
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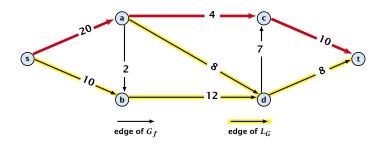


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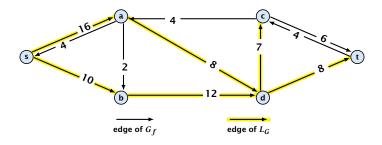


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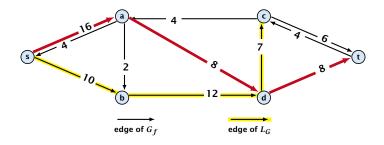


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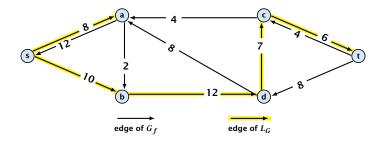


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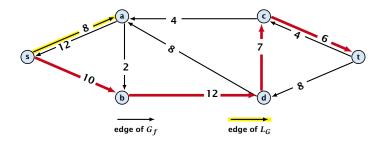


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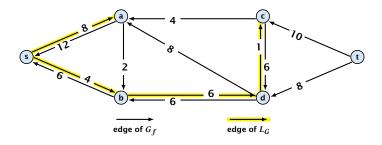


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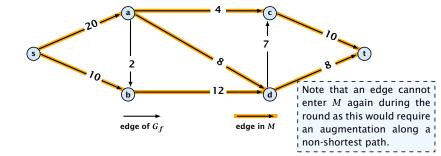
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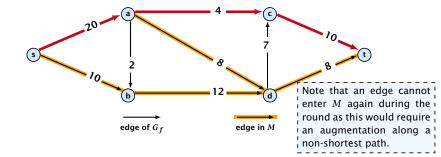
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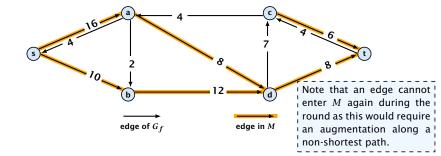
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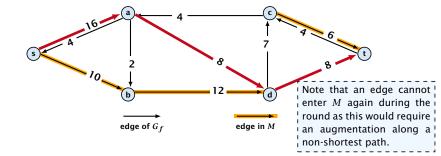
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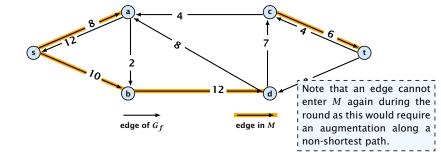
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Note:

There always exists a set of m augmentations that gives a maximum flow (why?).

When sticking to shortest augmenting paths we cannot improve (asymptotically) on the number of augmentations.

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However, we can improve the running time to $\mathcal{O}(mn^2)$ by improving the running time for finding an augmenting path (currently we assume $\mathcal{O}(m)$ per augmentation for this).

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Note that ${\cal M}$ is not the set of edges of the level graph but a subset of level-graph edges.

Suppose that the initial distance between s and t in G_f is k.

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There are at most n phases. Hence, total cost is $O(mn^2)$.

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- Choose path with maximum bottleneck capacity.
- Choose path with sufficiently large bottleneck capacity.
- Choose the shortest augmenting path.



Intuition:

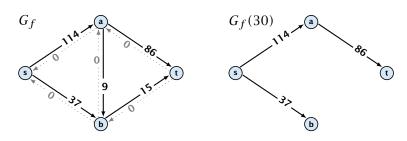
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```
Algorithm 1 maxflow(G, s, t, c)
 1: foreach e \in E do f_e \leftarrow 0;
 2: \Delta \leftarrow 2^{\lceil \log_2 C \rceil}
 3: while \Delta \geq 1 do
 4: G_f(\Delta) \leftarrow \Delta-residual graph
5: while there is augmenting path P in G_f(\Delta) do
6: f \leftarrow \text{augment}(f, c, P)
7: \text{update}(G_f(\Delta))
8: \Delta \leftarrow \Delta/2
 9: return f
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- ▶ In G_f this cut can have capacity at most $m\Delta$.
- This gives me an upper bound on the flow that I can still add.



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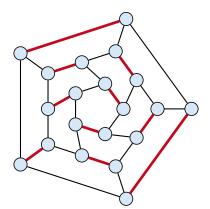
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Theorem 62

We need $\mathcal{O}(m \log C)$ augmentations. The algorithm can be implemented in time $\mathcal{O}(m^2 \log C)$.

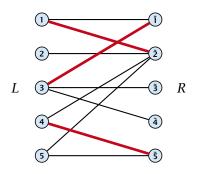
Matching

- ▶ Input: undirected graph G = (V, E).
- ► $M \subseteq E$ is a matching if each node appears in at most one edge in M.
- Maximum Matching: find a matching of maximum cardinality



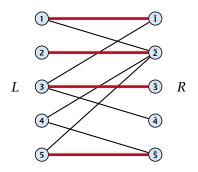
Bipartite Matching

- ▶ Input: undirected, bipartite graph $G = (L \uplus R, E)$.
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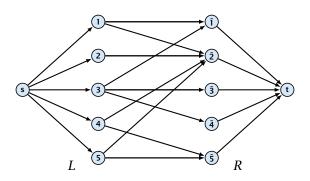
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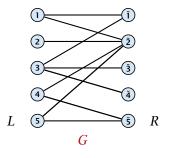
Maxflow Formulation

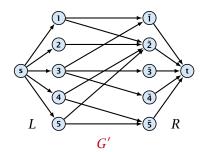
- ▶ Input: undirected, bipartite graph $G = (L \uplus R \uplus \{s, t\}, E')$.
- Direct all edges from L to R.
- Add source s and connect it to all nodes on the left.
- Add t and connect all nodes on the right to t.
- All edges have unit capacity.



Max cardinality matching in $G \le \text{value of maxflow in } G'$

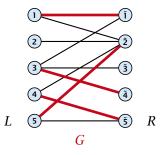
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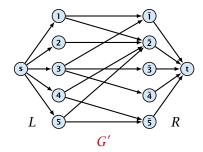




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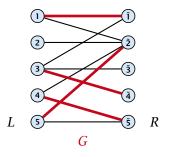


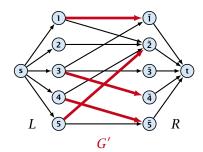


12.1 Matching 6. Feb. 202

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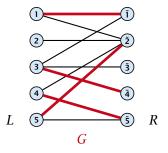
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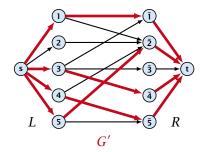




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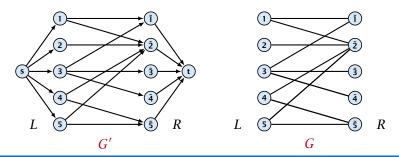
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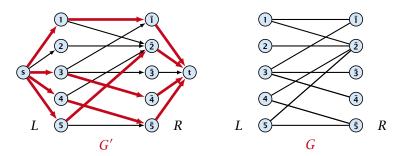
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- Consider M= set of edges from L to R with f(e) = 1.
- \blacktriangleright Each node in L and R participates in at most one edge in M.
- |M| = k, as the flow must use at least k middle edges.



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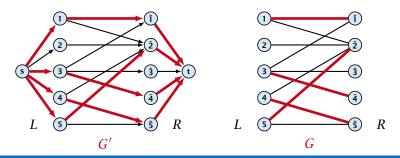
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12.1 Matching

Which flow algorithm to use?

- Generic augmenting path: $\mathcal{O}(m \operatorname{val}(f^*)) = \mathcal{O}(mn)$.
- Capacity scaling: $\mathcal{O}(m^2 \log C) = \mathcal{O}(m^2)$.
- Shortest augmenting path: $O(mn^2)$.

For unit capacity simple graphs shortest augmenting path can be implemented in time $\mathcal{O}(m\sqrt{n})$.

A graph is a unit capacity simple graph if

- every edge has capacity 1
- a node has either at most one leaving edge or at most one entering edge



team	wins	losses	remaining games			
i	w_i	ℓ_i	Atl	Phi	NY	Mon
Atlanta	83	71	_	1	6	1
Philadelphia	80	79	1	_	0	2
New York	78	78	6	0	_	0
Montreal	77	82	1	2	0	_

Which team can end the season with most wins?

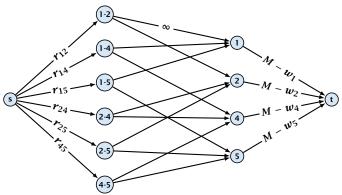
- Montreal is eliminated, since even after winning all remaining games there are only 80 wins.
- But also Philadelphia is eliminated. Why?



Formal definition of the problem:

- ▶ Given a set S of teams, and one specific team $z \in S$.
- ▶ Team x has already won w_x games.
- ► Team x still has to play team y, r_{xy} times.
- Does team z still have a chance to finish with the most number of wins.

Flow network for z = 3. M is number of wins Team 3 can still obtain.



Idea. Distribute the results of remaining games in such a way that no team gets too many wins.

Certificate of Elimination

Let $T \subseteq S$ be a subset of teams. Define

$$w(T) := \sum_{i \in T} w_i, \qquad r(T) := \sum_{i,j \in T, i < j} r_{ij}$$
 wins of teams in T remaining games among teams in T

If $\frac{w(T)+r(T)}{|T|}>M$ then one of the teams in T will have more than M wins in the end. A team that can win at most M games is therefore eliminated.

A team z is eliminated if and only if the flow network for z does not allow a flow of value $\sum_{i,j \in S \setminus \{z\}, i < j} \gamma_{i,j}$.

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► This gives M < (w(T) + r(T))/|T|, i.e., z is eliminated.

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- Hence, we found a set of results for the remaining games, such that no team obtains more than M wins in total.
- Hence, team z is not eliminated.

Project Selection

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Set P of possible projects. Project v has an associated profit p_v (can be positive or negative).

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- ▶ A subset *A* of projects is feasible if the prerequisites of every project in *A* also belong to *A*.

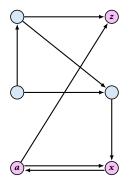
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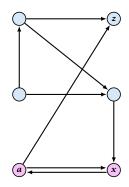
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- Dependencies are modelled in a graph. Edge (u, v) means "can't do project u without also doing project v."
- ▶ A subset *A* of projects is feasible if the prerequisites of every project in *A* also belong to *A*.

Goal: Find a feasible set of projects that maximizes the profit.

The prerequisite graph:

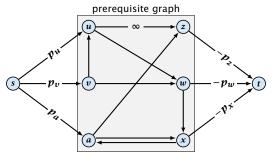
- \blacktriangleright {x, a, z} is a feasible subset.
- \triangleright {x, a} is infeasible.





Mincut formulation:

- Edges in the prerequisite graph get infinite capacity.
- Add edge (s, v) with capacity p_v for nodes v with positive profit.
- ► Create edge (v,t) with capacity $-p_v$ for nodes v with negative profit.



A is a mincut if $A \setminus \{s\}$ is the optimal set of projects.

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Proof.

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For the formula we define $p_s := 0$.

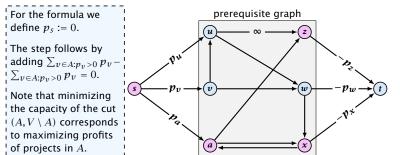
The step follows by adding $\sum_{v \in A: p_v > 0} p_v - \sum_{v \in A: p_v > 0} p_v = 0$.

Note that minimizing the capacity of the cut $(A, V \setminus A)$ corresponds to maximizing profits of projects in A.

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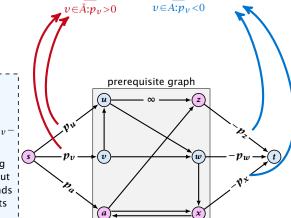
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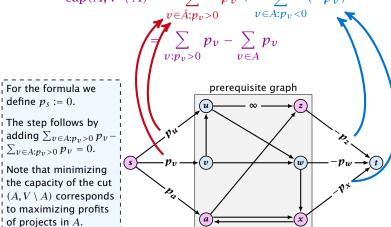
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Definition 65

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1. For each edge e

$$0 \leq f(e) \leq c(e) \ .$$

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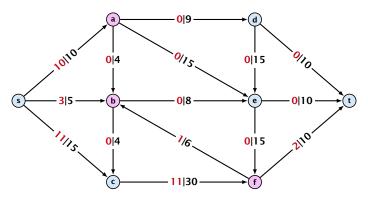
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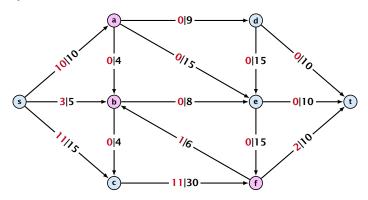
2. For each $v \in V \setminus \{s, t\}$

$$\sum_{e \in \text{out}(v)} f(e) \le \sum_{e \in \text{into}(v)} f(e) .$$

Example 66



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A node that has $\sum_{e \in \text{out}(v)} f(e) < \sum_{e \in \text{into}(v)} f(e)$ is called an active node.

Definition:

A labelling is a function $\ell: V \to \mathbb{N}$. It is valid for preflow f if

• $\ell(u) \le \ell(v) + 1$ for all edges (u, v) in the residual graph G_f (only non-zero capacity edges!!!)

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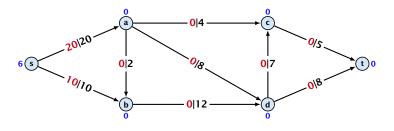
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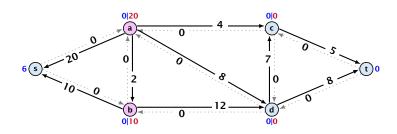
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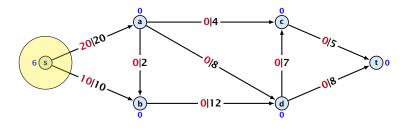
Intuition:

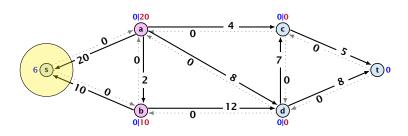
The labelling can be viewed as a height function. Whenever the height from node u to node v decreases by more than 1 (i.e., it goes very steep downhill from u to v), the corresponding edge must be saturated.

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Lemma 68

A flow that has a valid labelling is a maximum flow.



Idea:

start with some preflow and some valid labelling

Note that this is somewhat dual to an augmenting path algorithm. The former maintains the property that it has a feasible flow. It successively changes this flow until it saturates some cut in which case we conclude that the flow is maximum. A preflow push algorithm maintains the property that it has a saturated cut. The preflow is changed iteratively until it fulfills conservation constraints in which case we can conclude that we have a maximum flow.

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- start with some preflow and some valid labelling
- successively change the preflow while maintaining a valid labelling
- stop when you have a flow (i.e., no more active nodes)

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An arc (u,v) with $c_f(u,v)>0$ in the residual graph is admissible if $\ell(u)=\ell(v)+1$ (i.e., it goes downwards w.r.t. labelling ℓ).

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The push operation

Consider an active node u with excess flow

 $f(u) = \sum_{e \in \text{into}(u)} f(e) - \sum_{e \in \text{out}(u)} f(e)$ and suppose e = (u, v) is an admissible arc with residual capacity $c_f(e)$.

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We can send flow $\min\{c_f(e), f(u)\}$ along e and obtain a new preflow. The old labelling is still valid (!!!).

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- ▶ deactivating push: $min{f(u), c_f(e)} = f(u)$ the node u becomes inactive

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Increasing the label of u by 1 results in a valid labelling.

- ▶ Edges (w, u) incoming to u still fulfill their constraint $\ell(w) \le \ell(u) + 1$.
- An outgoing edge (u, w) had $\ell(u) < \ell(w) + 1$ before since it was not admissible. Now: $\ell(u) \le \ell(w) + 1$.

Intuition:

We want to send flow downwards, since the source has a height/label of n and the target a height/label of 0. If we see an active node u with an admissible arc we push the flow at u towards the other end-point that has a lower height/label. If we do not have an admissible arc but excess flow into u it should roughly mean that the level/height/label of u should rise. (If we consider the flow to be water then this would be natural.)

Note that the above intuition is very incorrect as the labels are integral, i.e., they cannot really be seen as the height of a node.

Reminder

- In a preflow nodes may not fulfill conservation constraints; a node may have more incoming flow than outgoing flow.
- Such a node is called active.
- ▶ A labelling is valid if for every edge (u, v) in the residual graph $\ell(u) \le \ell(v) + 1$.
- An arc (u, v) in residual graph is admissible if $\ell(u) = \ell(v) + 1$.
- A saturating push along e pushes an amount of c(e) flow along the edge, thereby saturating the edge (and making it dissappear from the residual graph).
- A deactivating push along e = (u, v) pushes a flow of f(u), where f(u) is the excess flow of u. This makes u inactive.

```
Algorithm 1 maxflow(G, s, t, c)

1: find initial preflow f

2: while there is active node u do

3: if there is admiss. arc e out of u then

4: push(G, e, f, c)

5: else

6: relabel(u)

7: return f
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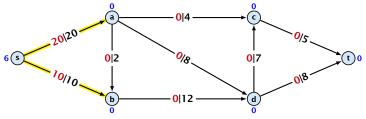
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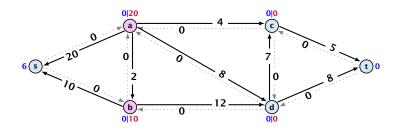
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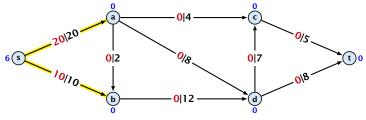
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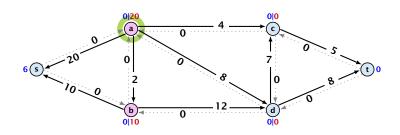
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In the following example we always stick to the same active node \boldsymbol{u} until it becomes inactive but this is not required.

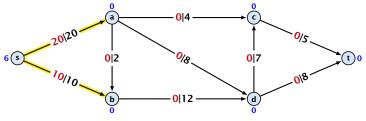




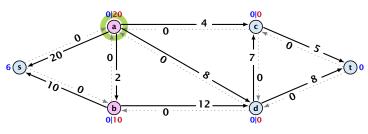


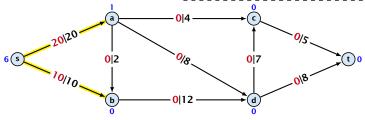


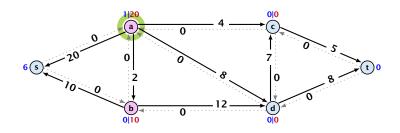
The yellow edges indicate the cut that is introduced by the smallest missing label.



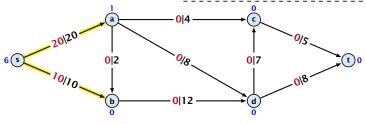
relabel to 1



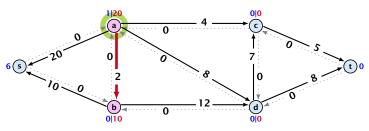


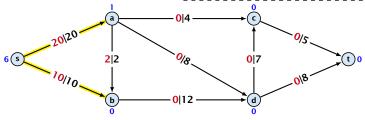


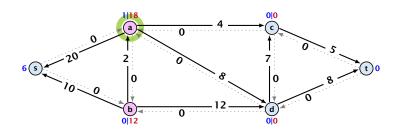
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saturating push

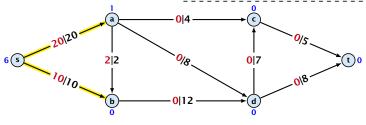




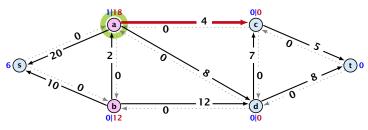


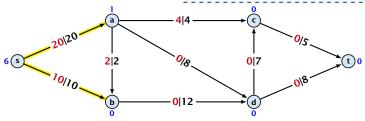


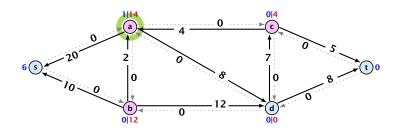
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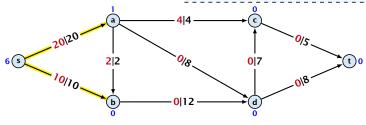
saturating push



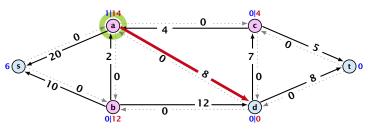


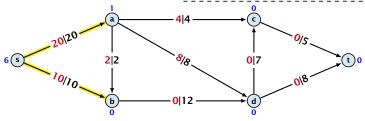


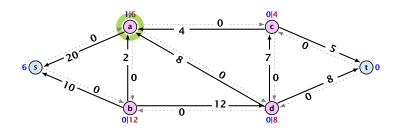
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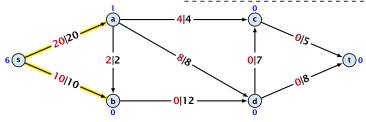
saturating push



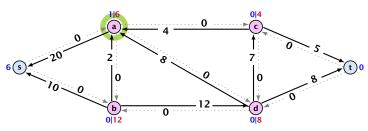


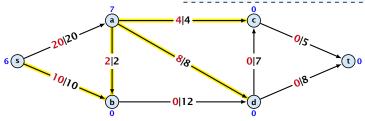


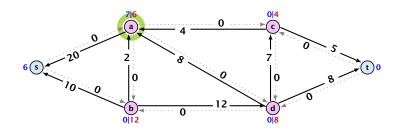
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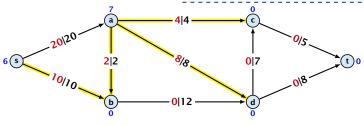
relabel to 7



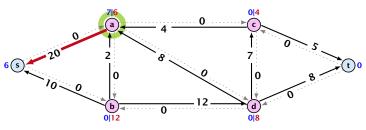


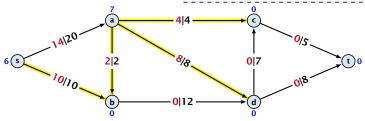


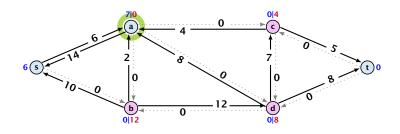
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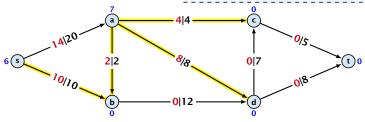


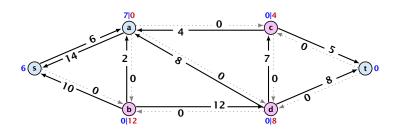
deactivating push

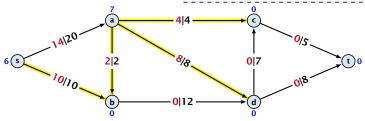


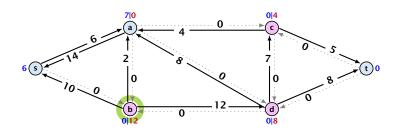






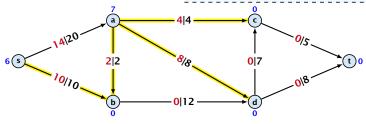




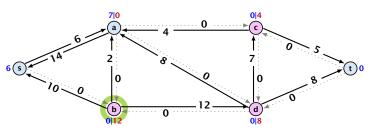


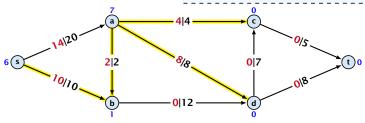


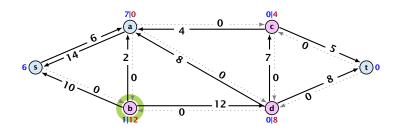
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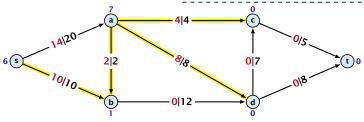
relabel to 1



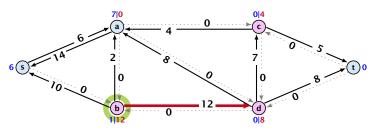


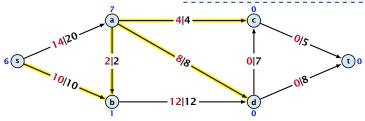


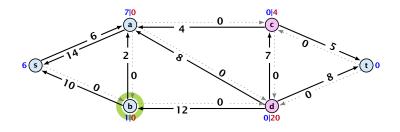
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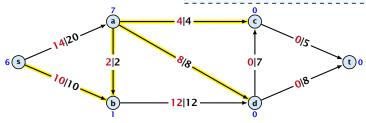


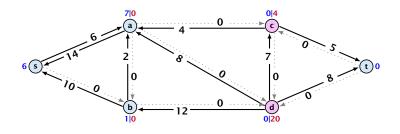
saturating and deactivating push

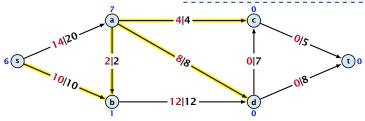


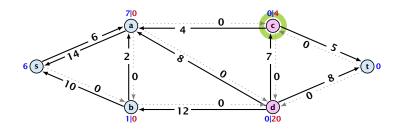






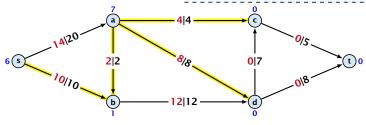




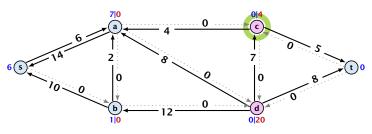


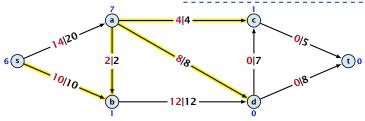


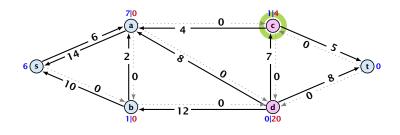
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relabel to 1

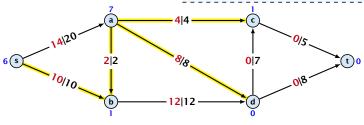




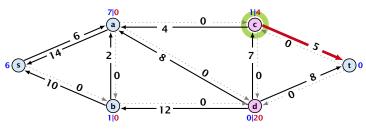




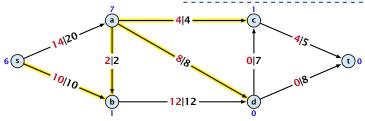
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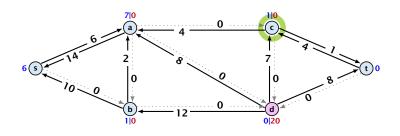


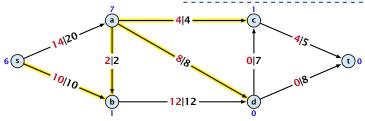
deactivating push

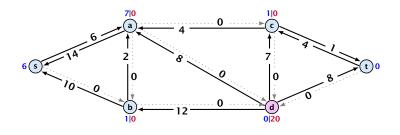


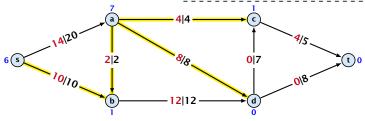


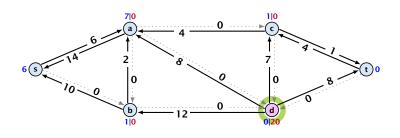




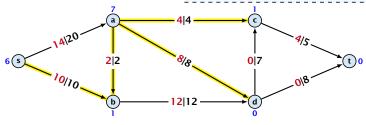




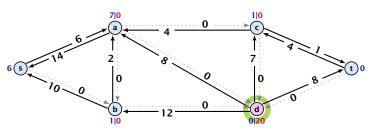


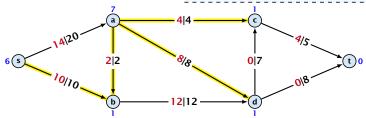


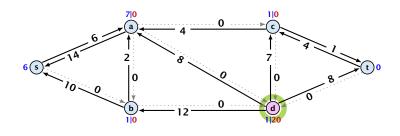
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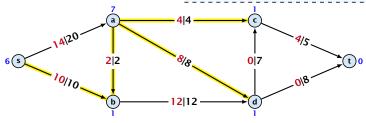
relabel to 1



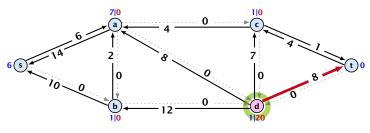


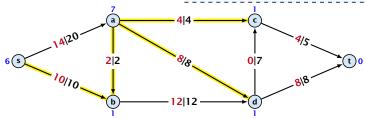


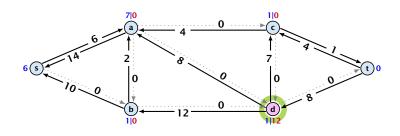
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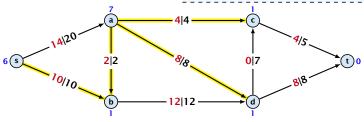
saturating push



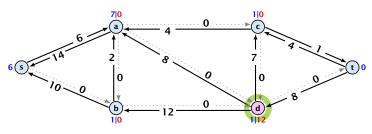


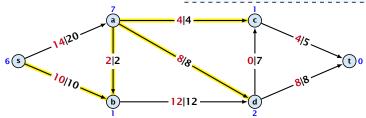


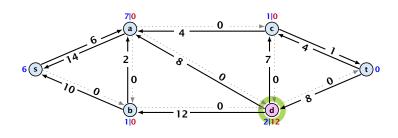
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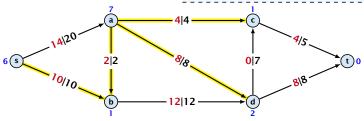
relabel to 2



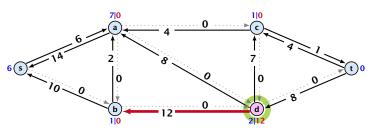


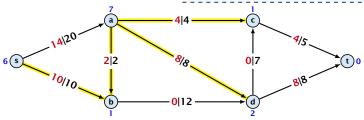


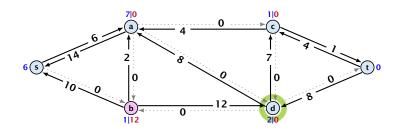
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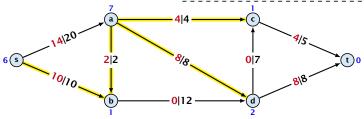


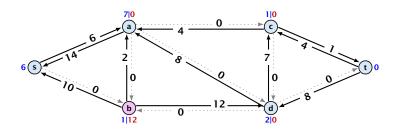
saturating and deactivating push

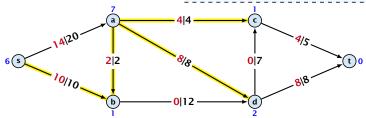


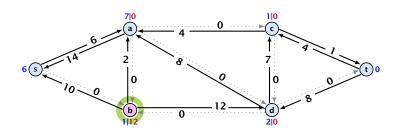




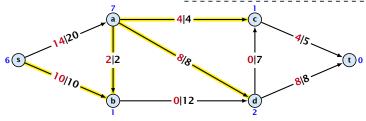




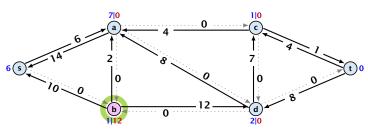


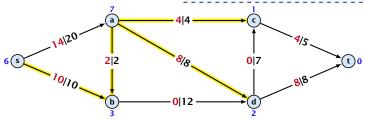


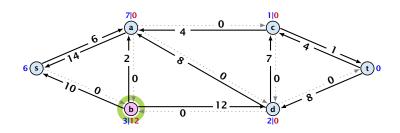
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relabel to 3

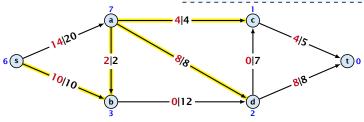




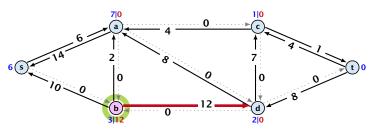


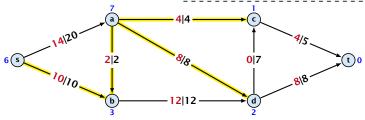


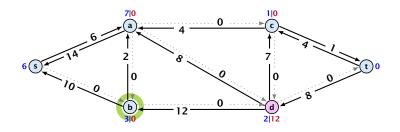
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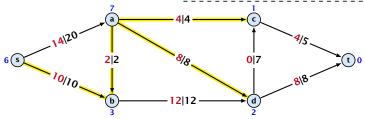


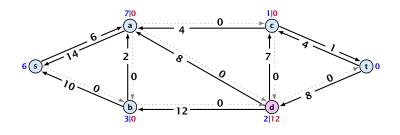
saturating and deactivating push

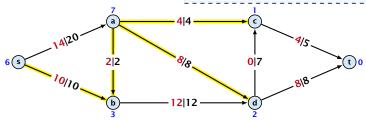


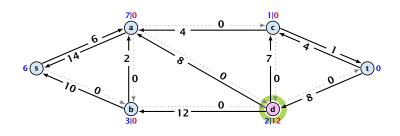




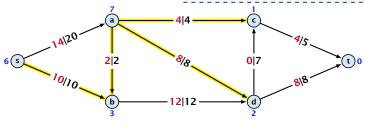




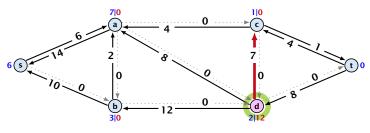


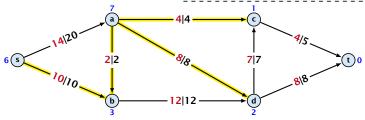


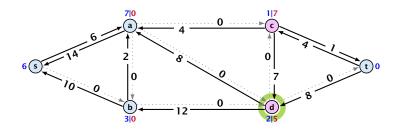
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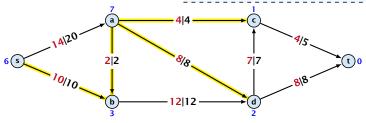
saturating push



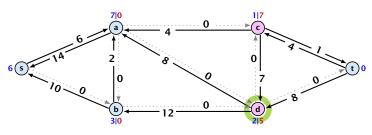


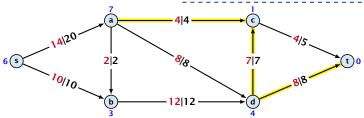


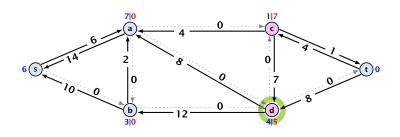
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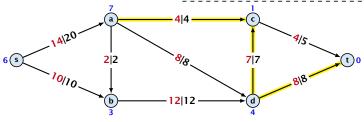
relabel to 4



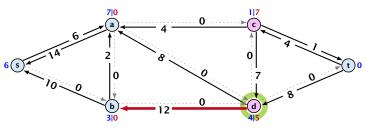


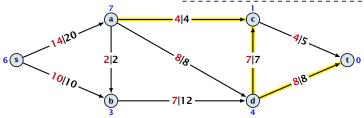


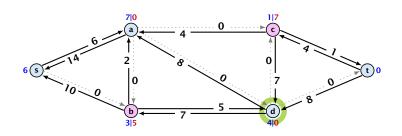
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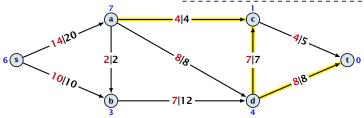


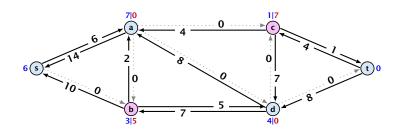
deactivating push

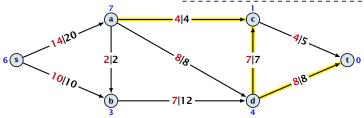


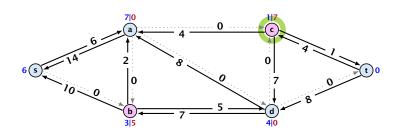




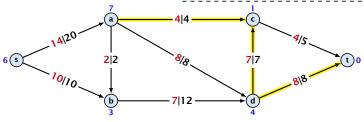




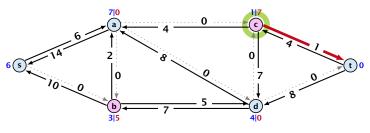


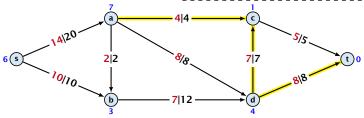


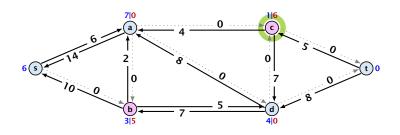
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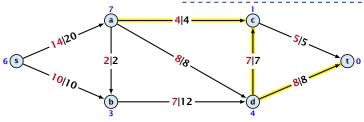
saturating push



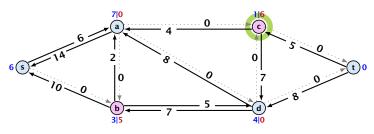


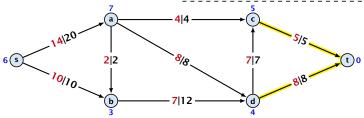


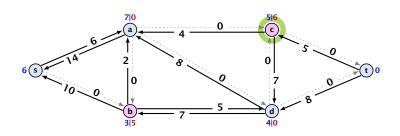
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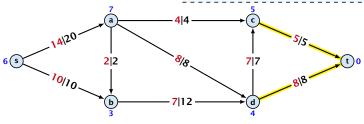
relabel to 5



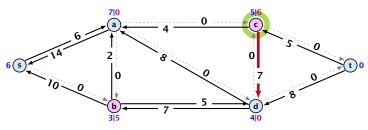


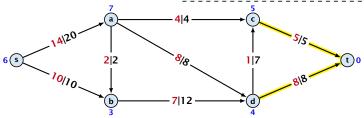


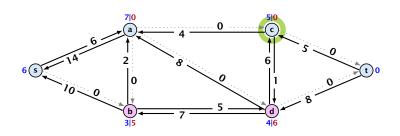
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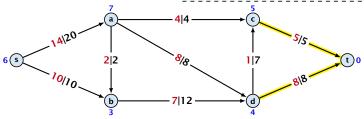


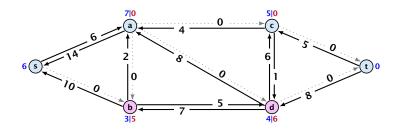
deactivating push

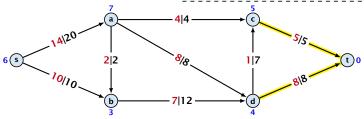


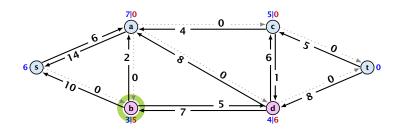




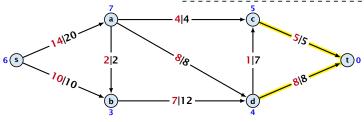




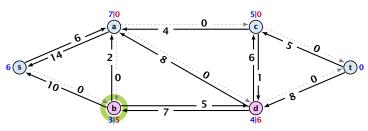


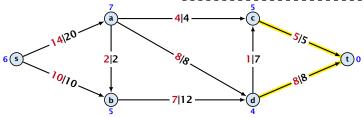


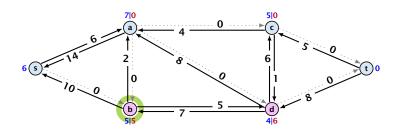
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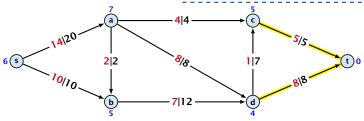
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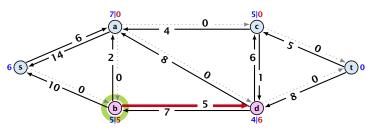


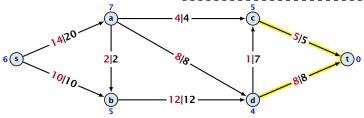


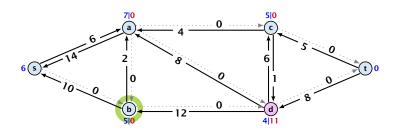
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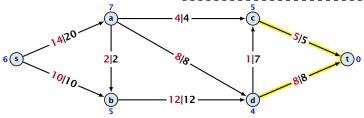


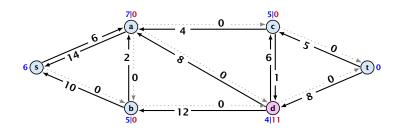
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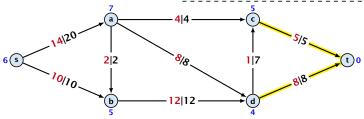


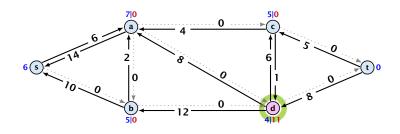




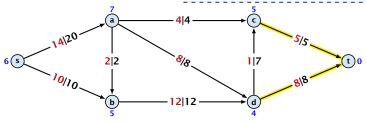




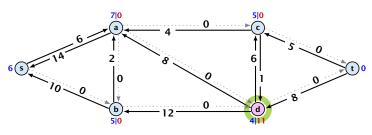


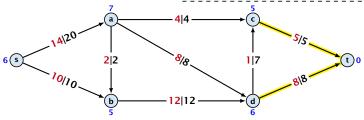


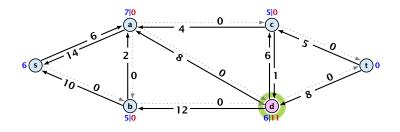
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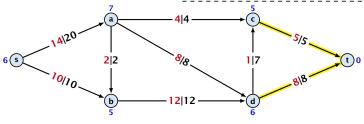
relabel to 6



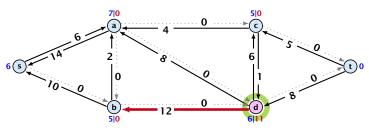


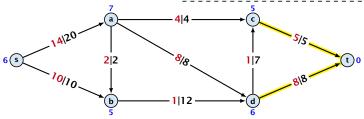


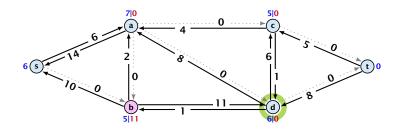
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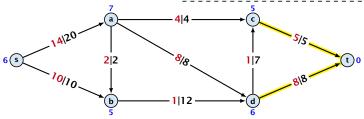


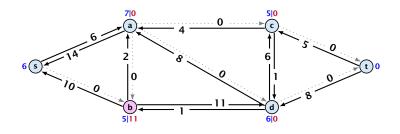
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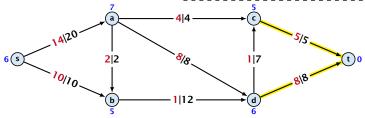


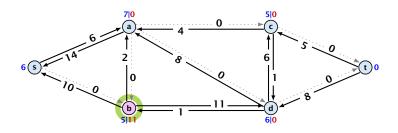




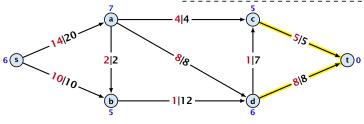




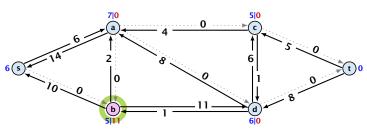


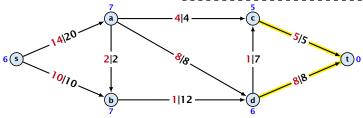


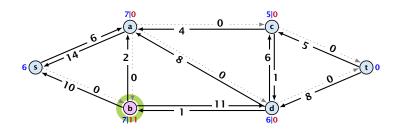
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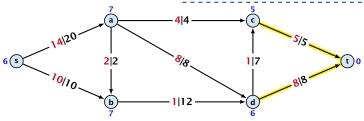
relabel to 7



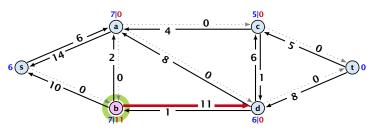


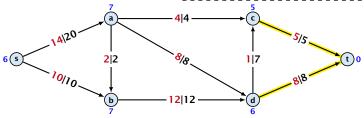


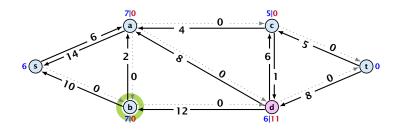
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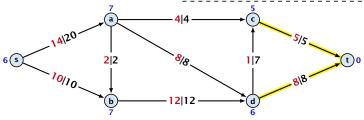


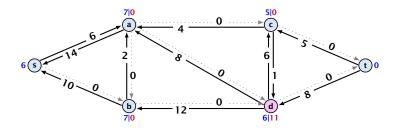
saturating and deactivating push

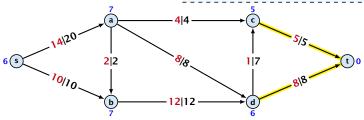


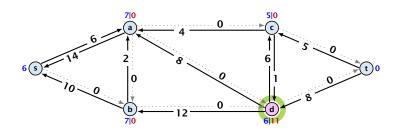




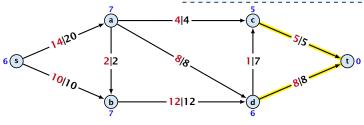




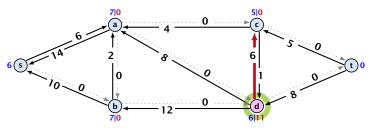


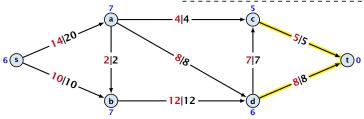


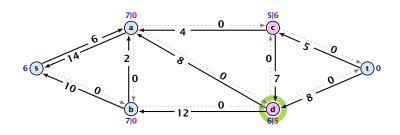
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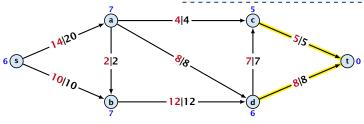
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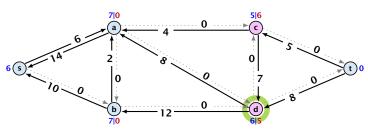


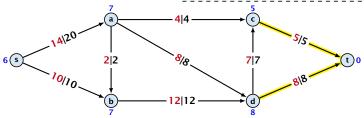


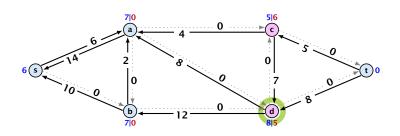
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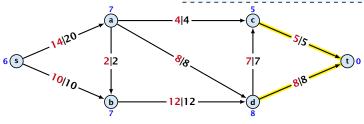
relabel to 8



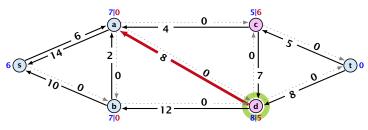


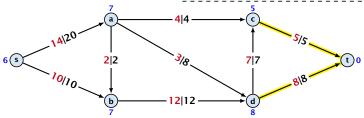


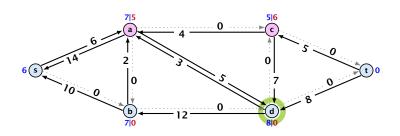
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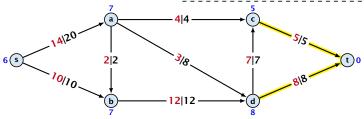


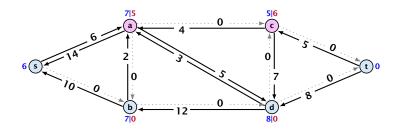
deactivating push

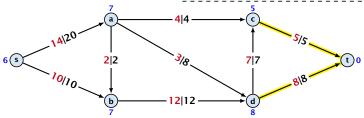


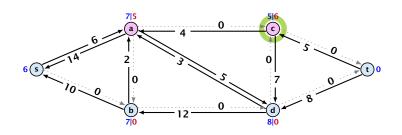




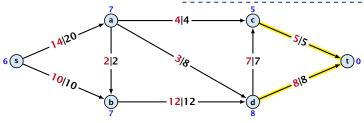




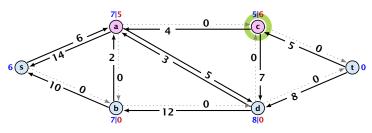


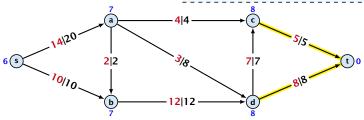


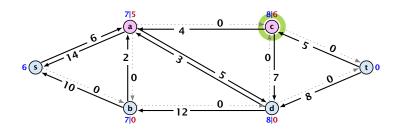
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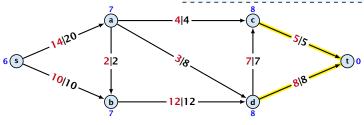
relabel to 8



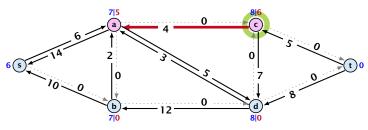


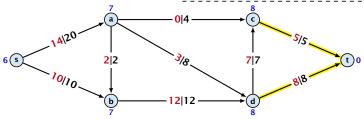


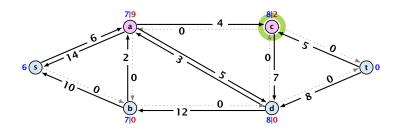
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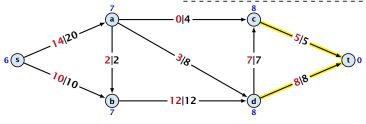
saturating push



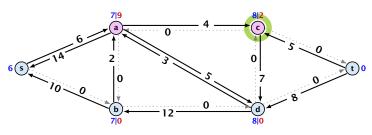


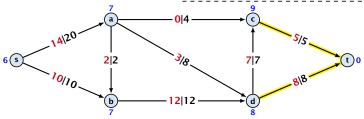


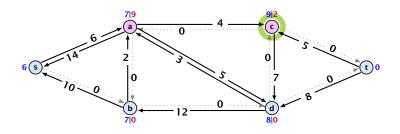
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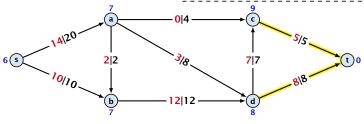
relabel to 9



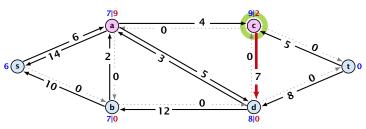


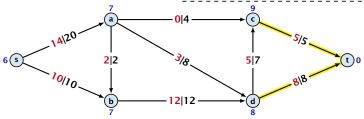


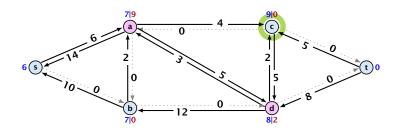
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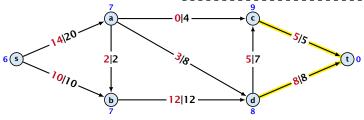


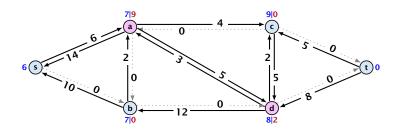
deactivating push

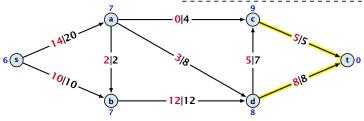


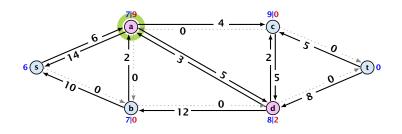




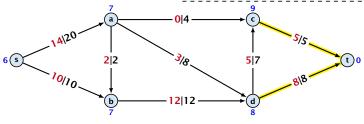




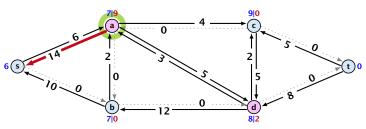


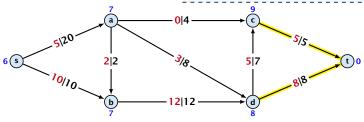


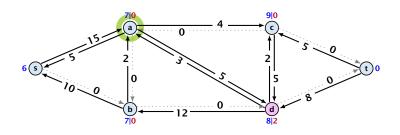
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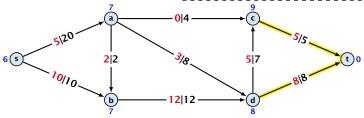


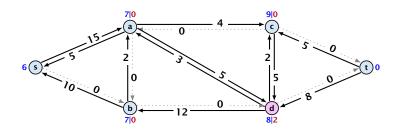
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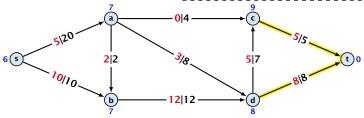


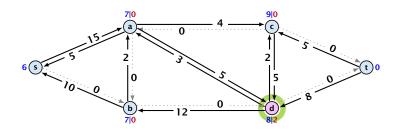




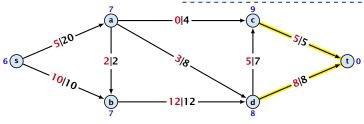




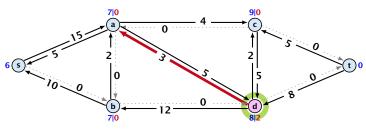


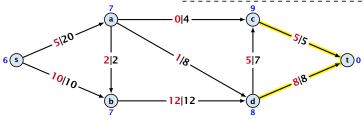


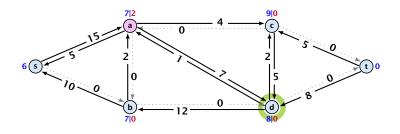
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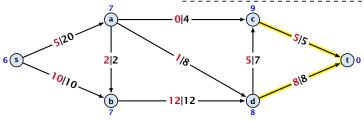


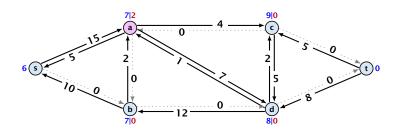
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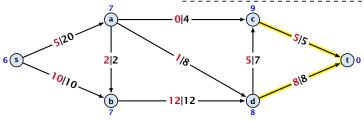


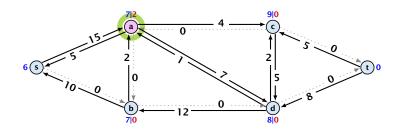




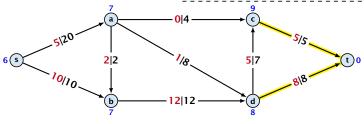




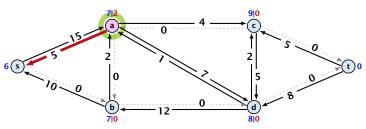


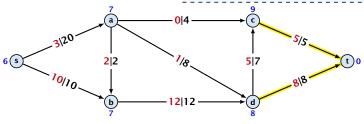


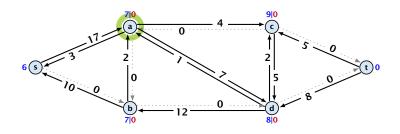
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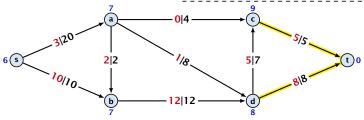


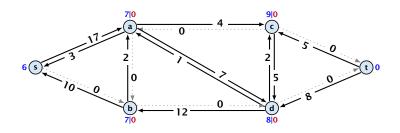
deactivating push











Note that the lemma is almost trivial. A node v having excess flow means that the current preflow ships something to v. The residual graph allows to undo flow. Therefore, there must exist a path that can undo the shipment and move it back to s. However, a formal proof is required.

Lemma 69

An active node has a path to s in the residual graph.



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Proof.

Let A denote the set of nodes that can reach s, and let B denote the remaining nodes. Note that $s \in A$.

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- ► In the residual graph there are no edges into *A*, and, hence, no edges leaving *A*/entering *B* can carry any flow.
- Let $f(B) = \sum_{v \in B} f(v)$ be the excess flow of all nodes in B.

$$f(x,y) = \left\{ \begin{array}{ll} 0 & (x,y) \notin E \\ f((x,y)) & (x,y) \in E \end{array} \right.$$

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$$= -\sum_{b \in B} \sum_{v \in A} \underbrace{f(b, v)}_{> 0}$$

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Hence, the excess flow f(b) must be 0 for every node $b \in B$.

Lemma 70

The label of a node cannot become larger than 2n-1.

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Proof.

When increasing the label at a node u there exists a path from u to s of length at most n-1. Along each edge of the path the height/label can at most drop by 1, and the label of the source is n.

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Lemma 71

There are only $O(n^2)$ relabel operations.

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- For a push from v to u the edge (v, u) must become admissible. The label of v must increase by at least 2.
- Since the label of v is at most 2n-1, there are at most n pushes along (u,v).

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- A relabel increases Φ by at most 1.
- A deactivating push decreases Φ by at least 1 as the node that is pushed from becomes inactive and has a label that is strictly larger than the target.
- Hence,

#deactivating_pushes $\leq \#$ relabels $+2n \cdot \#$ saturating_pushes $\leq \mathcal{O}(n^2m)$.

Theorem 74

There is an implementation of the generic push relabel algorithm with running time $\mathcal{O}(n^2m)$.

Proof:

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For every node maintain a list of admissible edges starting at that node. Further maintain a list of active nodes.

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check for all outgoing edges if they become admissible



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A relabel at a node u can be performed in time O(n)

- check for all outgoing edges if they become admissible
- check for all incoming edges if they become non-admissible



For special variants of push relabel algorithms we organize the neighbours of a node into a linked list (possible neighbours in the residual graph G_f). Then we use the discharge-operation:

```
Algorithm 2 discharge(u)
1: while u is active do
        v \leftarrow u.current-neighbour
2:
      if v = \text{null then}
3:
             relabel(u)
4:
5:
             u.current-neighbour ← u.neighbour-list-head
        else
6:
7:
             if (u, v) admissible then push(u, v)
             else u.current-neighbour \leftarrow v.next-in-list
8:
```

Note that *u.current-neighbour* is a global variable. It is only changed within the discharge routine, but keeps its value between consecutive calls to discharge.

If v = null in Line 3, then there is no outgoing admissible edge from u.

In order for e to become admissible the other end-point say v has to push flow to u (so that the edge (u,v) re-appears in the residual graph). For this the label of v needs to be larger than the label of u. Then in order to make (u,v) admissible the label of u has to increase.

Proof.

- While pushing from u the current-neighbour pointer is only advanced if the current edge is not admissible.
- ► The only thing that could make the edge admissible again would be a relabel at *u*.
- If we reach the end of the list (v = null) all edges are not admissible.

This shows that discharge(u) is correct, and that we can perform a relabel in Line 4.



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```
Algorithm 1 relabel-to-front(G, s, t)
1: initialize preflow
2: initialize node list L containing V \setminus \{s, t\} in any order
3: foreach u \in V \setminus \{s, t\} do
         u.current-neighbour \leftarrow u.neighbour-list-head
4:
5: u \leftarrow L.head
6: while u \neq \text{null do}
         old-height \leftarrow \ell(u)
7:
         discharge(u)
8:
         if \ell(u) > old-height then // relabel happened
9:
10:
               move u to the front of L
11:
         u \leftarrow u.next
```

Lemma 76 (Invariant)

In Line 6 of the relabel-to-front algorithm the following invariant holds.

- 1. The sequence L is topologically sorted w.r.t. the set of admissible edges; this means for an admissible edge (x,y) the node x appears before y in sequence L.
- **2.** No node before u in the list L is active.

Proof:

- Initialization:
 - 1. In the beginning s has label $n \ge 2$, and all other nodes have label 0. Hence, no edge is admissible, which means that any ordering L is permitted.
 - 2. We start with u being the head of the list; hence no node before u can be active

Maintenance:

- Pushes do no create any new admissible edges. Therefore, if discharge() does not relabel u, L is still topologically sorted.
 - After relabeling, u cannot have admissible incoming edges as such an edge (x, u) would have had a difference $\ell(x) \ell(u) \ge 2$ before the re-labeling (such edges do not exist in the residual graph).

Hence, moving u to the front does not violate the sorting property for any edge; however it fixes this property for all admissible edges leaving u that were generated by the relabeling.

Proof:

- Maintenance:
 - 2. If we do a relabel there is nothing to prove because the only node before u' (u in the next iteration) will be the current u; the discharge(u) operation only terminates when u is not active anymore.

For the case that we do not relabel, observe that the only way a predecessor could be active is that we push flow to it via an admissible arc. However, all admissible arc point to successors of u.

Note that the invariant means that for u = null we have a preflow with a valid labelling that does not have active nodes. This means we have a maximum flow.

Lemma 77

There are at most $O(n^3)$ calls to discharge(u).

Every discharge operation without a relabel advances u (the current node within list L). Hence, if we have n discharge operations without a relabel we have $u = \mathrm{null}$ and the algorithm terminates.

Therefore, the number of calls to discharge is at most $n(\#relabels + 1) = \mathcal{O}(n^3)$.

Lemma 78

The cost for all relabel-operations is only $O(n^2)$.

A relabel-operation at a node is constant time (increasing the label and resetting u.current-neighbour). In total we have $\mathcal{O}(n^2)$ relabel-operations.

Recall that a saturating push operation $(\min\{c_f(e),f(u)\}=c_f(e))$ can also be a deactivating push operation $(\min\{c_f(e),f(u)\}=f(u))$.

Lemma 79

The cost for all saturating push-operations that are **not** deactivating is only O(mn).

Note that such a push-operation leaves the node u active but makes the edge e disappear from the residual graph. Therefore the push-operation is immediately followed by an increase of the pointer u.current-neighbour.

This pointer can traverse the neighbour-list at most $\mathcal{O}(n)$ times (upper bound on number of relabels) and the neighbour-list has only degree(u) + 1 many entries (+1 for null-entry).

Lemma 80

The cost for all deactivating push-operations is only $O(n^3)$.

A deactivating push-operation takes constant time and ends the current call to discharge(). Hence, there are only $\mathcal{O}(n^3)$ such operations.

Theorem 81

The push-relabel algorithm with the rule relabel-to-front takes time $\mathcal{O}(n^3)$.

Algorithm 1 highest-label (G, s, t)

- 1: initialize preflow
- 2: foreach $u \in V \setminus \{s, t\}$ do
- 3: $u.current-neighbour \leftarrow u.neighbour-list-head$
- 4: **while** \exists active node u **do**
- 5: select active node u with highest label
- 6: $\operatorname{discharge}(u)$

Lemma 82

When using highest label the number of deactivating pushes is only $\mathcal{O}(n^3)$.

A push from a node on level ℓ can only "activate" nodes on levels strictly less than ℓ .

This means, after a deactivating push from \boldsymbol{u} a relabel is required to make \boldsymbol{u} active again.

Hence, after n deactivating pushes without an intermediate relabel there are no active nodes left.

Therefore, the number of deactivating pushes is at most $n(\#relabels + 1) = O(n^3)$.

Since a discharge-operation is terminated by a deactivating push this gives an upper bound of $\mathcal{O}(n^3)$ on the number of discharge-operations.

The cost for relabels and saturating pushes can be estimated in exactly the same way as in the case of the generic push-relabel algorithm.

Question:

How do we find the next node for a discharge operation?

Maintain lists L_i , $i \in \{0, ..., 2n\}$, where list L_i contains active nodes with label i (maintaining these lists induces only constant additional cost for every push-operation and for every relabel-operation).

After a discharge operation terminated for a node u with label k, traverse the lists $L_k, L_{k-1}, \ldots, L_0$, (in that order) until you find a non-empty list.

Unless the last (deactivating) push was to s or t the list k-1 must be non-empty (i.e., the search takes constant time).

Hence, the total time required for searching for active nodes is at most

$$O(n^3) + n(\# deactivating-pushes-to-s-or-t)$$

Lemma 83

The number of deactivating pushes to s or t is at most $O(n^2)$.

With this lemma we get

Theorem 84

The push-relabel algorithm with the rule highest-label takes time $\mathcal{O}(n^3)$.

Proof of the Lemma.

- We only show that the number of pushes to the source is at most $\mathcal{O}(n^2)$. A similar argument holds for the target.
- After a node v (which must have $\ell(v) = n+1$) made a deactivating push to the source there needs to be another node whose label is increased from $\leq n+1$ to n+2 before v can become active again.
- This happens for every push that v makes to the source. Since, every node can pass the threshold n + 2 at most once, v can make at most n pushes to the source.
- As this holds for every node the total number of pushes to the source is at most $\mathcal{O}(n^2)$.

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Mincost Flow

Problem Definition:

$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & 0 \leq f(e) \leq u(e) \\ & & \forall v \in V: & f(v) = b(v) \end{aligned}$$

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- G = (V, E) is a directed graph.
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```

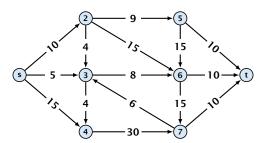
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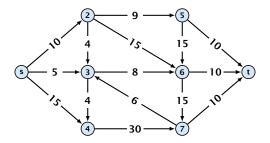
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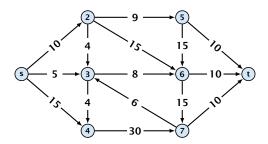
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- G = (V, E) is a directed graph.
- ▶ $u: E \to \mathbb{R}_0^+ \cup \{\infty\}$ is the capacity function.
- ► $c: E \to \mathbb{R}$ is the cost function (note that c(e) may be negative).
- ▶ $b: V \to \mathbb{R}$, $\sum_{v \in V} b(v) = 0$ is a demand function.

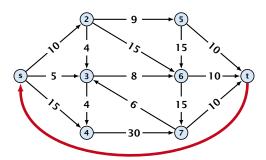




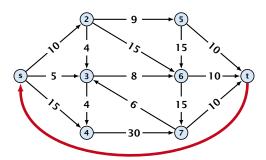
Given a flow network for a standard maxflow problem.



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- Given a flow network for a standard maxflow problem.
- Set b(v) = 0 for every node. Keep the capacity function u for all edges. Set the cost c(e) for every edge to 0.
- ▶ Add an edge from t to s with infinite capacity and cost -1.



- Given a flow network for a standard maxflow problem.
- Set b(v) = 0 for every node. Keep the capacity function u for all edges. Set the cost c(e) for every edge to 0.
- ▶ Add an edge from t to s with infinite capacity and cost -1.
- ► Then, $val(f^*) = -cost(f_{min})$, where f^* is a maxflow, and f_{min} is a mincost-flow.



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Solve decision version of maxflow:

• Given a flow network for a standard maxflow problem, and a value k.

Solve decision version of maxflow:

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Solve decision version of maxflow:

- Given a flow network for a standard maxflow problem, and a value k.
- Set b(v) = 0 for every node apart from s or t. Set b(s) = -k and b(t) = k.
- Set edge-costs to zero, and keep the capacities.
- There exists a maxflow of value at least k if and only if the mincost-flow problem is feasible.

Generalization

Our model:

min
$$\sum_{e} c(e) f(e)$$

s.t. $\forall e \in E: 0 \le f(e) \le u(e)$
 $\forall v \in V: f(v) = b(v)$

where
$$b: V \to \mathbb{R}$$
, $\sum_{v} b(v) = 0$; $u: E \to \mathbb{R}_0^+ \cup \{\infty\}$; $c: E \to \mathbb{R}$;

Generalization

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where
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A more general model?

$$\begin{aligned} & \text{min} & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & & \forall v \in V: & a(v) \leq f(v) \leq b(v) \end{aligned}$$

where
$$a: V \to \mathbb{R}$$
, $b: V \to \mathbb{R}$; $\ell: E \to \mathbb{R} \cup \{-\infty\}$, $u: E \to \mathbb{R} \cup \{\infty\}$ $c: E \to \mathbb{R}$;

Generalization

Differences

- Flow along an edge e may have non-zero lower bound $\ell(e)$.
- Flow along e may have negative upper bound u(e).
- ▶ The demand at a node v may have lower bound a(v) and upper bound b(v) instead of just lower bound = upper bound = b(v).

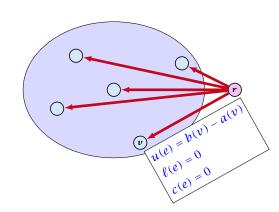
$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V: & a(v) \leq f(v) \leq b(v) \end{aligned}$$

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We can assume that a(v) = b(v):

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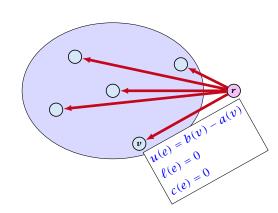
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We can assume that a(v) = b(v):

Add new node r.

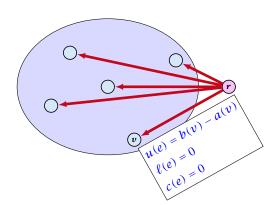


$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V: & a(v) \leq f(v) \leq b(v) \end{aligned}$$

We can assume that a(v) = b(v):

Add new node r.

 $\text{Add edge } (r,v) \text{ for all } v \in V.$



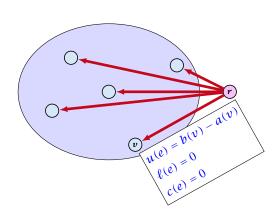
$$\begin{aligned} & \min \quad \sum_{e} c(e) f(e) \\ & \text{s.t.} \quad \forall e \in E : \ \ell(e) \leq f(e) \leq u(e) \\ & \quad \forall v \in V : \ a(v) \leq f(v) \leq b(v) \end{aligned}$$

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Add edge (r, v) for all $v \in V$.

Set $\ell(e) = c(e) = 0$ for these edges.



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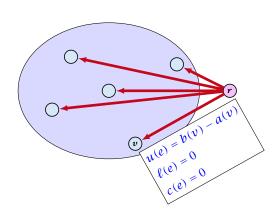
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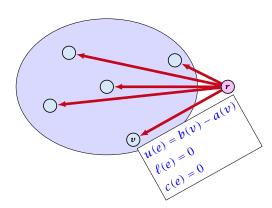
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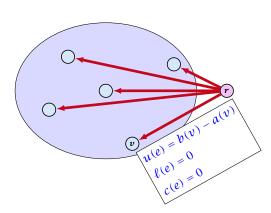
Add edge (r, v) for all $v \in V$.

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Set u(e) = b(v) - a(v) for edge (r, v).

Set a(v) = b(v) for all $v \in V$.

Set $b(r) = -\sum_{v \in V} b(v)$.



$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E : & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V : & a(v) \leq f(v) \leq b(v) \end{aligned}$$

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Add edge (r, v) for all $v \in V$.

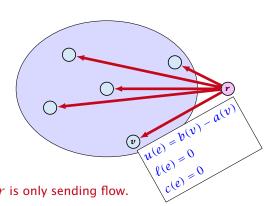
Set $\ell(e) = c(e) = 0$ for these edges.

Set u(e) = b(v) - a(v) for edge (r, v).

Set a(v) = b(v) for all $v \in V$.

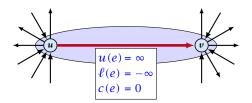
Set
$$b(r) = -\sum_{v \in V} b(v)$$
.

 $-\sum_{v}b(v)$ is negative; hence r is only sending flow.



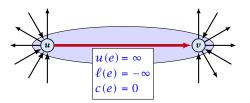
$$\begin{aligned} & \text{min} & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V: & f(v) = b(v) \end{aligned}$$

We can assume that either $\ell(e) \neq -\infty$ or $u(e) \neq \infty$:



$$\begin{aligned} & \text{min} & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V: & f(v) = b(v) \end{aligned}$$

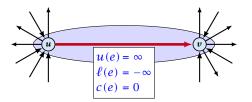
We can assume that either $\ell(e) \neq -\infty$ or $u(e) \neq \infty$:



If c(e) = 0 we can contract the edge/identify nodes u and v.

$$\begin{aligned} & \text{min} & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & \forall v \in V: & f(v) = b(v) \end{aligned}$$

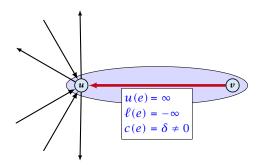
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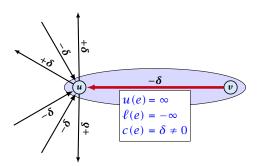
If c(e) = 0 we can contract the edge/identify nodes u and v.

If $c(e) \neq 0$ we can transform the graph so that c(e) = 0.

We can transform any network so that a particular edge has c(e) = 0:

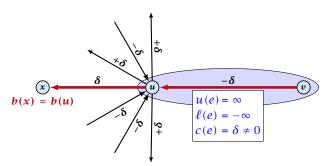


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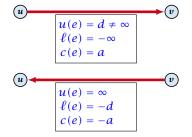


Additionally we set b(u) = 0.

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$$\begin{aligned} & \min & \quad \sum_{e} c(e) f(e) \\ & \text{s.t.} & \quad \forall e \in E : \quad \ell(e) \leq f(e) \leq u(e) \\ & \quad \forall v \in V : \quad f(v) = b(v) \end{aligned}$$

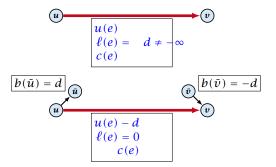
We can assume that $\ell(e) \neq -\infty$:



Replace the edge by an edge in opposite direction.

$$\begin{aligned} & \min & & \sum_{e} c(e) f(e) \\ & \text{s.t.} & & \forall e \in E: & \ell(e) \leq f(e) \leq u(e) \\ & & & \forall v \in V: & f(v) = b(v) \end{aligned}$$

We can assume that $\ell(e) = 0$:



The added edges have infinite capacity and cost c(e)/2.

Applications

Caterer Problem

• She needs to supply r_i napkins on N successive days.

Applications

Caterer Problem

- She needs to supply r_i napkins on N successive days.
- ▶ She can buy new napkins at p cents each.

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- She can use a slow laundry that takes k > m days and costs s cents each.

Applications

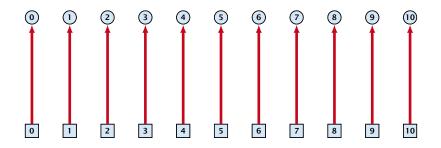
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- At the end of each day she should determine how many to send to each laundry and how many to buy in order to fulfill demand.

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- She can use a slow laundry that takes k > m days and costs s cents each.
- At the end of each day she should determine how many to send to each laundry and how many to buy in order to fulfill demand.
- Minimize cost.

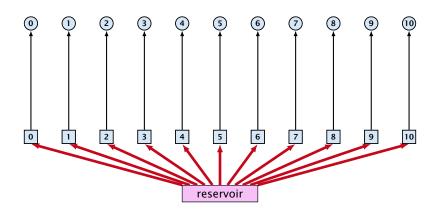


day edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = r_i$;

cost: c(e) = 0

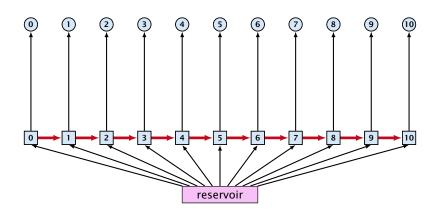


buy edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = p

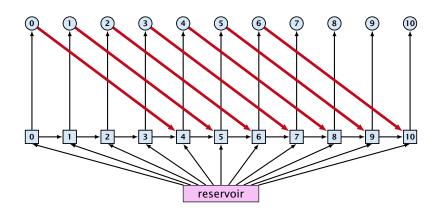


forward edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = 0

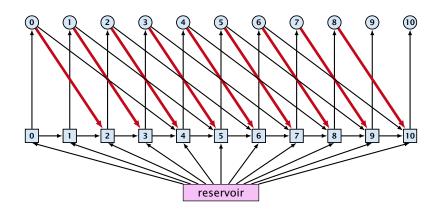


slow edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = s

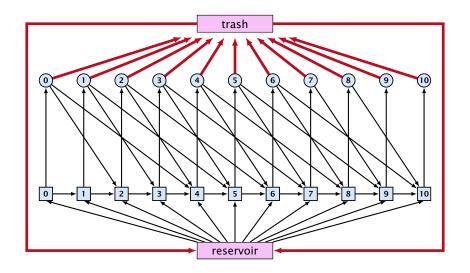


fast edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = f

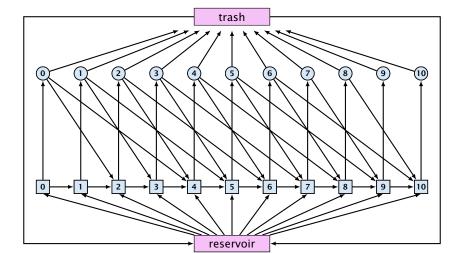


trash edges:

upper bound: $u(e_i) = \infty$;

lower bound: $\ell(e_i) = 0$;

cost: c(e) = 0



Residual Graph

Version A:

The residual graph G' for a mincost flow is just a copy of the graph G.

If we send f(e) along an edge, the corresponding edge e' in the residual graph has its lower and upper bound changed to $\ell(e') = \ell(e) - f(e)$ and u(e') = u(e) - f(e).

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If we send f(e) along an edge, the corresponding edge e' in the residual graph has its lower and upper bound changed to $\ell(e') = \ell(e) - f(e)$ and u(e') = u(e) - f(e).

Version B:

The residual graph for a mincost flow is exactly defined as the residual graph for standard flows, with the only exception that one needs to define a cost for the residual edge.

For a flow of z from u to v the residual edge (v,u) has capacity z and a cost of -c((u,v)).



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A circulation in a graph G=(V,E) is a function $f:E\to\mathbb{R}^+$ that has an excess flow f(v)=0 for every node $v\in V$.

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A circulation is feasible if it fulfills capacity constraints, i.e., $f(e) \le u(e)$ for every edge of G.

A given flow is a mincost-flow if and only if the corresponding residual graph G_f does not have a feasible circulation of negative cost.

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Then f+g is a feasible flow with cost $\cos t(f)+\cos t(g)<\cos t(f)$. Hence, f is not minimum cost.

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 \Rightarrow Suppose that g is a feasible circulation of negative cost in the residual graph.

Then f + g is a feasible flow with cost cost(f) + cost(g) < cost(f). Hence, f is not minimum cost.

 \Leftarrow Let f be a non-mincost flow, and let f^* be a min-cost flow. We need to show that the residual graph has a feasible circulation with negative cost.

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 \leftarrow Let f be a non-mincost flow, and let f^* be a min-cost flow. We need to show that the residual graph has a feasible circulation with negative cost.

Clearly $f^* - f$ is a circulation of negative cost. One can also easily see that it is feasible for the residual graph. (after sending -f in the residual graph (pushing all flow back) we arrive at the original graph; for this f^* is clearly feasible)

Lemma 86

A graph (without zero-capacity edges) has a feasible circulation of negative cost if and only if it has a negative cycle w.r.t. edge-weights $c: E \to \mathbb{R}$.

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Proof.

Suppose that we have a negative cost circulation.

Lemma 86

A graph (without zero-capacity edges) has a feasible circulation of negative cost if and only if it has a negative cycle w.r.t. edge-weights $c : E \to \mathbb{R}$.

- Suppose that we have a negative cost circulation.
- Find directed cycle only using edges that have non-zero flow.

Lemma 86

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- Suppose that we have a negative cost circulation.
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- If this cycle has negative cost you are done.

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- Otherwise send flow in opposite direction along the cycle until the bottleneck edge(s) does not carry any flow.

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- Find directed cycle only using edges that have non-zero flow.
- If this cycle has negative cost you are done.
- Otherwise send flow in opposite direction along the cycle until the bottleneck edge(s) does not carry any flow.
- You still have a circulation with negative cost.

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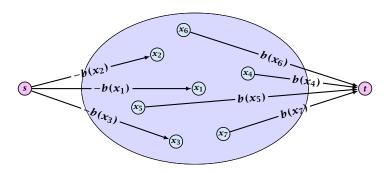
- Suppose that we have a negative cost circulation.
- Find directed cycle only using edges that have non-zero flow.
- If this cycle has negative cost you are done.
- Otherwise send flow in opposite direction along the cycle until the bottleneck edge(s) does not carry any flow.
- You still have a circulation with negative cost.
- Repeat.



Algorithm 48 CycleCanceling(G = (V, E), c, u, b)

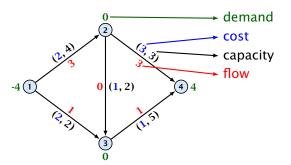
- 1: establish a feasible flow f in G
- 2: **while** G_f contains negative cycle **do**
- 3: use Bellman-Ford to find a negative circuit Z
- 4: $\delta \leftarrow \min\{u_f(e) \mid e \in Z\}$
- 5: augment δ units along Z and update G_f

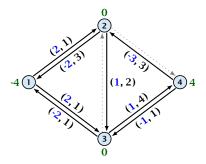
How do we find the initial feasible flow?

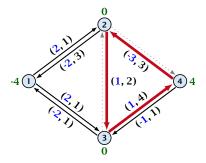


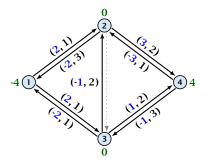
- Connect new node s to all nodes with negative b(v)-value.
- Connect nodes with positive b(v)-value to a new node t.
- ► There exist a feasible flow in the original graph iff in the resulting graph there exists an *s-t* flow of value

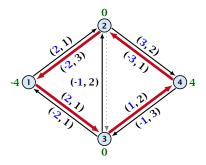
$$\sum_{v:b(v)<0} (-b(v)) = \sum_{v:b(v)>0} b(v) .$$

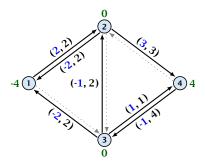












Lemma 87

The improving cycle algorithm runs in time $O(nm^2CU)$, for integer capacities and costs, when for all edges e, $|c(e)| \le C$ and $|u(e)| \le U$.

- Running time of Bellman-Ford is O(mn).
- ▶ Pushing flow along the cycle can be done in time O(n).
- Each iteration decreases the total cost by at least 1.
- ► The true optimum cost must lie in the interval [-mCU,...,+mCU].

Note that this lemma is weak since it does not allow for edges with infinite capacity.

14 Mincost Flow

A general mincost flow problem is of the following form:

$$\begin{aligned} & \min \quad \sum_{e} c(e) f(e) \\ & \text{s.t.} \quad \forall e \in E : \quad \ell(e) \leq f(e) \leq u(e) \\ & \quad \forall v \in V : \quad a(v) \leq f(v) \leq b(v) \end{aligned}$$
 where $a: V \to \mathbb{R}, \ b: V \to \mathbb{R}; \ \ell: E \to \mathbb{R} \cup \{-\infty\}, \ u: E \to \mathbb{R} \cup \{\infty\}$ $c: E \to \mathbb{R};$

Lemma 88 (without proof)

A general mincost flow problem can be solved in polynomial time.

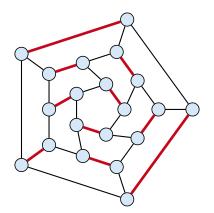
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Part V

Matchings

Matching

- ▶ Input: undirected graph G = (V, E).
- ► $M \subseteq E$ is a matching if each node appears in at most one edge in M.
- Maximum Matching: find a matching of maximum cardinality



16 Bipartite Matching via Flows

Which flow algorithm to use?

- Generic augmenting path: $\mathcal{O}(m \operatorname{val}(f^*)) = \mathcal{O}(mn)$.
- Capacity scaling: $\mathcal{O}(m^2 \log C) = \mathcal{O}(m^2)$.
- Shortest augmenting path: $O(mn^2)$.

For unit capacity simple graphs shortest augmenting path can be implemented in time $\mathcal{O}(m\sqrt{n})$.

Definitions.

Given a matching M in a graph G, a vertex that is not incident to any edge of M is called a free vertex w.r..t. M.

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- For a matching *M* a path *P* in *G* is called an alternating path if edges in *M* alternate with edges not in *M*.

Definitions.

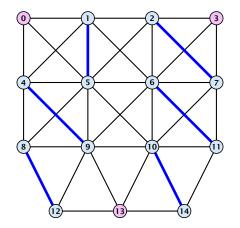
- Given a matching M in a graph G, a vertex that is not incident to any edge of M is called a free vertex w.r..t. M.
- For a matching M a path P in G is called an alternating path if edges in M alternate with edges not in M.
- An alternating path is called an augmenting path for matching M if it ends at distinct free vertices.

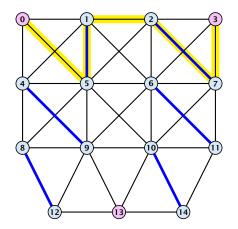
Definitions.

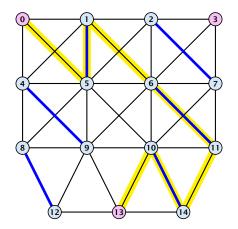
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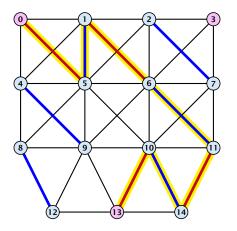
Theorem 89

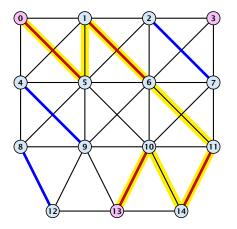
A matching M is a maximum matching if and only if there is no augmenting path w.r.t. M.

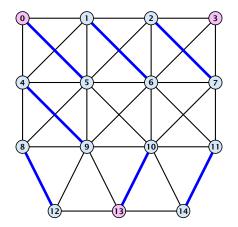












Proof.

 \Rightarrow If M is maximum there is no augmenting path P, because we could switch matching and non-matching edges along P. This gives matching $M' = M \oplus P$ with larger cardinality.

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Proof.

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Each vertex can be incident to at most two edges (one from M and one from M'). Hence, the connected components are alternating cycles or alternating path.

Proof.

- \Rightarrow If M is maximum there is no augmenting path P, because we could switch matching and non-matching edges along P. This gives matching $M' = M \oplus P$ with larger cardinality.
- \Leftarrow Suppose there is a matching M' with larger cardinality. Consider the graph H with edge-set $M' \oplus M$ (i.e., only edges that are in either M or M' but not in both).

Each vertex can be incident to at most two edges (one from M and one from M'). Hence, the connected components are alternating cycles or alternating path.

As |M'| > |M| there is one connected component that is a path P for which both endpoints are incident to edges from M'. P is an alternating path.

Algorithmic idea:

As long as you find an augmenting path augment your matching using this path. When you arrive at a matching for which no augmenting path exists you have a maximum matching.

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As long as you find an augmenting path augment your matching using this path. When you arrive at a matching for which no augmenting path exists you have a maximum matching.

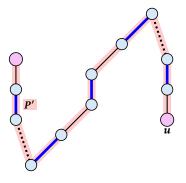
Theorem 90

Let G be a graph, M a matching in G, and let u be a free vertex w.r.t. M. Further let P denote an augmenting path w.r.t. M and let $M' = M \oplus P$ denote the matching resulting from augmenting M with P. If there was no augmenting path starting at u in M then there is no augmenting path starting at u in M'.

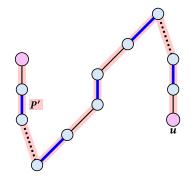
The above theorem allows for an easier implementation of an augmenting path algorithm. Once we checked for augmenting paths starting from u we don't have to check for such paths in future rounds.

Proof

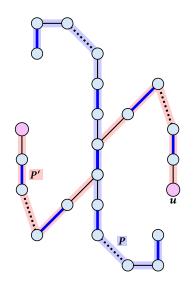
Assume there is an augmenting path P' w.r.t. M' starting at u.



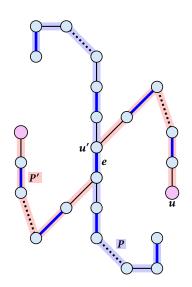
- Assume there is an augmenting path P' w.r.t. M' starting at u.
- If P' and P are node-disjoint, P' is also augmenting path w.r.t. M (∮).



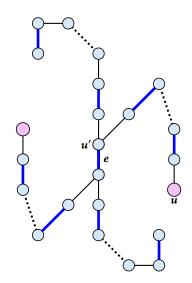
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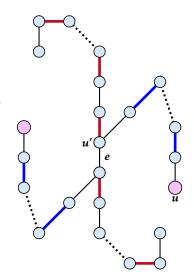
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- Let u' be the first node on P' that is in P, and let e be the matching edge from M' incident to u'.



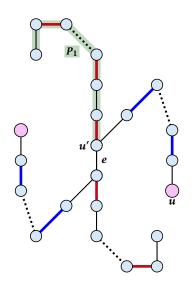
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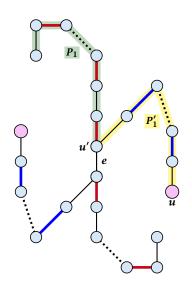
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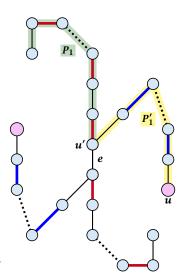
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- u' splits P into two parts one of which does not contain e. Call this part P_1 . Denote the sub-path of P' from u to u' with P'_1 .



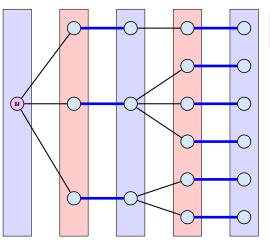
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- Assume there is an augmenting path P' w.r.t. M' starting at u.
- If P' and P are node-disjoint, P' is also augmenting path w.r.t. M (∮).
- Let u' be the first node on P' that is in P, and let e be the matching edge from M' incident to u'.
- u' splits P into two parts one of which does not contain e. Call this part P_1 . Denote the sub-path of P' from u to u' with P'_1 .
- $P_1 \circ P_1'$ is augmenting path in M (3).

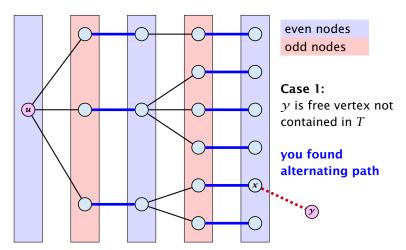


Construct an alternating tree.

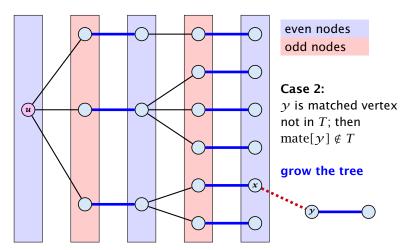


even nodes odd nodes

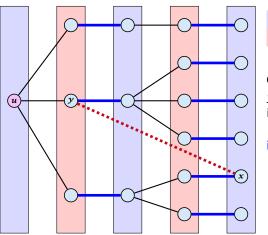
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Construct an alternating tree.



Construct an alternating tree.

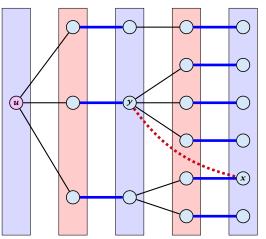


even nodes odd nodes

Case 3: *y* is already contained in *T* as an odd vertex

ignore successor y

Construct an alternating tree.



even nodes odd nodes

Case 4:

y is already contained in T as an even vertex

can't ignore ${m y}$

does not happen in bipartite graphs

```
Algorithm 49 BiMatch(G, match)
 1: for x \in V do mate[x] \leftarrow 0;
2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
4: r \leftarrow r + 1
 5: if mate[r] = 0 then
6:
          for i = 1 to n do parent[i'] \leftarrow 0
    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
7:
8:
    while aug = false and Q \neq \emptyset do
9:
              x \leftarrow Q. dequeue();
10:
              for y \in A_x do
11:
                  if mate[\gamma] = 0 then
12:
                      augm(mate, parent, y);
13:
                     aua ← true:
                     free \leftarrow free - 1:
14:
15:
                  else
16:
                     if parent[y] = 0 then
                         parent[y] \leftarrow x;
17:
```

Q. enqueue(mate[y]);

18:

graph $G = (S \cup S', E)$ $S = \{1, \dots, n\}$ $S' = \{1', \dots, n'\}$

```
Algorithm 49 BiMatch(G, match)
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14:
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16:
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                         parent[y] \leftarrow x;
17:
```

Q. enqueue(mate[y]);

18:

start with an empty matching

```
Algorithm 49 BiMatch(G, match)
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    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
 7:
8:
    while aug = false and Q \neq \emptyset do
9:
              x \leftarrow Q. dequeue();
```

for $y \in A_x$ do

else

if mate[y] = 0 then

aua ← true: $free \leftarrow free - 1$:

augm(mate, parent, y);

if parent[y] = 0 then $parent[y] \leftarrow x$;

10:

11:

12:

13:

14: 15:

16:

17:

18:

free: number of unmatched nodes in S r: root of current tree

```
Q. enqueue(mate[y]);
```

Algorithm 49 BiMatch(*G*, *match*) 1: **for** $x \in V$ **do** $mate[x] \leftarrow 0$;

2: $r \leftarrow 0$; free $\leftarrow n$;

3: while $free \ge 1$ and r < n do

 $\gamma \leftarrow \gamma + 1$ 4:

6:

7: 8:

9:

10:

11:

12:

13:

14: 15:

16:

17:

18:

5: **if** mate[r] = 0 **then**

for i = 1 to n do $parent[i'] \leftarrow 0$

 $Q \leftarrow \emptyset$; Q. append(r); $aug \leftarrow false$;

 $x \leftarrow Q$. dequeue();

for $y \in A_x$ do

else

while aug = false and $Q \neq \emptyset$ do

if mate[y] = 0 then

augm(mate, parent, y);

aua ← true:

 $free \leftarrow free - 1$:

if parent[y] = 0 then

 $parent[y] \leftarrow x$; Q. enqueue(mate[y]);

as long as there are unmatched nodes and we did not yet try to grow from all nodes we continue

```
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12:
                      augm(mate, parent, y);
13:
                      aua ← true:
                      free \leftarrow free - 1:
14:
15:
                  else
```

if parent[y] = 0 then $parent[y] \leftarrow x$;

Q. enqueue(mate[y]);

16:

17: 18: r is the new node that we grow from.

```
Algorithm 49 BiMatch(G, match)
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2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
    \gamma \leftarrow \gamma + 1
 4:
 5: if mate[r] = 0 then
           for i = 1 to n do parent[i'] \leftarrow 0
6:
    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
 7:
 8:
          while aug = false and Q \neq \emptyset do
9:
              x \leftarrow Q. dequeue();
10:
               for y \in A_x do
                  if mate[y] = 0 then
11:
12:
                      augm(mate, parent, y);
```

else

 $aug \leftarrow true;$ free $\leftarrow free - 1$:

if parent[y] = 0 then $parent[y] \leftarrow x$;

O. enqueue(mate[v]);

13:

14: 15:

16:

17: 18: If *r* is free start tree construction

```
Algorithm 49 BiMatch(G, match)

1: for x \in V do mate[x] \leftarrow 0;

2: r \leftarrow 0; free \leftarrow n;

3: while free \geq 1 and r < n do

4: r \leftarrow r + 1

5: if mate[r] = 0 then

6: for i = 1 to n do parent[i'] \leftarrow 0

7: Q \leftarrow \varnothing; Q. append(r); auq \leftarrow false;
```

while aug = false and $Q \neq \emptyset$ do

if mate[y] = 0 then

 $aug \leftarrow true;$ $free \leftarrow free - 1:$

augm(mate, parent, y);

if parent[y] = 0 then $parent[y] \leftarrow x$;

Q. enqueue(mate[y]);

 $x \leftarrow Q$. dequeue();

for $y \in A_x$ do

else

8:

9:

10:

11:

12:

13:

14: 15:

16:

17:

18:

Initialize an empty tree. Note that only nodes i' have parent pointers.

```
Algorithm 49 BiMatch(G, match)
1: for x \in V do mate[x] \leftarrow 0;
```

2: $r \leftarrow 0$; free $\leftarrow n$;

3: while $free \ge 1$ and r < n do

4: $r \leftarrow r + 1$

5: **if** mate[r] = 0 **then**

6:

7:

8:

9:

10:

11:

12:

13:

14: 15:

16:

17: 18:

for i = 1 to n do $parent[i'] \leftarrow 0$

 $Q \leftarrow \emptyset$; Q. append(r); aug \leftarrow false;

while aug = false and $Q \neq \emptyset$ do

 $x \leftarrow Q$. dequeue();

for $y \in A_x$ do

if mate[y] = 0 then

augm(mate, parent, y);

aua ← true:

 $free \leftarrow free - 1$:

else if parent[y] = 0 then $parent[y] \leftarrow x$; Q. enqueue(mate[y]);

Q is a queue (BFS!!!). aug is a Boolean that stores whether we already found an

augmenting path.

Algorithm 49 BiMatch(*G*, *match*) 1: **for** $x \in V$ **do** $mate[x] \leftarrow 0$;

2: $r \leftarrow 0$; free $\leftarrow n$;

6:

7: 8:

9:

10: 11:

12:

13:

14: 15:

16:

17:

18:

3: while $free \ge 1$ and r < n do

4: $r \leftarrow r + 1$

5: **if** mate[r] = 0 **then**

for i = 1 to n do $parent[i'] \leftarrow 0$

 $Q \leftarrow \emptyset$; Q. append(r); aug \leftarrow false;

while auq = false and $Q \neq \emptyset$ do

 $x \leftarrow Q$. dequeue(); for $\gamma \in A_{\chi}$ do

if $mate[\gamma] = 0$ then

augm(mate, parent, y);

aua ← true:

 $free \leftarrow free - 1$:

else if parent[y] = 0 then $parent[y] \leftarrow x$; Q. enqueue(mate[y]);

as long as we did not augment and there are still unexamined leaves continue...

```
Algorithm 49 BiMatch(G, match)
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11:
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13:
                     aua ← true:
                     free \leftarrow free - 1:
14:
15:
                  else
16:
                     if parent[y] = 0 then
```

17: 18: $parent[y] \leftarrow x$;

O. enqueue(mate[v]);

take next unexamined leaf

```
Algorithm 49 BiMatch(G, match)

1: for x \in V do mate[x] \leftarrow 0;
2: r \leftarrow 0; free \leftarrow n;
3: while free \geq 1 and r < n do
4: r \leftarrow r + 1
5: if mate[r] = 0 then
6: for i = 1 to n do parent[i'] \leftarrow 0
7: Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
```

while aug = false and $Q \neq \emptyset$ do

if mate[v] = 0 then

 $aug \leftarrow true;$ free $\leftarrow free - 1$:

augm(mate, parent, y);

if parent[y] = 0 then $parent[y] \leftarrow x$;

Q. enqueue(mate[y]);

 $x \leftarrow Q$. dequeue();

for $y \in A_x$ do

else

8:

9:

10:

11:

12:

13:

14: 15:

16:

17:

18:

if x has unmatched neighbour we found an augmenting path (note that $y \neq r$ because we are in a bipartite graph)

```
Algorithm 49 BiMatch(G, match)
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4: r \leftarrow r + 1
5: if mate[r] = 0 then
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    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
7:
8:
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9:
              x \leftarrow Q. dequeue();
10:
              for y \in A_x do
11:
                 if mate[v] = 0 then
12:
                     augm(mate, parent, y);
13:
                     aua ← true:
                     free \leftarrow free - 1:
14:
15:
                 else
16:
                     if parent[y] = 0 then
                        parent[y] \leftarrow x;
17:
18:
                        Q. enqueue(mate[y]);
```

do an augmentation...

Algorithm 49 BiMatch(*G*, *match*) 1: **for** $x \in V$ **do** $mate[x] \leftarrow 0$; 2: $r \leftarrow 0$; free $\leftarrow n$;

3: while $free \ge 1$ and r < n do

6:

7:

8:

9:

10:

11:

12:

13:

14: 15:

16:

17:

18:

4: $r \leftarrow r + 1$

5: **if** mate[r] = 0 **then**

for i = 1 to n do $parent[i'] \leftarrow 0$

 $x \leftarrow Q$. dequeue();

for $y \in A_x$ do

else

aua ← true: $free \leftarrow free - 1$:

augm(mate, parent, y);

if parent[y] = 0 then $parent[y] \leftarrow x$;

Q. enqueue(mate[y]);

 $Q \leftarrow \emptyset$; Q. append(r); $aug \leftarrow false$;

setting aug = true

ensures that the tree

construction will not

continue

while aug = false and $Q \neq \emptyset$ do

if mate[y] = 0 then

```
Algorithm 49 BiMatch(G, match)
 1: for x \in V do mate[x] \leftarrow 0;
2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
 4: r \leftarrow r + 1
 5: if mate[r] = 0 then
6:
          for i = 1 to n do parent[i'] \leftarrow 0
    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
 7:
8:
    while aug = false and Q \neq \emptyset do
9:
              x \leftarrow Q. dequeue();
10:
              for y \in A_x do
```

if $mate[\gamma] = 0$ then

free \leftarrow free -1;

aua ← true:

else

augm(mate, parent, y);

if parent[y] = 0 then $parent[y] \leftarrow x$;

Q. enqueue(mate[y]);

11:

12:

13:

14:

15:

16:

17:

18:

```
reduce number of free
nodes
```

```
Algorithm 49 BiMatch(G, match)
 1: for x \in V do mate[x] \leftarrow 0;
2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
 4: r \leftarrow r + 1
 5: if mate[r] = 0 then
6:
          for i = 1 to n do parent[i'] \leftarrow 0
    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
7:
8:
    while aug = false and Q \neq \emptyset do
9:
              x \leftarrow Q. dequeue();
10:
              for y \in A_x do
                  if mate[y] = 0 then
11:
12:
                     augm(mate, parent, y);
13:
                     aua ← true:
                     free \leftarrow free - 1;
14:
15:
                  else
16:
                     if parent[y] = 0
                                         then
                         parent[y] \leftarrow x;
17:
18:
                         Q. enqueue(mate[y]);
```

if y is not in the tree yet

```
Algorithm 49 BiMatch(G, match)
 1: for x \in V do mate[x] \leftarrow 0;
2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
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              x \leftarrow Q. dequeue();
10:
              for y \in A_x do
11:
                  if mate[\gamma] = 0 then
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                     augm(mate, parent, y);
13:
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                     free \leftarrow free - 1:
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                  else
16:
                     if parent[y] = 0 then
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```

...put it into the tree

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6:

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for i = 1 to n do $parent[i'] \leftarrow 0$

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 $Q \leftarrow \emptyset$; Q. append(r); $aug \leftarrow false$;

while aug = false and $Q \neq \emptyset$ do

 $x \leftarrow Q$. dequeue();

if mate[y] = 0 then

augm(mate, parent, y);

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 $free \leftarrow free - 1$:

if parent[y] = 0 then $parent[y] \leftarrow x$;

15: else 16: 17:

18: Q. enqueue(mate[y]); add its buddy to the set of unexamined leaves

18 Weighted Bipartite Matching

Weighted Bipartite Matching/Assignment

- ▶ Input: undirected, bipartite graph $G = L \cup R, E$.
- ▶ an edge $e = (\ell, r)$ has weight $w_e \ge 0$
- find a matching of maximum weight, where the weight of a matching is the sum of the weights of its edges

Simplifying Assumptions (wlog [why?]):

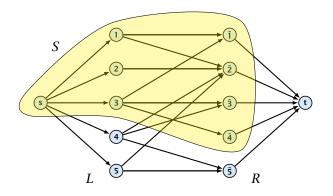
- ightharpoonup assume that |L| = |R| = n
- assume that there is an edge between every pair of nodes $(\ell,r) \in V \times V$
- can assume goal is to construct maximum weight perfect matching

Weighted Bipartite Matching

Theorem 91 (Halls Theorem)

A bipartite graph $G = (L \cup R, E)$ has a perfect matching if and only if for all sets $S \subseteq L$, $|\Gamma(S)| \ge |S|$, where $\Gamma(S)$ denotes the set of nodes in R that have a neighbour in S.

18 Weighted Bipartite Matching



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 - ► This gives $R_S \ge |\Gamma(L_S)|$.
 - ▶ The size of the cut is $|L| |L_S| + |R_S|$.
 - ▶ Using the fact that $|\Gamma(L_S)| \ge L_S$ gives that this is at least |L|.

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We introduce a node weighting \vec{x} . Let for a node $v \in V$, $x_v \in \mathbb{R}$ denote the weight of node v.

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- Try to compute a perfect matching in the subgraph $H(\vec{x})$. If you are successful you found an optimal matching.

Reason:

▶ The weight of your matching M^* is

$$\sum_{(u,v)\in M^*} w_{(u,v)} = \sum_{(u,v)\in M^*} (x_u + x_v) = \sum_v x_v \ .$$

Any other perfect matching M (in G, not necessarily in $H(\vec{x})$) has

$$\sum_{(u,v)\in M} w_{(u,v)} \leq \sum_{(u,v)\in M} (x_u + x_v) = \sum_v x_v \ .$$

What if you don't find a perfect matching?

Then, Halls theorem guarantees you that there is a set $S \subseteq L$, with $|\Gamma(S)| < |S|$, where Γ denotes the neighbourhood w.r.t. the subgraph $H(\vec{x})$.

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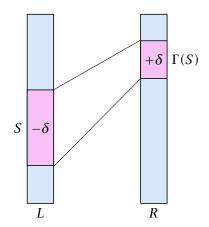
Idea: reweight such that:

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If we can do this we have an algorithm that terminates with an optimal solution (we analyze the running time later).

Changing Node Weights

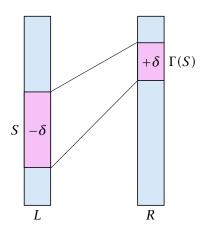
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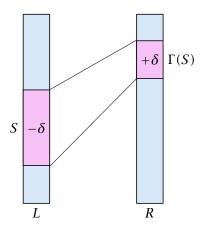
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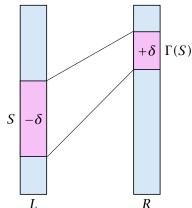
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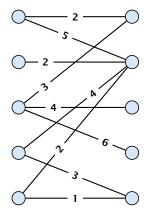


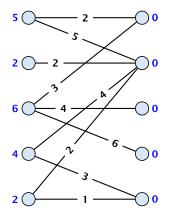
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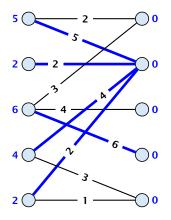
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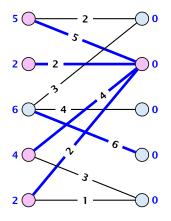
- Total node-weight decreases.
- ▶ Only edges from S to $R \Gamma(S)$ decrease in their weight.
- Since, none of these edges is tight (otw. the edge would be contained in $H(\vec{x})$, and hence would go between S and $\Gamma(S)$) we can do this decrement for small enough $\delta>0$ until a new edge gets tight.



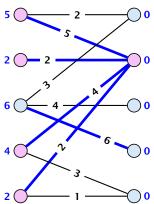


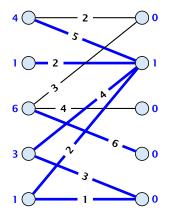


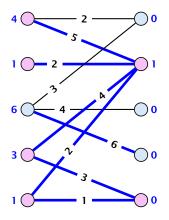




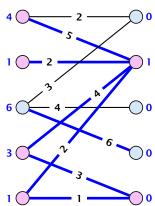


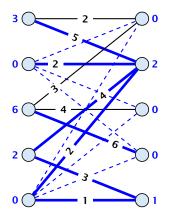


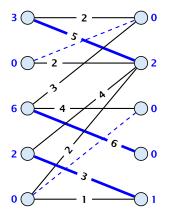


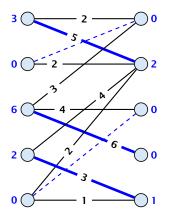












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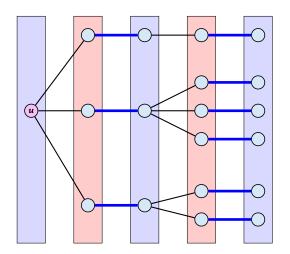
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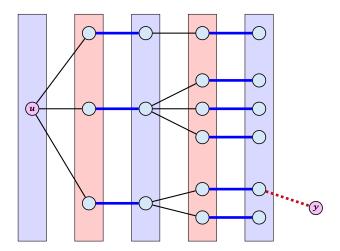
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- ► This matching is still contained in the new graph, because all its edges either go between $\Gamma(S)$ and S or between L-S and $R-\Gamma(S)$.
- Hence, reweighting does not decrease the size of a maximum matching in the tight sub-graph.

- ▶ We will show that after at most *n* reweighting steps the size of the maximum matching can be increased by finding an augmenting path.
- This gives a polynomial running time.

Construct an alternating tree.



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Start on the left and compute an alternating tree, starting at any free node u.

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- The set of even vertices is on the left and the set of odd vertices is on the right and contains all neighbours of even nodes.
- All odd vertices are matched to even vertices. Furthermore, the even vertices additionally contain the free vertex u. Hence, $|V_{\rm odd}| = |\Gamma(V_{\rm even})| < |V_{\rm even}|$, and all odd vertices are saturated in the current matching.

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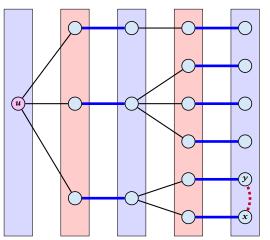
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- A more careful implementation of the algorithm obtains a running time of $\mathcal{O}(n^3)$.

Construct an alternating tree.



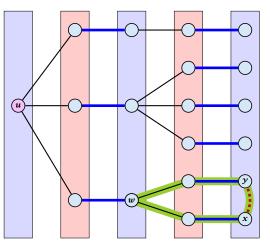
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Case 4:

 \boldsymbol{y} is already contained in T as an even vertex

can't ignore ${m y}$

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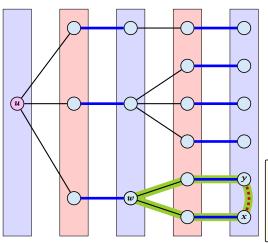
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The cycle $w \leftrightarrow y - x \leftrightarrow w$ is called a blossom. w is called the base of the blossom (even node!!!). The path u-w is called the stem of the blossom.



Flowers and Blossoms

Definition 92

A flower in a graph G=(V,E) w.r.t. a matching M and a (free) root node r, is a subgraph with two components:

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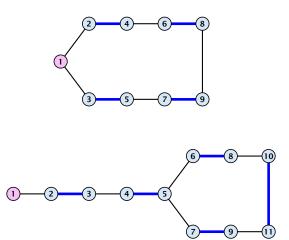
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Flowers and Blossoms

Definition 92

A flower in a graph G = (V, E) w.r.t. a matching M and a (free) root node r, is a subgraph with two components:

- A stem is an even length alternating path that starts at the root node γ and terminates at some node w. We permit the possibility that r = w (empty stem).
- A blossom is an odd length alternating cycle that starts and terminates at the terminal node w of a stem and has no other node in common with the stem. w is called the base of the blossom.



Properties:

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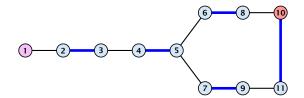
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- **2.** A blossom spans 2k + 1 nodes and contains k matched edges for some integer $k \ge 1$. The matched edges match all nodes of the blossom except the base.
- 3. The base of a blossom is an even node (if the stem is part of an alternating tree starting at r).

Properties:

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- 5. The even alternating path to x terminates with a matched edge and the odd path with an unmatched edge.



When during the alternating tree construction we discover a blossom B we replace the graph G by G' = G/B, which is obtained from G by contracting the blossom B.

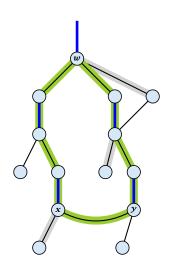
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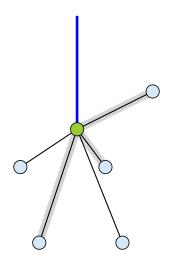
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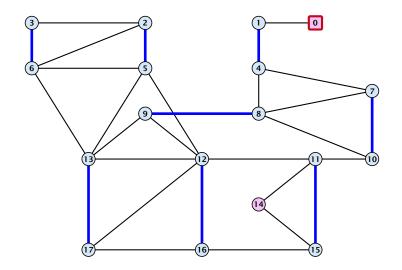
- Delete all vertices in B (and its incident edges) from G.
- Add a new (pseudo-)vertex b. The new vertex b is connected to all vertices in $V \setminus B$ that had at least one edge to a vertex from B.

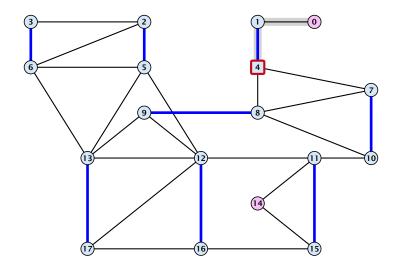
- Edges of T that connect a node u not in B to a node in B become tree edges in T' connecting u to b.
- Matching edges (there is at most one) that connect a node u not in B to a node in B become matching edges in M'.
- Nodes that are connected in G to at least one node in B become connected to b in G'.

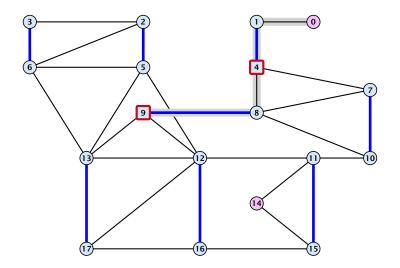


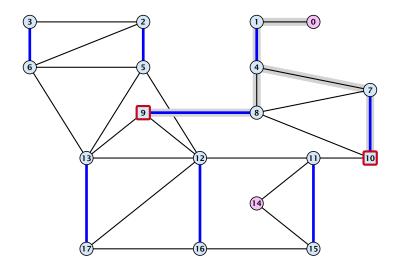
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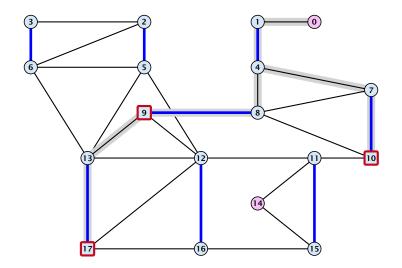


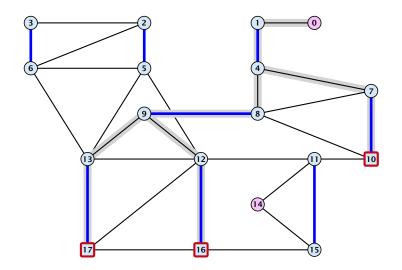


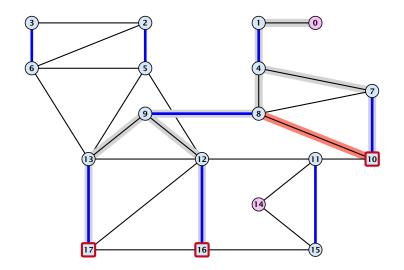


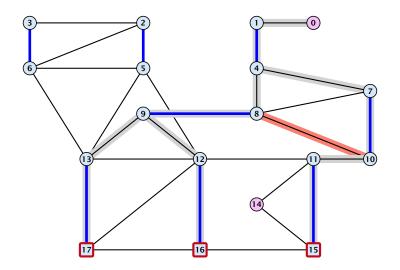


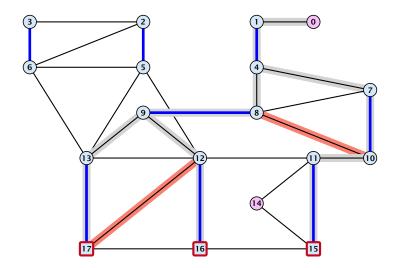


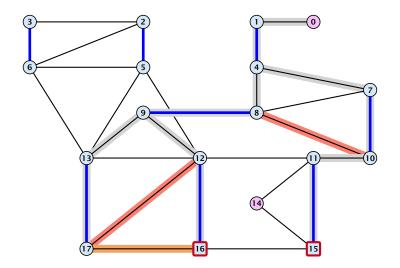


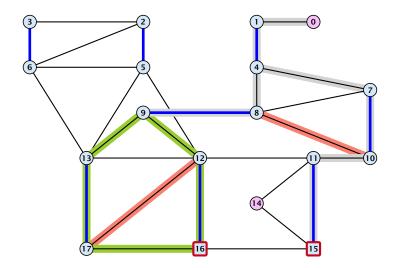


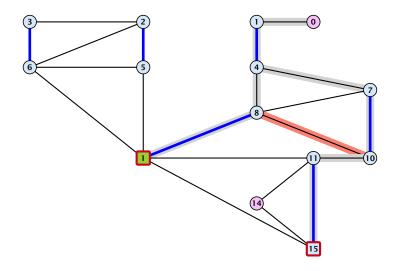


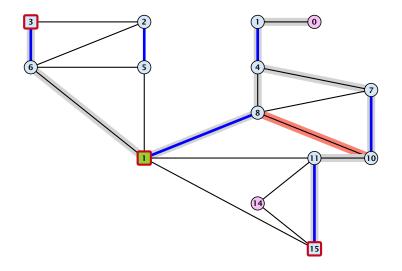


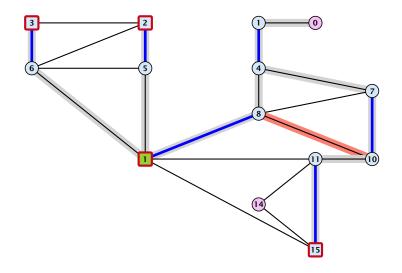


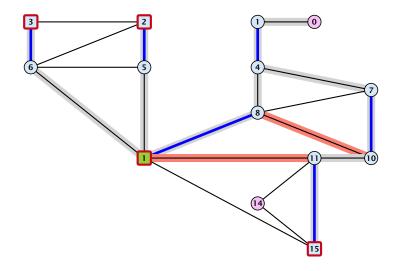


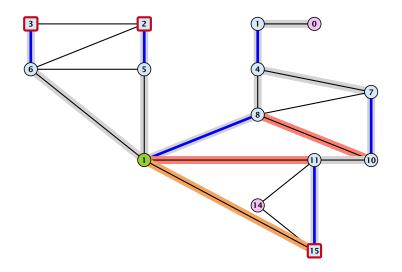


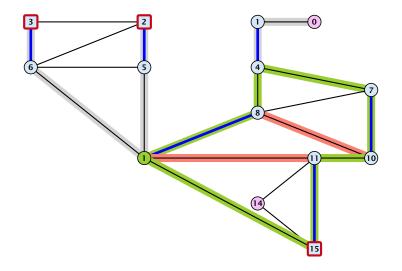


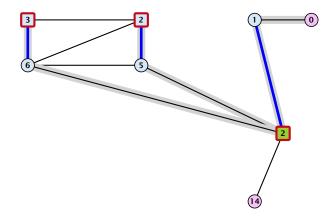


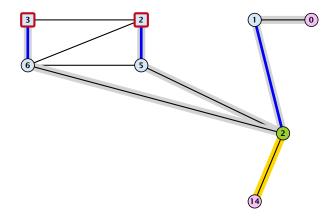


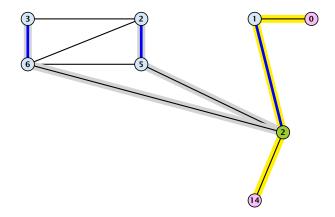


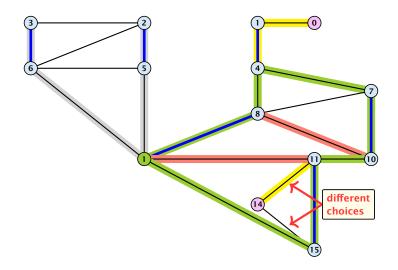


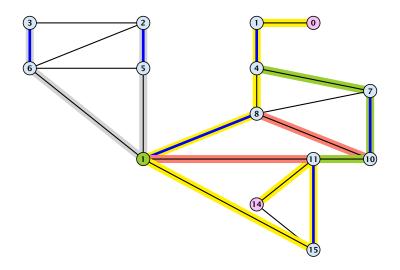


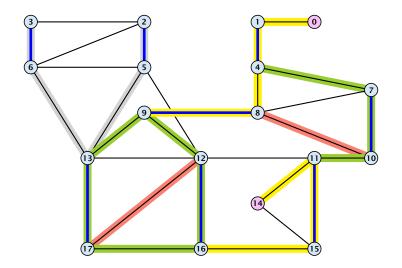


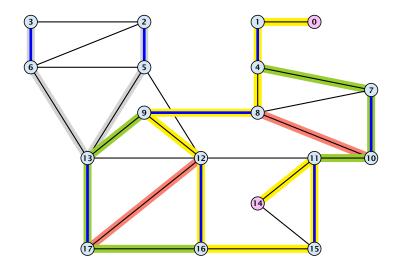




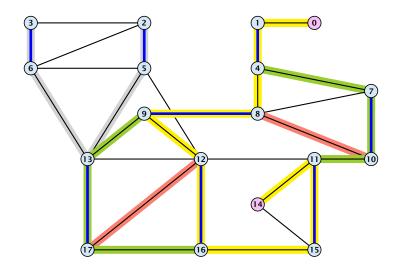








Example: Blossom Algorithm



Assume that in G we have a flower w.r.t. matching M. Let r be the root, B the blossom, and w the base. Let graph G' = G/B with pseudonode b. Let M' be the matching in the contracted graph.

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Lemma 93

If G' contains an augmenting path P' starting at r (or the pseudo-node containing r) w.r.t. the matching M' then G contains an augmenting path starting at r w.r.t. matching M.

Proof.

If P' does not contain b it is also an augmenting path in G.

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Case 1: non-empty stem

Next suppose that the stem is non-empty.

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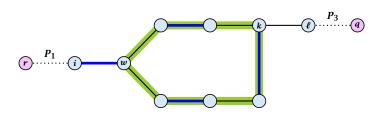
Proof.

If P' does not contain b it is also an augmenting path in G.

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Next suppose that the stem is non-empty.





- After the expansion ℓ must be incident to some node in the blossom. Let this node be k.
- ▶ If $k \neq w$ there is an alternating path P_2 from w to k that ends in a matching edge.
- ▶ $P_1 \circ (i, w) \circ P_2 \circ (k, \ell) \circ P_3$ is an alternating path.
- ▶ If k = w then $P_1 \circ (i, w) \circ (w, \ell) \circ P_3$ is an alternating path.

Proof.

Case 2: empty stem

If the stem is empty then after expanding the blossom, w = r.

Proof.

Case 2: empty stem

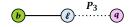
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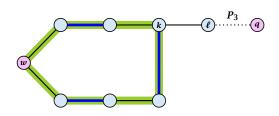


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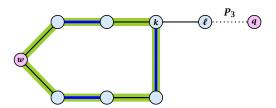


Proof.

Case 2: empty stem

If the stem is empty then after expanding the blossom, w = r.





▶ The path $r \circ P_2 \circ (k, \ell) \circ P_3$ is an alternating path.

Lemma 94

If G contains an augmenting path P from r to q w.r.t. matching M then G' contains an augmenting path from r (or the pseudo-node containing r) to q w.r.t. M'.

Proof.

▶ If *P* does not contain a node from *B* there is nothing to prove.

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Let i be the last node on the path P that is part of the blossom.

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Case 1: empty stem

Let i be the last node on the path P that is part of the blossom.

P is of the form $P_1 \circ (i, j) \circ P_2$, for some node j and (i, j) is unmatched.

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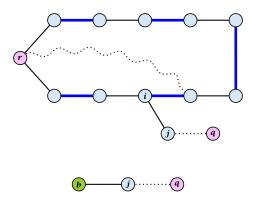
Case 1: empty stem

Let i be the last node on the path P that is part of the blossom.

P is of the form $P_1 \circ (i, j) \circ P_2$, for some node j and (i, j) is unmatched.

 $(b, j) \circ P_2$ is an augmenting path in the contracted network.

Illustration for Case 1:



Case 2: non-empty stem

Case 2: non-empty stem

Let P_3 be alternating path from r to w; this exists because r and w are root and base of a blossom. Define $M_+ = M \oplus P_3$.

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In M_+ , γ is matched and w is unmatched.

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In M_+ , γ is matched and w is unmatched.

G must contain an augmenting path w.r.t. matching M_+ , since M and M_+ have same cardinality.

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In M_+ , r is matched and w is unmatched.

G must contain an augmenting path w.r.t. matching M_+ , since M and M_+ have same cardinality.

This path must go between w and q as these are the only unmatched vertices w.r.t. $M_{\rm +}$.

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For M'_+ the blossom has an empty stem. Case 1 applies.

Case 2: non-empty stem

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For M'_+ the blossom has an empty stem. Case 1 applies.

G' has an augmenting path w.r.t. M'_+ . It must also have an augmenting path w.r.t. M', as both matchings have the same cardinality.

Case 2: non-empty stem

Let P_3 be alternating path from r to w; this exists because r and w are root and base of a blossom. Define $M_+ = M \oplus P_3$.

In M_+ , γ is matched and w is unmatched.

G must contain an augmenting path w.r.t. matching M_+ , since M and M_+ have same cardinality.

This path must go between w and q as these are the only unmatched vertices w.r.t. M_{+} .

For M'_+ the blossom has an empty stem. Case 1 applies.

G' has an augmenting path w.r.t. M'_+ . It must also have an augmenting path w.r.t. M', as both matchings have the same cardinality.

This path must go between r and q.

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
 - 2: found ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$
- 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(*i*, *found*)
- 8: **if** *found* = true **then return**

Search for an augmenting path starting at r.

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
- 2: found ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$ 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(*i*, *found*)
- 8: **if** *found* = true **then return**

A(i) contains neighbours of node i. We create a copy $\bar{A}(i)$ so that we later can shrink blossoms.

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
- 2: found ← false

8:

- unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$
- 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(*i*, *found*)
 - if found = true then return

found is just a Boolean that allows to abort the search process...

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
- 2: *found* ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$
- 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(i, found)
- 8: **if** *found* = true **then return**

In the beginning no node is in the tree.

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
 - 2: *found* ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$
- 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(i, found)
- 8: **if** *found* = true **then return**

Put the root in the tree.

list could also be a set or a stack.

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
- 2: *found* ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$ 5: while $list \neq \emptyset$ do
- 6: delete a node *i* from *list*
- 7: examine(i, found)
- 8: **if** *found* = true **then return**

As long as there are nodes with unexamined neighbours...

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
 - 2: found ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$
- 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(i, found)
- 8: **if** found = true **then return**

...examine the next one

- 1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i
- 2: *found* ← false
- 3: unlabel all nodes;
- 4: give an even label to r and initialize $list \leftarrow \{r\}$
- 5: while $list \neq \emptyset$ do
- 6: delete a node i from list
- 7: examine(i, found)
- 8: **if** found = true **then return**

If you found augmenting path abort and start from next root.

Algorithm 51 examine(*i*, *found*) 1: for all $j \in \bar{A}(i)$ do if j is even then contract(i, j) and return 2: **if** j is unmatched **then** 3: 4: $q \leftarrow j$; $pred(q) \leftarrow i$; 5: 6: *found* ← true; 7: return if j is matched and unlabeled then 8: $pred(j) \leftarrow i$; 9: 10: $pred(mate(j)) \leftarrow j$; add mate(j) to *list* 11:

```
Algorithm 51 examine(i, found)
1: for all j \in \bar{A}(i) do
        if j is even then contract(i, j) and return
 2:
      if j is unmatched then
3:
4:
             q \leftarrow j;
             pred(q) \leftarrow i;
5:
             found ← true;
6:
7:
             return
        if j is matched and unlabeled then
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             pred(j) \leftarrow i;
9:
             pred(mate(j)) \leftarrow j;
10:
             add mate(j) to list
11:
```

For all neighbours j do...

```
Algorithm 51 examine(i, found)
1: for all j \in \bar{A}(i) do
        if j is even then contract(i, j) and return
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```

You have found a blossom...

Algorithm 51 examine(*i*, *found*) 1: for all $j \in \bar{A}(i)$ do if j is even then contract(i, j) and return 2: if i is unmatched then 3: 4: $q \leftarrow j$; $pred(q) \leftarrow i$; 5: 6: *found* ← true; 7: return if j is matched and unlabeled then 8: 9: $pred(j) \leftarrow i$; 10: $pred(mate(j)) \leftarrow j$; add mate(j) to *list* 11:

You have found a free node which gives you an augmenting path.

```
Algorithm 51 examine(i, found)
1: for all j \in \bar{A}(i) do
        if j is even then contract(i, j) and return
 2:
      if j is unmatched then
3:
4:
             q \leftarrow j;
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5:
6:
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7:
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8:
         if i is matched and unlabeled then
9:
             pred(j) \leftarrow i;
10:
             pred(mate(j)) \leftarrow j;
             add mate(j) to list
11:
```

If you find a matched node that is not in the tree you grow...

```
Algorithm 51 examine(i, found)
1: for all j \in \bar{A}(i) do
        if j is even then contract(i, j) and return
 2:
      if j is unmatched then
3:
4:
             q \leftarrow j;
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5:
6:
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        if j is matched and unlabeled then
8:
             pred(j) \leftarrow i;
9:
10:
             pred(mate(j)) \leftarrow j;
             add mate(j) to list
11:
```

mate(j) is a new node from
which you can grow further.

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label *b* even and add to *list*
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Contract blossom identified by nodes *i* and *j*

- 1: trace pred-indices of i and j to identify a blossom B
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Get all nodes of the blossom.

Time: $\mathcal{O}(m)$

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
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- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Identify all neighbours of b.

Time: $\mathcal{O}(m)$ (how?)

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

b will be an even node, and it has unexamined neighbours.

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label *b* even and add to *list*
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Every node that was adjacent to a node in B is now adjacent to b

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
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- 5: form a circular double linked list of nodes in B
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Only for making a blossom expansion easier.

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in *B* from the graph

Only delete links from nodes not in B to B.

When expanding the blossom again we can recreate these links in time O(m).

A contraction operation can be performed in time $\mathcal{O}(m)$. Note, that any graph created will have at most m edges.

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- The time between two contraction-operation is basically a BFS/DFS on a graph. Hence takes time $\mathcal{O}(m)$.

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- ► There are at most *n* contractions as each contraction reduces the number of vertices.

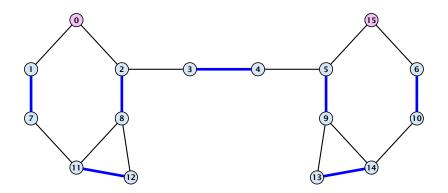
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- The time between two contraction-operation is basically a BFS/DFS on a graph. Hence takes time $\mathcal{O}(m)$.
- ► There are at most *n* contractions as each contraction reduces the number of vertices.
- ► The expansion can trivially be done in the same time as needed for all contractions.

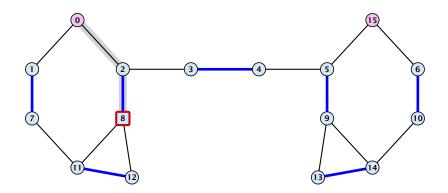
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- The expansion can trivially be done in the same time as needed for all contractions.
- An augmentation requires time $\mathcal{O}(n)$. There are at most n of them.

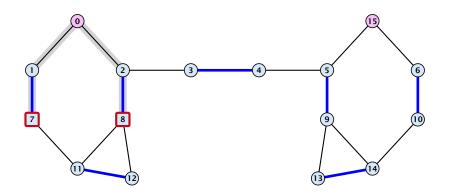
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- ► There are at most *n* contractions as each contraction reduces the number of vertices.
- The expansion can trivially be done in the same time as needed for all contractions.
- An augmentation requires time $\mathcal{O}(n)$. There are at most n of them.
- In total the running time is at most

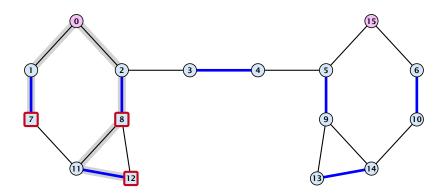
$$n \cdot (\mathcal{O}(mn) + \mathcal{O}(n)) = \mathcal{O}(mn^2)$$
.

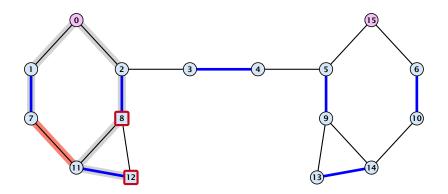


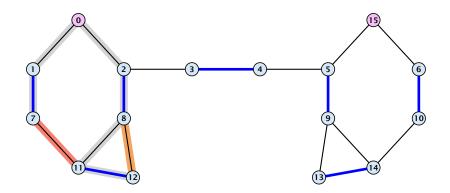


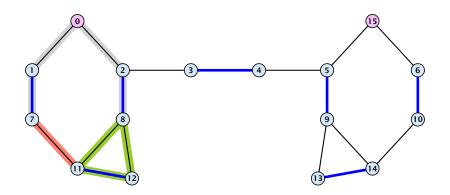


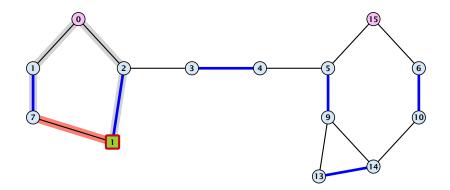


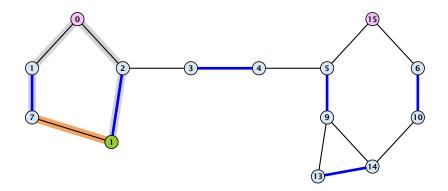


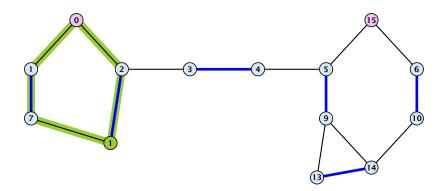


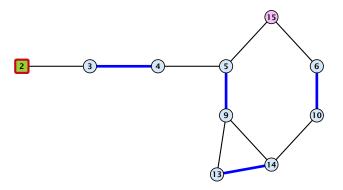


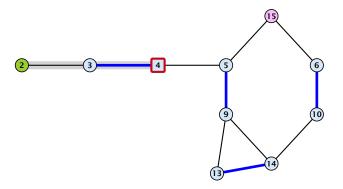


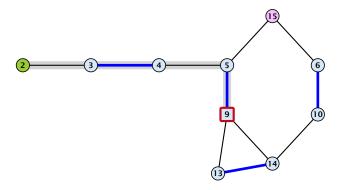


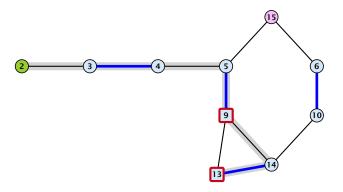


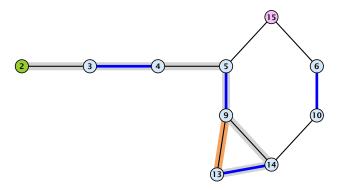


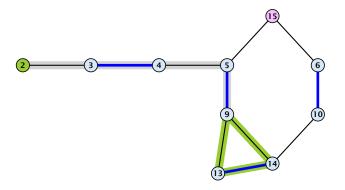


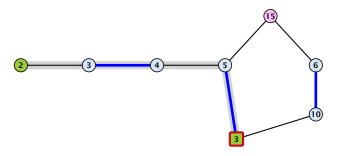


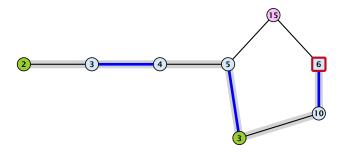


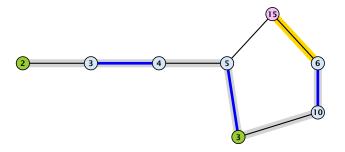


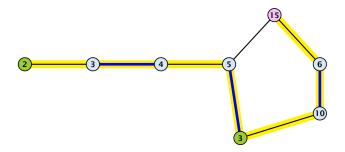


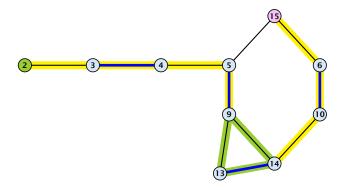


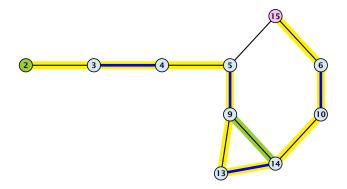


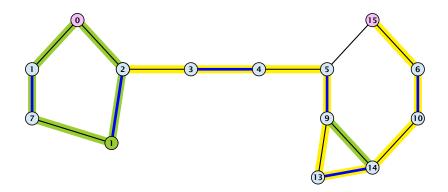


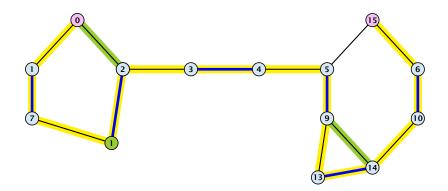


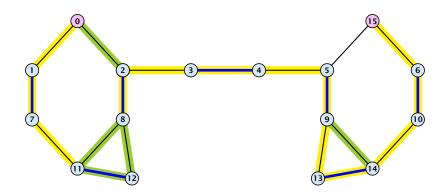


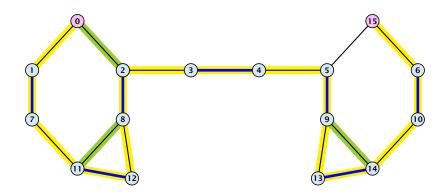


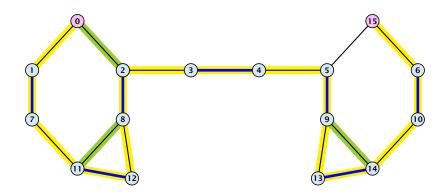












A Fast Matching Algorithm

```
Algorithm 53 Bimatch-Hopcroft-Karp(G)

1: M \leftarrow \varnothing

2: repeat

3: let \mathcal{P} = \{P_1, \dots, P_k\} be maximal set of

4: vertex-disjoint, shortest augmenting path w.r.t. M.

5: M \leftarrow M \oplus (P_1 \cup \cdots \cup P_k)

6: until \mathcal{P} = \varnothing

7: return M
```

We call one iteration of the repeat-loop a phase of the algorithm.

Lemma 95

Given a matching M and a matching M^* with $|M^*| - |M| \ge 0$. There exist $|M^*| - |M|$ vertex-disjoint augmenting path w.r.t. M.

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Similar to the proof that a matching is optimal iff it does not contain an augmenting path.

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- Consider the graph $G = (V, M \oplus M^*)$, and mark edges in this graph blue if they are in M and red if they are in M^* .

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- Consider the graph $G = (V, M \oplus M^*)$, and mark edges in this graph blue if they are in M and red if they are in M^* .
- ▶ The connected components of *G* are cycles and paths.
- ▶ The graph contains $k ext{ \leq } |M^*| |M|$ more red edges than blue edges.
- ▶ Hence, there are at least *k* components that form a path starting and ending with a red edge. These are augmenting paths w.r.t. *M*.

Let P_1, \ldots, P_k be a maximal collection of vertex-disjoint, shortest augmenting paths w.r.t. M (let $\ell = |P_i|$).

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- Let P be an augmenting path in M'.

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Lemma 96

The set $A \stackrel{\text{def}}{=} M \oplus (M' \oplus P) = (P_1 \cup \cdots \cup P_k) \oplus P$ contains at least $(k+1)\ell$ edges.

Proof.

► The set describes exactly the symmetric difference between matchings M and $M' \oplus P$.

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- **Each** of these paths is of length at least ℓ .

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P is of length at least $\ell+1$. This shows that the length of a shortest augmenting path increases between two phases of the Hopcroft-Karp algorithm.

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If P does not intersect any of the P_1, \ldots, P_k , this follows from the maximality of the set $\{P_1, \ldots, P_k\}$.

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- If P does not intersect any of the P_1, \ldots, P_k , this follows from the maximality of the set $\{P_1, \ldots, P_k\}$.
- ▶ Otherwise, at least one edge from P coincides with an edge from paths $\{P_1, \ldots, P_k\}$.

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- ▶ Otherwise, at least one edge from P coincides with an edge from paths $\{P_1, \ldots, P_k\}$.
- This edge is not contained in A.
- ▶ Hence, $|A| \le k\ell + |P| 1$.
- ► The lower bound on |A| gives $(k+1)\ell \le |A| \le k\ell + |P| 1$, and hence $|P| \ge \ell + 1$.

If the shortest augmenting path w.r.t. a matching M has ℓ edges then the cardinality of the maximum matching is of size at most $|M| + \frac{|V|}{\ell+1}$.

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Proof.

The symmetric difference between M and M^* contains $|M^*| - |M|$ vertex-disjoint augmenting paths. Each of these paths contains at least $\ell+1$ vertices. Hence, there can be at most $\frac{|V|}{\ell+1}$ of them.

Lemma 98

The Hopcroft-Karp algorithm requires at most $2\sqrt{|V|}$ phases.

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- ▶ After iteration $\lfloor \sqrt{|V|} \rfloor$ the length of a shortest augmenting path must be at least $\lfloor \sqrt{|V|} \rfloor + 1 \ge \sqrt{|V|}$.
- ► Hence, there can be at most $|V|/(\sqrt{|V|}+1) \le \sqrt{|V|}$ additional augmentations.

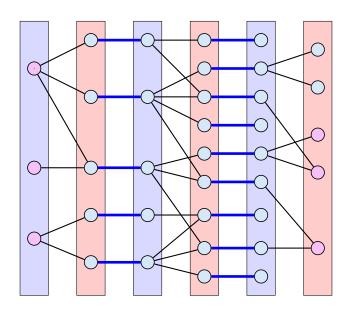
Lemma 99

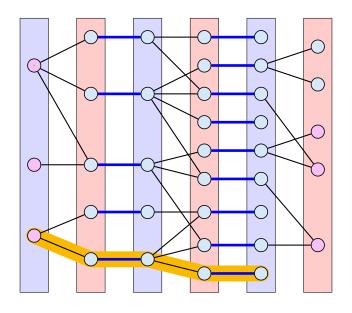
One phase of the Hopcroft-Karp algorithm can be implemented in time O(m).

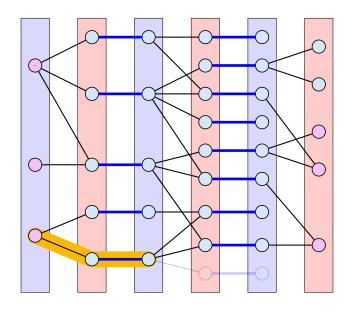
construct a "level graph" G':

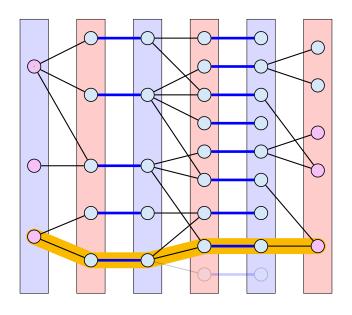
- construct Level 0 that includes all free vertices on left side L
- construct Level 1 containing all neighbors of Level 0
- construct Level 2 containing matching neighbors of Level 1
- construct Level 3 containing all neighbors of Level 2
- **.**...
- > stop when a level (apart from Level 0) contains a free vertex can be done in time $\mathcal{O}(m)$ by a modified BFS

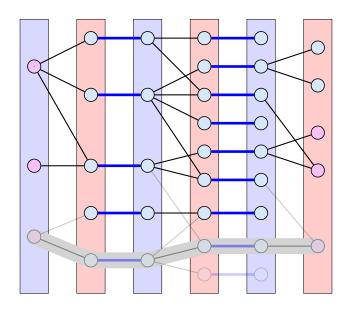
- a shortest augmenting path must go from Level 0 to the last layer constructed
- it can only use edges between layers
- construct a maximal set of vertex disjoint augmenting path connecting the layers
- for this, go forward until you either reach a free vertex or you reach a "dead end" \boldsymbol{v}
- if you reach a free vertex delete the augmenting path and all incident edges from the graph
- if you reach a dead end backtrack and delete \boldsymbol{v} together with its incident edges

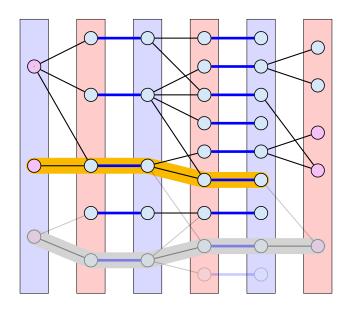


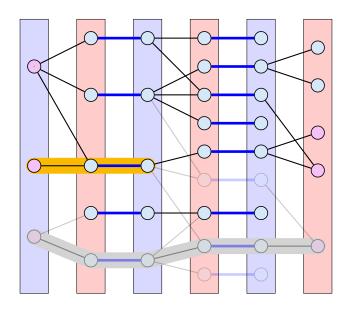


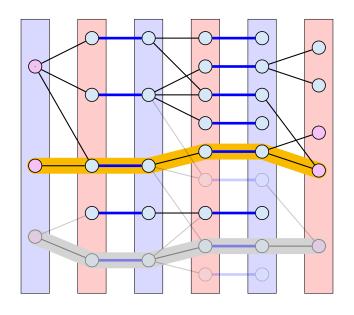


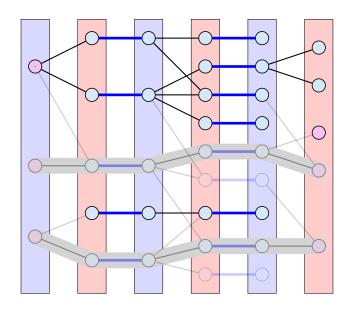


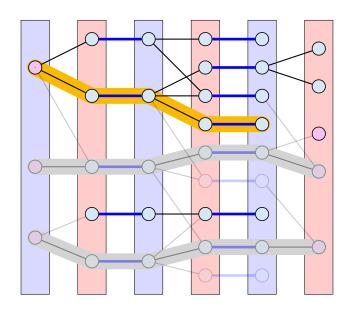


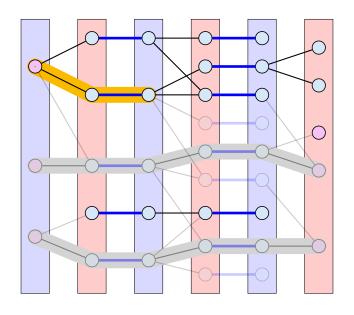


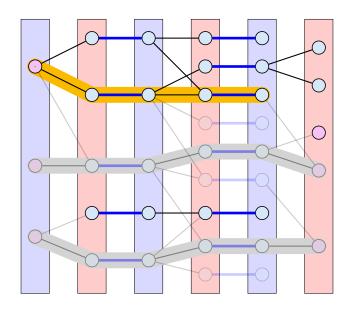


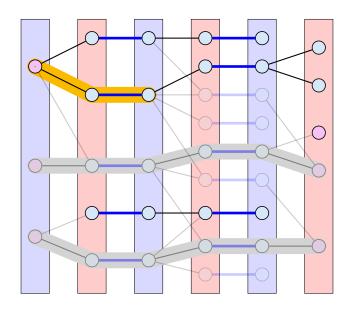


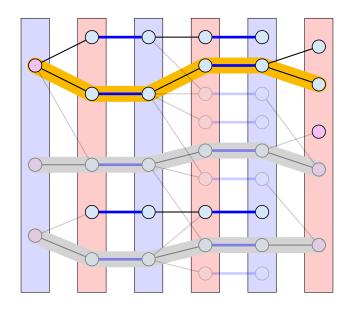


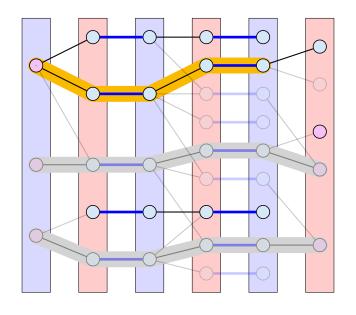


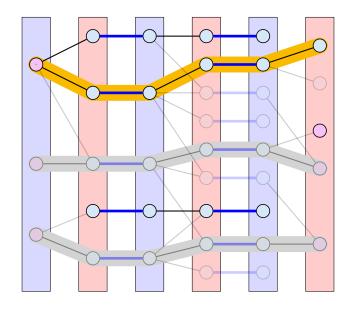


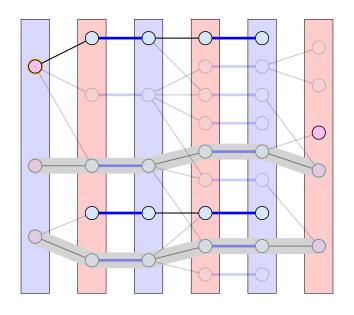


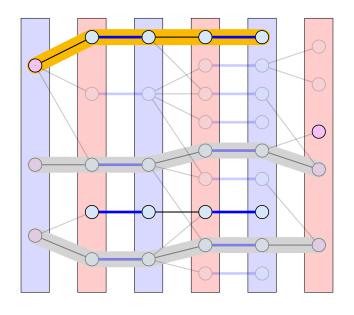


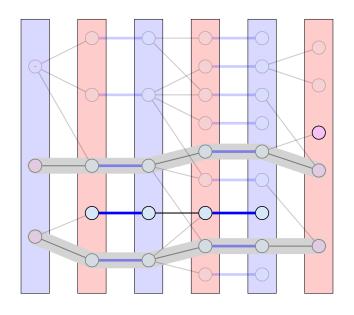


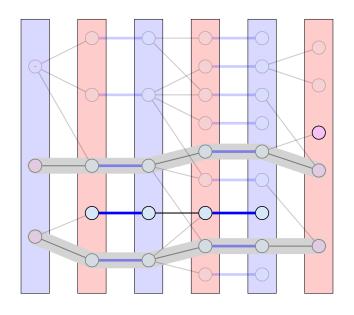












Analysis: Shortest Augmenting Path for Flows

cost for searches during a phase is O(mn)

- ightharpoonup a search (successful or unsuccessful) takes time O(n)
- a search deletes at least one edge from the level graph

there are at most n phases

Time: $\mathcal{O}(mn^2)$.

Analysis for Unit-capacity Simple Networks

cost for searches during a phase is O(m)

an edge/vertex is traversed at most twice

need at most $\mathcal{O}(\sqrt{n})$ phases

- after \sqrt{n} phases there is a cut of size at most \sqrt{n} in the residual graph
- lacktriangle hence at most \sqrt{n} additional augmentations required

Time: $\mathcal{O}(m\sqrt{n})$.

21 Gomory Hu Trees

Given an undirected, weighted graph G = (V, E, c) a cut-tree T = (V, F, w) is a tree with edge-set F and capacities w that fulfills the following properties.

- **1. Equivalent Flow Tree:** For any pair of vertices $s, t \in V$, f(s,t) in G is equal to $f_T(s,t)$.
- **2. Cut Property:** A minimum *s-t* cut in *T* is also a minimum cut in *G*.

Here, f(s,t) is the value of a maximum s-t flow in G, and $f_T(s,t)$ is the corresponding value in T.

The algorithm maintains a partition of V, (sets S_1, \ldots, S_t), and a spanning tree T on the vertex set $\{S_1, \ldots, S_t\}$.

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In each such split-operation it chooses a set S_i with $|S_i| \ge 2$ and splits this set into two non-empty parts X and Y.

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- ▶ In each such split-operation it chooses a set S_i with $|S_i| \ge 2$ and splits this set into two non-empty parts X and Y.
- ▶ S_i is then removed from T and replaced by X and Y.

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- ➤ X and Y are connected by an edge, and the edges that before the split were incident to S_i are attached to either X or Y.

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- ➤ *X* and *Y* are connected by an edge, and the edges that before the split were incident to *S*_i are attached to either *X* or *Y*.

In the end this gives a tree on the vertex set V.



▶ Select S_i that contains at least two nodes a and b.

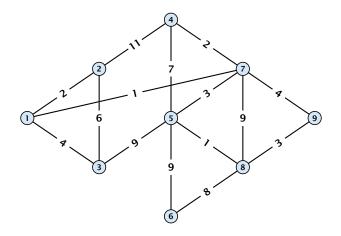
- Select S_i that contains at least two nodes a and b.
- Compute the connected components of the forest obtained from the current tree T after deleting S_i . Each of these components corresponds to a set of vertices from V.

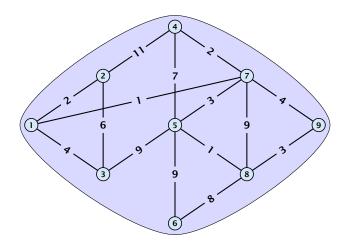
- Select S_i that contains at least two nodes a and b.
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- ► Consider the graph *H* obtained from *G* by contracting these connected components into single nodes.

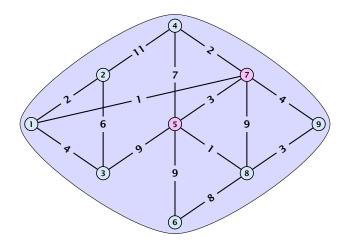
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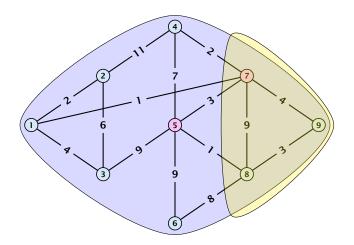
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- ▶ Split S_i in T into two sets/nodes $S_i^a = S_i \cap A$ and $S_i^b = S_i \cap B$ and add edge $\{S_i^a, S_i^b\}$ with capacity $f_H(a, b)$.

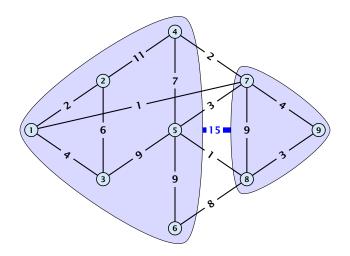
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- ▶ Replace an edge $\{S_i, S_x\}$ by $\{S_i^a, S_x\}$ if $S_x \subset A$ and by $\{S_i^b, S_x\}$ if $S_x \subset B$.

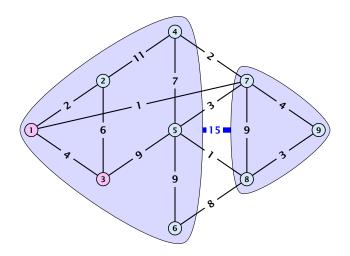


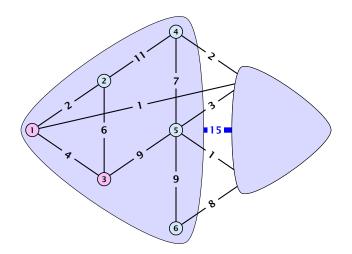


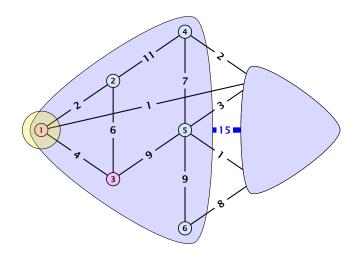


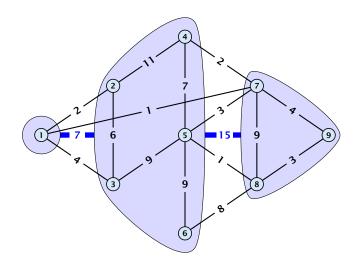


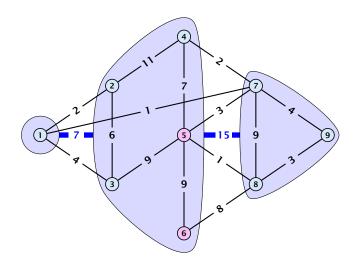


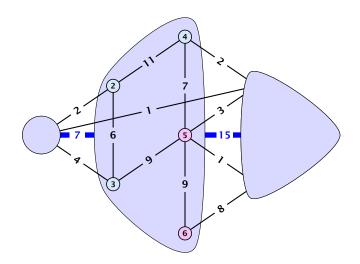


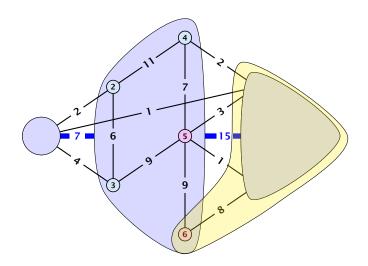


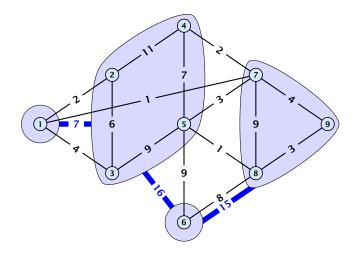


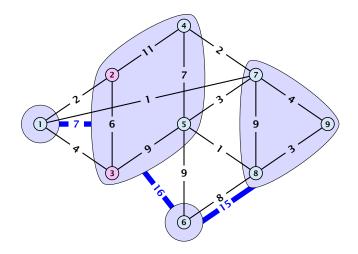


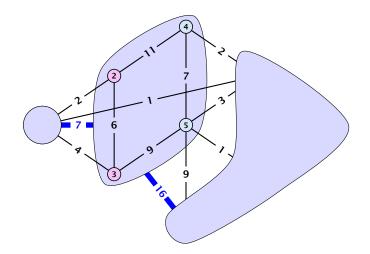


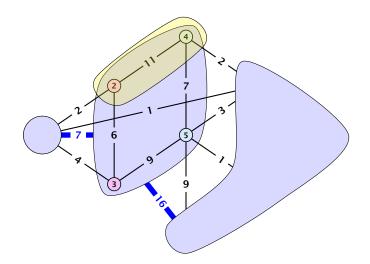


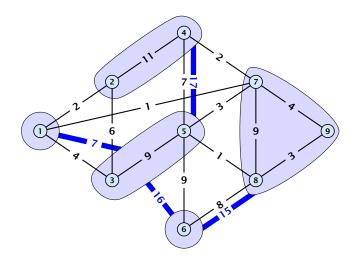


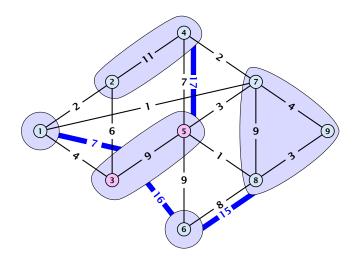


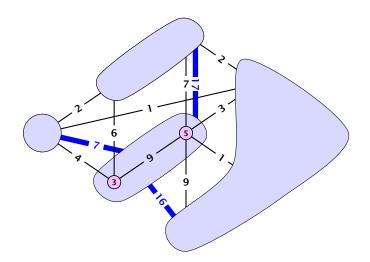


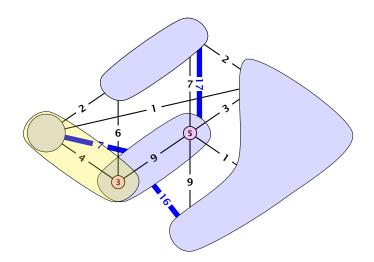


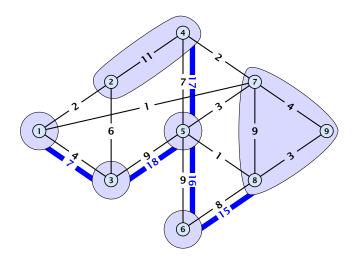


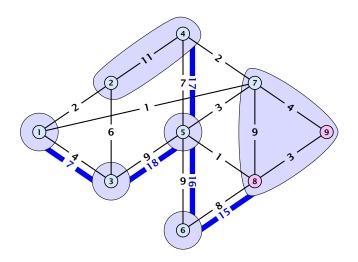


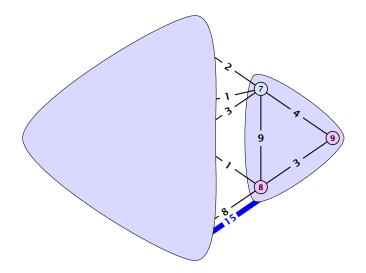


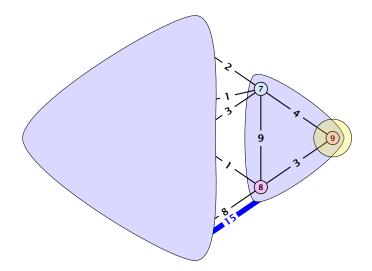


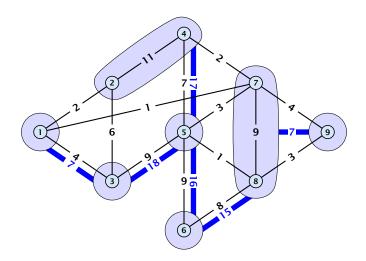


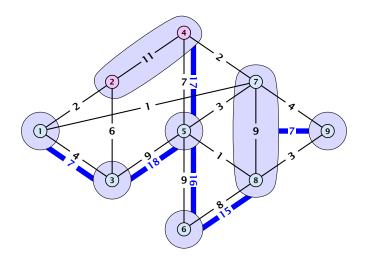


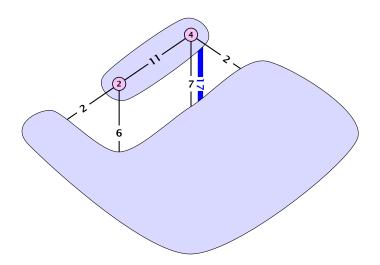


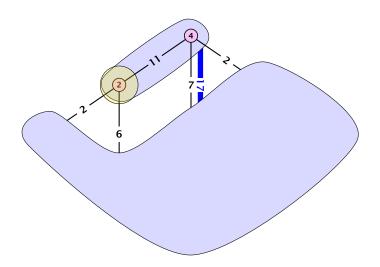


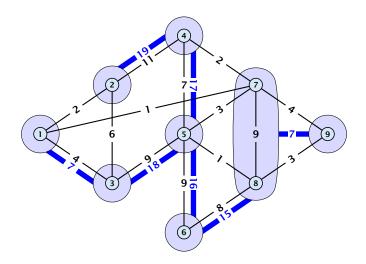


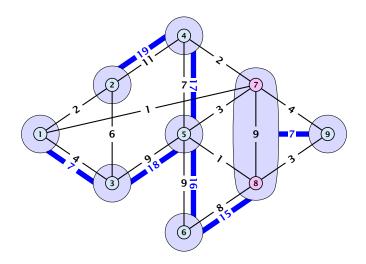


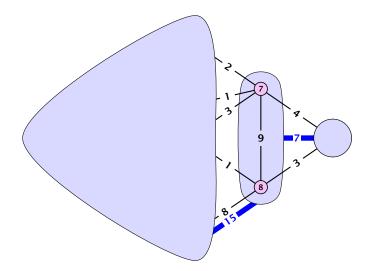


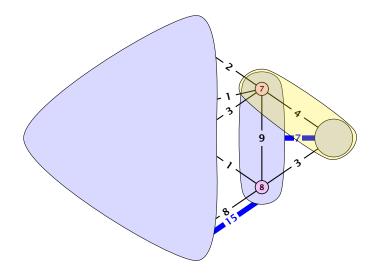


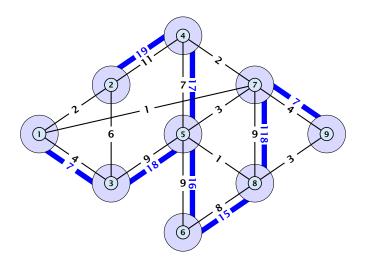












Analysis

Lemma 100

For nodes $s, t, x \in V$ we have $f(s, t) \ge \min\{f(s, x), f(x, t)\}$

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Lemma 101

For nodes $s, t, x_1, \dots, x_k \in V$ we have

$$f(s,t) \ge \min\{f(s,x_1), f(x_1,x_2), \dots, f(x_{k-1},x_k), f(x_k,t)\}$$

Lemma 102

Let S be some minimum r-s cut for some nodes $r, s \in V$ ($s \in S$), and let $v, w \in S$. Then there is a minimum v-w-cut T with $T \subset S$.

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Proof: Let X be a minimum v-w cut with $X \cap S \neq \emptyset$ and $X \cap (V \setminus S) \neq \emptyset$.

Let S be some minimum r-s cut for some nodes $r, s \in V$ ($s \in S$), and let $v, w \in S$. Then there is a minimum v-w-cut T with $T \subset S$.

Proof: Let X be a minimum $v \cdot w$ cut with $X \cap S \neq \emptyset$ and $X \cap (V \setminus S) \neq \emptyset$. Note that $S \setminus X$ and $S \cap X$ are $v \cdot w$ cuts inside S.

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First case $r \in X$.

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- ▶ $cap(X \setminus S) \ge cap(S)$ because $X \setminus S$ is an r-s cut.

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- ▶ $cap(X \cup S) \ge cap(S)$ because $X \cup S$ is an r-s cut.

Let S be some minimum r-s cut for some nodes r, $s \in V$ ($s \in S$), and let v, $w \in S$. Then there is a minimum v-w-cut T with $T \subset S$.

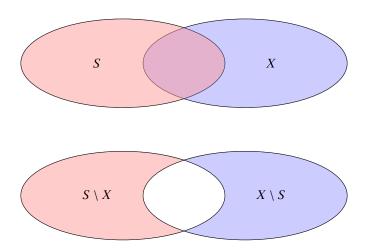
Proof: Let X be a minimum v-w cut with $X \cap S \neq \emptyset$ and $X \cap (V \setminus S) \neq \emptyset$. Note that $S \setminus X$ and $S \cap X$ are v-w cuts inside S. We may assume w.l.o.g. $S \in X$.

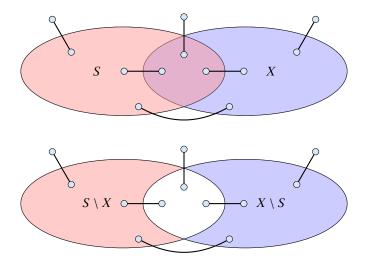
First case $r \in X$.

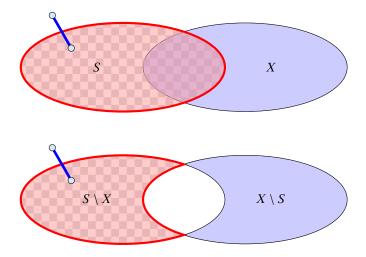
- ▶ $cap(X \setminus S) \ge cap(S)$ because $X \setminus S$ is an r-s cut.
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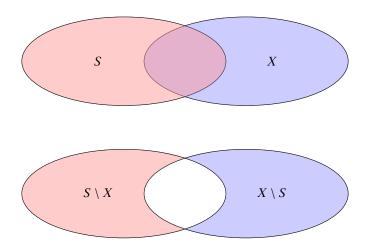
Second case $r \notin X$.

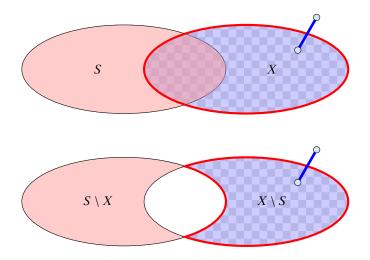
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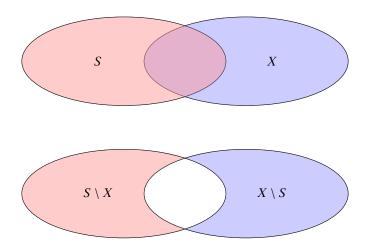


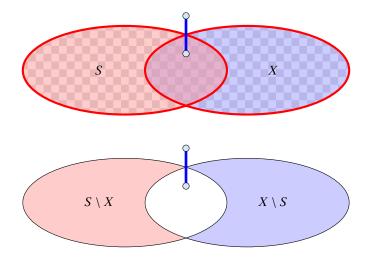


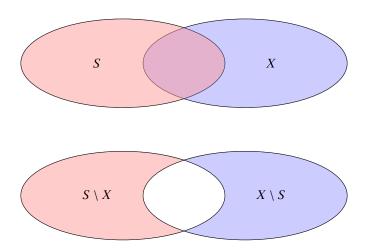


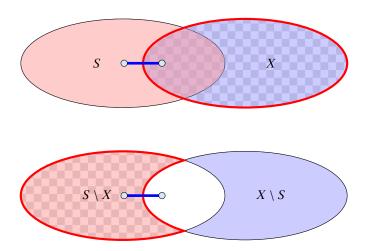


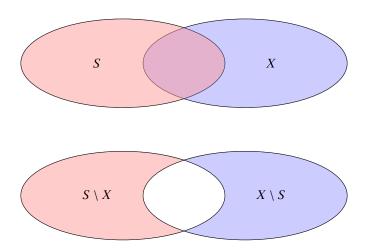


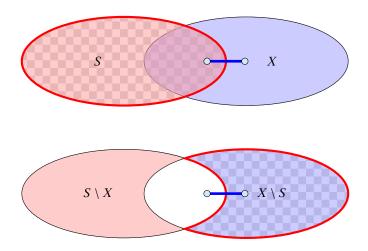


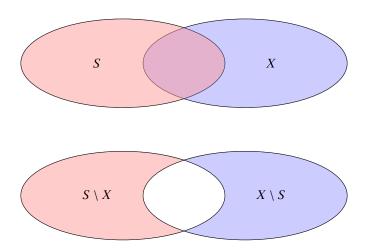


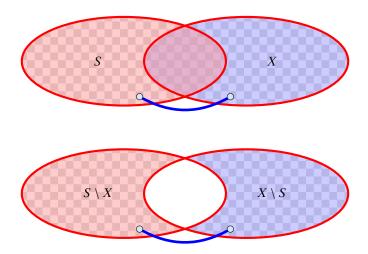


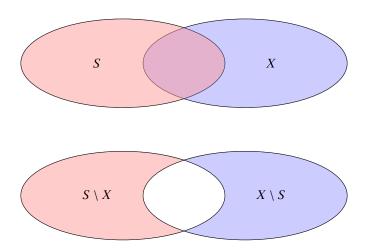


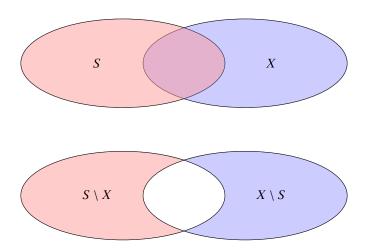


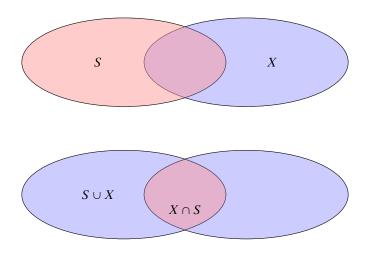


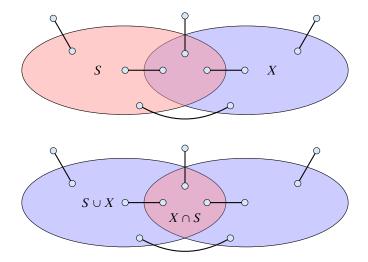


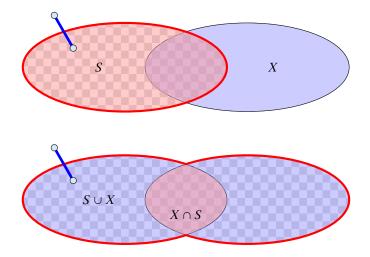


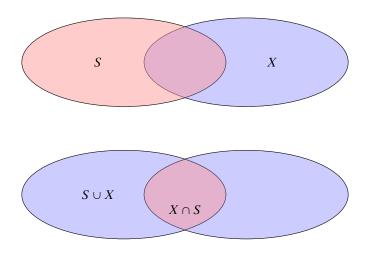


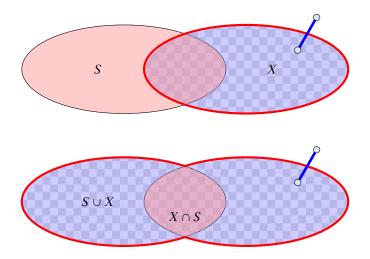


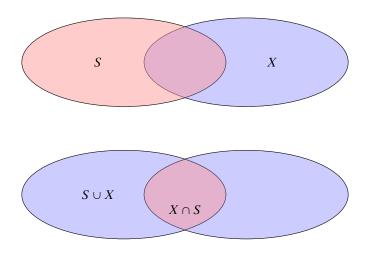


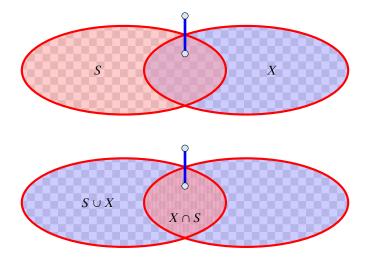


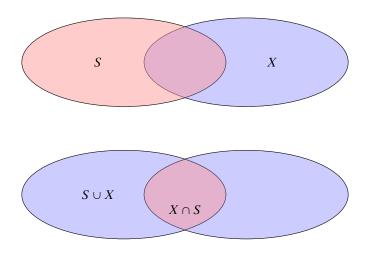


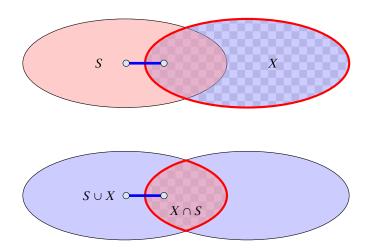


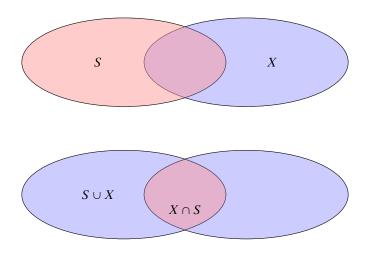


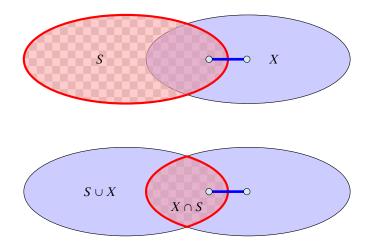


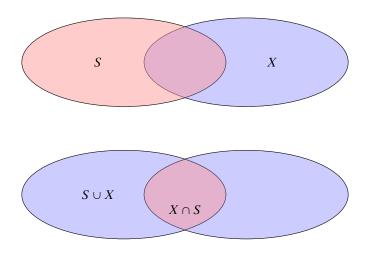


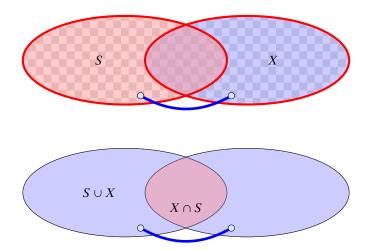


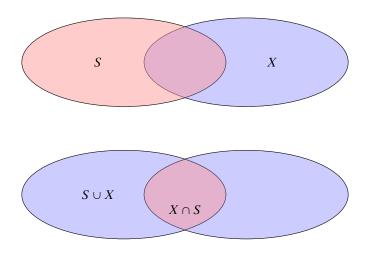


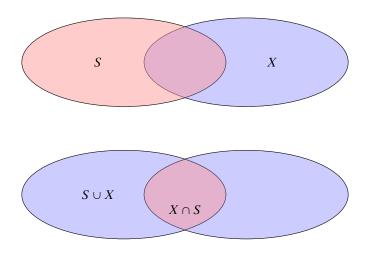












Lemma 102 tells us that if we have a graph G = (V, E) and we contract a subset $X \subset V$ that corresponds to some mincut, then the value of f(s,t) does not change for two nodes $s,t \notin X$.

We will show (later) that the connected components that we contract during a split-operation each correspond to some mincut and, hence, $f_H(s,t)=f(s,t)$, where $f_H(s,t)$ is the value of a minimum s-t mincut in graph H.

Invariant [existence of representatives]:

For any edge $\{S_i, S_j\}$ in T, there are vertices $a \in S_i$ and $b \in S_j$ such that $w(S_i, S_j) = f(a, b)$ and the cut defined by edge $\{S_i, S_j\}$ is a minimum a-b cut in G.

We first show that the invariant implies that at the end of the algorithm T is indeed a cut-tree.

Let $s = x_0, x_1, \dots, x_{k-1}, x_k = t$ be the unique simple path from s to t in the final tree T. From the invariant we get that $f(x_i, x_{i+1}) = w(x_i, x_{i+1})$ for all j.

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$$f_T(s,t)$$

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- Then

$$\begin{split} f_T(s,t) &= \min_{i \in \{0,\dots,k-1\}} \{w(x_i,x_{i+1})\} \\ &= \min_{i \in \{0,\dots,k-1\}} \{f(x_i,x_{i+1})\} \leq f(s,t) \ . \end{split}$$

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- Let $\{x_i, x_{i+1}\}$ be the edge with minimum weight on the path.
- Since by the invariant this edge induces an s-t cut with capacity $f(x_i, x_{i+1})$ we get $f(s, t) \le f(x_i, x_{i+1}) = f_T(s, t)$.

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► Hence, $f_T(s,t) = f(s,t)$ (flow equivalence).

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- ▶ By invariant, it forms a cut with capacity $f(x_j, x_{j+1})$ in G (which separates s and t).

- ► Hence, $f_T(s,t) = f(s,t)$ (flow equivalence).
- ▶ The edge $\{x_j, x_{j+1}\}$ is a mincut between s and t in T.
- ▶ By invariant, it forms a cut with capacity $f(x_j, x_{j+1})$ in G (which separates s and t).
- Since, we can send a flow of value $f(x_j, x_{j+1})$ btw. s and t, this is an s-t mincut (cut property).

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Now, we show that it holds after a split-operation provided that it was true before the operation.

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Let S_i denote our selected cluster with nodes a and b. Because of the invariant all edges leaving $\{S_i\}$ in T correspond to some mincuts.

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Now, we show that it holds after a split-operation provided that it was true before the operation.

Let S_i denote our selected cluster with nodes a and b. Because of the invariant all edges leaving $\{S_i\}$ in T correspond to some mincuts.

Therefore, contracting the connected components does not change the mincut btw. a and b due to Lemma 102.

The invariant obviously holds at the beginning of the algorithm.

Now, we show that it holds after a split-operation provided that it was true before the operation.

Let S_i denote our selected cluster with nodes a and b. Because of the invariant all edges leaving $\{S_i\}$ in T correspond to some mincuts.

Therefore, contracting the connected components does not change the mincut btw. a and b due to Lemma 102.

After the split we have to choose representatives for all edges. For the new edge $\{S_i^a, S_i^b\}$ with capacity $w(S_i^a, S_i^b) = f_H(a,b)$ we can simply choose a and b as representatives.

For edges that are not incident to S_i we do not need to change representatives as the neighbouring sets do not change.

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If $s \in S_i^a$ we can keep x and s as representatives.

Otherwise, we choose x and a as representatives. We need to show that f(x,a) = f(x,s).

Because the invariant was true before the split we know that the edge $\{X, S_i\}$ induces a cut in G of capacity f(x, s). Since, x and a are on opposite sides of this cut, we know that $f(x, a) \le f(x, s)$.

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The set B forms a mincut separating a from b. Contracting all nodes in this set gives a new graph G' where the set B is represented by node v_B . Because of Lemma 102 we know that f'(x,a) = f(x,a) as $x, a \notin B$.

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Since $s \in B$ we have $f'(v_B, x) \ge f(s, x)$.

Also, $f'(a, v_B) \ge f(a, b) \ge f(x, s)$ since the a-b cut that splits S_i into S_i^a and S_i^b also separates s and x.

