Introduction to Optimization Lectures 6 and 7: Discrete Optimization

October 19, 2018 and October 26, 2018 TC2 - Optimisation Université Paris-Saclay, Orsay, France



Dimo Brockhoff Inria Saclay – Ile-de-France

Course Overview

1	Mon, 17.9.2018	Monday's lecture: introduction, example problems, problem types		
	Thu, 20.9.2018 groups defined via wiki			
		everybody went (actively!) through the github.com/numbbo/coco	e Getting Started part of	
2	Fri, 21.9.2018	lecture "Benchmarking", final adjustments of groups everybody can run and postprocess the example experiment (~1h for final questions/help during the lecture)		
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4	Fri, 5.10.2018	lecture "Gradient-Based Algorithms" + DFO		
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7	Fri, 26.10.2018	final lecture "Discrete Optimization II: dyn. progr., B&B, heuristics"		
	29.102.11.2018	vacation aka learning for the exams		
	Thu, 8.11.2018 / Fri, 9.11.2018	oral presentations (individual time slots)		
	Fri, 16.11.2018	written exam	All deadlines:	
			23:59pm Paris time	

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Context discrete optimization:

- discrete variables
- or optimization over discrete structures (e.g. graphs)
- search space often finite, but typically too large for enumeration
- → need for smart algorithms

Algorithms for discrete problems:

- typically problem-specific
- but some general concepts are repeatedly used:
 - greedy algorithms
 - branch and bound
 - dynamic programming
 - randomized search heuristics

Motivation for this Part:

get an idea of the most common algorithm design principles

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before 2 excursions: the O-notation & graph theory

Remark: Coping with Difficult Problems

Exact

- brute-force often too slow
- better strategies such as dynamic programming & branch and bound
- still: often exponential runtime

Approximation Algorithms

- guarantee of low run time
- guarantee of high quality solution
- obstacle: difficult to prove these guarantees

Heuristics

- intuitive algorithms
- guarantee to run in short time
- often no guarantees on solution quality

we will see all 3 kinds of algorithms here...

Excursion: The O-Notation

Motivation:

- we often want to characterize how quickly a function f(x) grows asymptotically
- e.g. when we say an algorithm takes quadratically many steps (in the input size) to find the optimum of a problem with n (binary) variables, it is most likely not exactly n², but maybe n²+1 or (n+1)²

Big-O Notation

should be known, here mainly restating the definition:

Definition 1 We write f(x) = O(g(x)) iff there exists a constant c > 0 and an $x_0 > 0$ such that $f(x) \le c|g(x)|$ holds for all $x > x_0$.

we also view O(g(x)) as a set of functions growing at most as quick as g(x) and write $f(x) \in O(g(x))$

Big-O: Examples

- f(x) + c = O(f(x)) [if f(x) does not go to zero for x to infinity]
- $c \cdot f(x) = O(f(x))$
- $f(x) \cdot g(x) = O(f(x) \cdot g(x))$
- $3n^4 + n^2 7 = O(n^4)$

Intuition of the Big-O:

- if f(x) = O(g(x)) then g(x) gives an upper bound (asymptotically) for f
 excluding constants and lower order terms
- With Big-O, you should have '≤' in mind
- An algorithm that solves a problem in polynomial time is "efficient"
- An algorithm that solves a problem in exponential time is not
- But be aware:

In practice, often the line between efficient and non-efficient lies around $n \log n$ or even n (or even $\log n$ in the big data context) and the constants matter!!!

Excursion: The O-Notation

Further definitions to generalize from ' \leq ' to ' \geq ' and '=':

•
$$f(x) = \Omega(g(x))$$
 if $g(x) = O(f(x))$

• $f(x) = \Theta(g(x))$ if f(x) = O(g(x)) and g(x) = O(f(x))

Note: extensions to '<' and '>' exist as well, but are not needed here.

Example:

- Algo A solves problem P in time O(n)
- Algo B solves problem P in time O(n²)
- which one is faster?

only proving upper bounds to compare algorithms is not sufficient!

Excursion: The O-Notation

Further definitions to generalize from ' \leq ' to ' \geq ' and '=':

•
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• $f(x) = \Theta(g(x))$ if f(x) = O(g(x)) and g(x) = O(f(x))

Note: extensions to '<' and '>' exist as well, but are not needed here.

Example:

- Algo A solves problem P in time O(n)
- Algo B solves problem P in time $O(n^2) \Omega(n^2)$
- which one is faster?

only proving upper bounds to compare algorithms is not sufficient!

- Please order the following functions in terms of their asymptotic behavior (from smallest to largest):
 - exp(n²)
 - log n
 - In n / In In n
 - n
 - n log n
 - exp(n)
 - In n!

Pick one pair of runtimes and give a formal proof for the relation.

Exercise O-Notation (Solution)

Correct ordering: $\frac{\ln(n)}{\ln(\ln(n))} = O(\log n)$ $\log n = O(n)$ $n = O(n \log n)$ $n \log n = O(\ln(n!))$ $\ln(n!) = O(e^n)$ $e^n = O(e^{n^2})$

but for example $e^{n^2} \neq O(e^n)$

One exemplary proof: $\frac{\ln(n)}{\ln(\ln(n))} = O(\log n):$

$$\frac{\ln(n)}{\ln(\ln(n))} = \frac{\log(n)}{\log(e)\ln(\ln(n))} \leq \frac{3\log(n)}{\ln(\ln(n))} \leq 3|\log(n)|$$
for $n > 1$ for $n > 15$

Exercise O-Notation (Solution)

One additional proof: In n! = O(n log n)

Stirling's approximation:

$$n! \sim \sqrt{2\pi n} \left(n/e\right)^n$$
 or even

$$\sqrt{2\pi} \, n^{n+1/2} e^{-n} \le n! \le e \, n^{n+1/2} e^{-n}$$

•
$$\ln n! \leq \ln(en^{n+\frac{1}{2}}e^{-n}) = 1 + \left(n + \frac{1}{2}\right)\ln n - n$$

 $\leq \left(n + \frac{1}{2}\right)\ln n \leq 2n\ln n = 2n\frac{\log n}{\log e} = c \cdot n\log n$
okay for $c = 2/\log e$ and all $n \in \mathbb{N}$

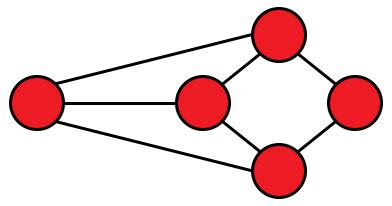
n ln n = O(ln n!) proven in a similar vein

Excursion: Basic Concepts of Graph Theory

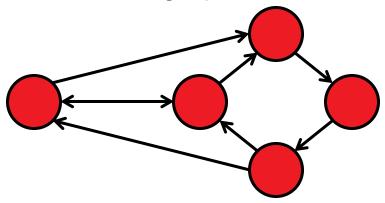
[following for example http://math.tut.fi/~ruohonen/GT_English.pdf]

Graphs

Definition 1 An undirected graph G is a tupel G = (V, E) of edges $e = \{u, v\} \in E$ over the vertex set V (i.e., $u, v \in V$).

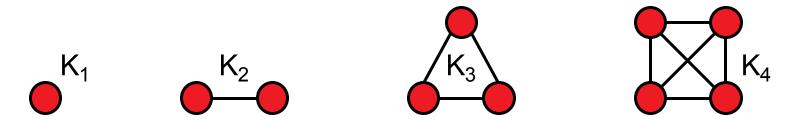


- vertices = nodes
- edges = lines
- Note: edges cover two unordered vertices (undirected graph)
 - if they are *ordered*, we call G a *directed* graph



Graphs: Basic Definitions

- G is called *empty* if E empty
- u and v are end vertices of an edge {u,v}
- Edges are *adjacent* if they share an end vertex
- Vertices u and v are *adjacent* if {u,v} is in E
- The *degree* of a vertex is the number of times it is an end vertex
- A complete graph contains all possible edges (once):





Walks, Paths, and Circuits

Definition 1 A walk in a graph G = (V, E) is a sequence

$$v_{i_0}, e_{i_1} = (v_{i_0}, v_{i_1}), v_{i_1}, e_{i_2} = (v_{i_1}, v_{i_2}), \dots, e_{i_k}, v_{i_k},$$

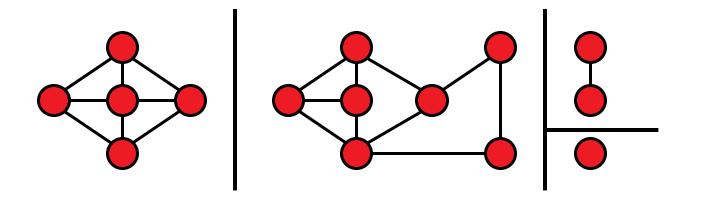
alternating vertices and adjacent edges of G.

A walk is

- closed if first and last node coincide
- a trail if each edge traversed at most once
- a path if each vertex is visited at most once
- a closed path is a *circuit* or *cycle*
- a closed path involving all vertices of G is a *Hamiltonian cycle*

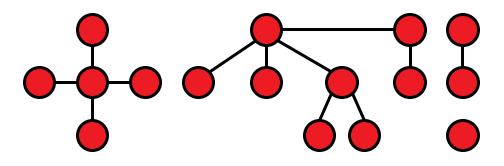
Graphs: Connectedness

- Two vertices are called *connected* if there is a walk between them in G
- If all vertex pairs in G are connected, G is called connected
- The connected components of G are the (maximal) subgraphs which are connected.

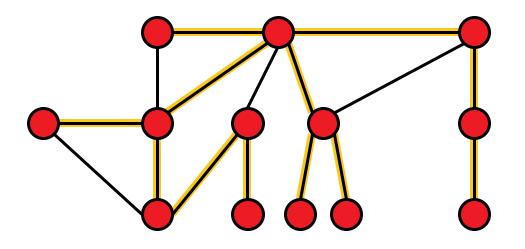


Trees and Forests

- A forest is a cycle-free graph
- A *tree* is a connected forest



A *spanning tree* of a connected graph G is a tree in G which contains all vertices of G



Greedy Algorithms

From Wikipedia:

"A *greedy algorithm* is an algorithm that follows the problem solving *heuristic* of making the locally optimal choice at each stage with the hope of finding a global optimum."

- Note: typically greedy algorithms do not find the global optimum
- We will see later when this is the case

Lecture Outline Greedy Algorithms

What we will see:

- Example 1: Money Change problem
- Example 2: Minimal Spanning Trees (MST) and the algorithm of Kruskal
- Example 3: An approximation algorithm for Bin Packing

Change-making problem

- Given n coins of distinct values w₁=1, w₂, ..., w_n and a total change W (where w₁, ..., w_n, and W are integers).
- Minimize the total amount of coins Σx_i such that $\Sigma w_i x_i = W$ and where x_i is the number of times, coin i is given back as change.

Greedy Algorithm

Unless total change not reached:

add the largest coin which is not larger than the remaining amount to the change

Note: only optimal for standard coin sets, not for arbitrary ones!

Related Problem:

finishing darts (from 501 to 0 with 9 darts)

Example 2: Minimal Spanning Trees (MST)

Outline:

- problem definition
- Kruskal's algorithm
- analysis of its running time
- proof of its correctness

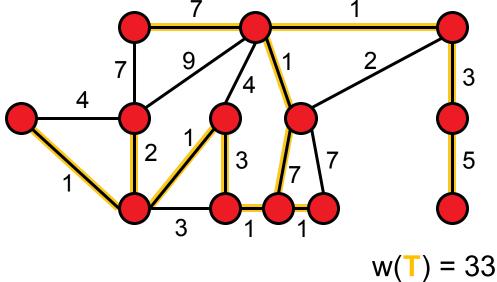
Minimum Spanning Trees (MST)

Minimum Spanning Tree problem:

Given a graph G=(V,E) with edge weights w_i for each edge e_i . Find the spanning tree with the smallest weight among all spanning trees.

weight of a spanning tree:

$$w(T) = \sum_{e_i \text{ in } T} w_i$$



Applications

Setting up a new wired telecommunication/water supply/electricity network

Constructing minimal delay trees for broadcasting in networks

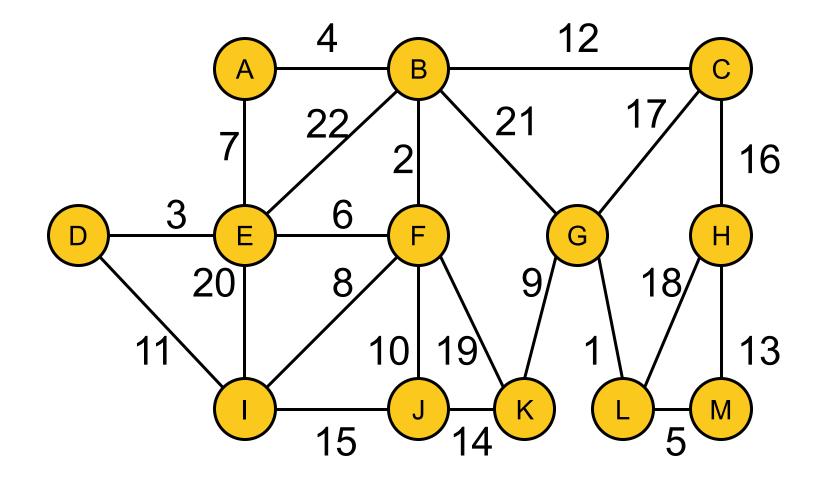
Kruskal's Algorithm: Idea

Algorithm, see [1]

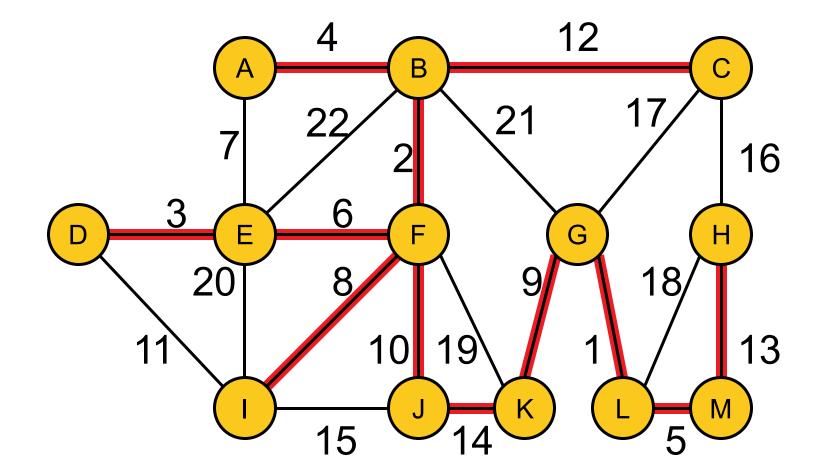
- Create forest F = (V,{}) with n components and no edge
- Put sorted edges (such that w.l.o.g. $w_1 < w_2 < ... < w_{|E|}$) into set S
- While S non-empty and F not spanning:
 - delete cheapest edge from S
 - add it to F if no cycle is introduced

[1] Kruskal, J. B. (1956). "On the shortest spanning subtree of a graph and the traveling salesman problem". *Proceedings of the American Mathematical Society* 7: 48–50. doi:10.1090/S0002-9939-1956-0078686-7

Kruskal's Algorithm: Example

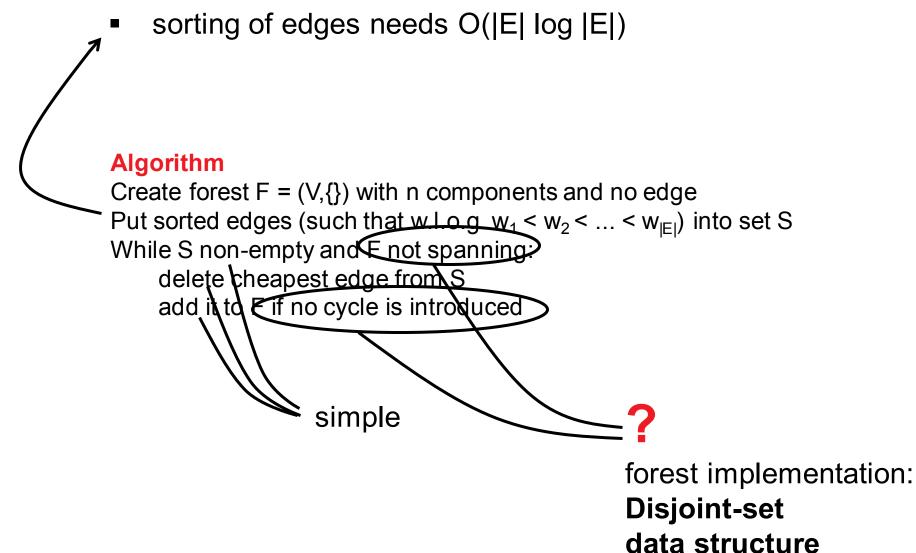


Kruskal's Algorithm: Example



Kruskal's Algorithm: Runtime Considerations

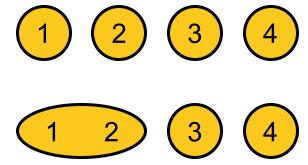
First question: how to implement the algorithm?



Disjoint-set Data Structure ("Union&Find")

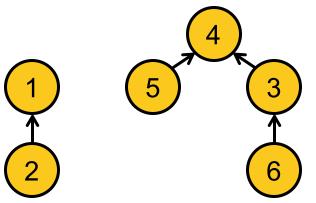
Data structure: ground set 1...N grouped to disjoint sets

- FIND(i): to which set does i belong?
- UNION(i,j): union the sets of i and j!



Implemented as trees:

- UNION(T1, T2): hang root node of smaller tree under root node of larger tree (constant time), thus
- FIND(u): traverse tree from u to root (to return a representative of u's set) takes logarithmic time in total number of nodes



Implementation of Kruskal's Algorithm

Algorithm, rewritten with UNION-FIND:

- Create initial disjoint-set data structure, i.e. for each vertex v_i, store v_i as representative of its set
- Create empty forest F = {}
- Sort edges such that w.l.o.g. $w_1 < w_2 < ... < w_{|E|}$
- for each edge e_i={u,v} starting from i=1:
 - if $FIND(u) \neq FIND(v)$: # no cycle introduced?
 - $F = F \cup \{\{u,v\}\}$
 - UNION(u,v)
- return F

Back to Runtime Considerations

- Sorting of edges needs O(|E| log |E|)
- forest: Disjoint-set data structure
 - initialization: O(|V|)
 - log |V| to find out whether the minimum-cost edge {u,v} connects two sets (no cycle induced) or is within a set (cycle would be induced)
 - 2x FIND + potential UNION needs to be done O(|E|) times
 - total O(|E| log |V|)
- Overall: O(|E| log |E|)

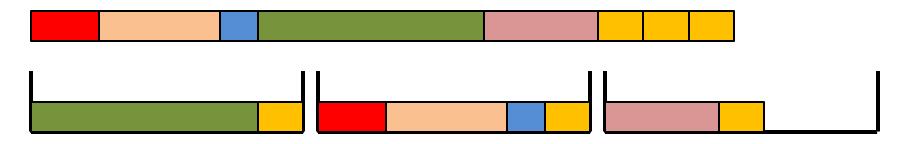
Kruskal's Algorithm: Proof of Correctness

Two parts needed:

- Algo always produces a spanning tree final F contains no cycle and is connected by definition
- Algo always produces a *minimum* spanning tree
 - argument by induction
 - P: If F is forest at a given stage of the algorithm, then there is some minimum spanning tree that contains F.
 - clearly true for F = (V, {})
 - assume that P holds when new edge e is added to F and be T a MST that contains F
 - if e in T, fine
 - if e not in T: T + e has cycle C with edge f in C but not in F (otherwise e would have introduced a cycle in F)
 - now T f + e is a tree with same weight as T (since T is a MST and f was not chosen to F)
 - hence T f + e is MST including F + e (i.e. P holds)

Bin Packing Problem

Given a set of n items with sizes a_1 , a_2 , ..., a_n . Find an assignment of the a_i 's to bins of size V such that the number of bins is minimal and the sum of the sizes of all items assigned to each bin is $\leq V$.

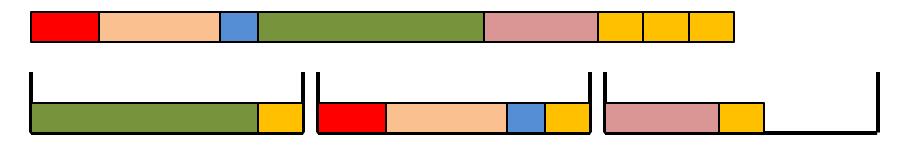


Applications

similar to multiprocessor scheduling of n jobs to m processors

Bin Packing Problem

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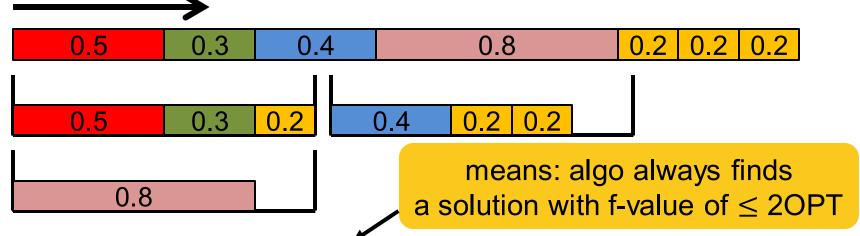


Known Facts

- no optimization algorithm reaches a better than 3/2 approximation in polynomial time (not shown here)
- greedy first-fit approach already yields an approximation algorithm with approximation ratio of 2

First-Fit Algorithm

- without sorting the items do:
 - put each item into the first bin where it fits
 - if it does not fit anywhere, open a new bin



Theorem: First-Fit algorithm is a 2-approximation algorithm

Proof: Assume First Fit uses m bins. Then, at least m-1 bins are more than half full (otherwise, move items).

$$OPT > \frac{m-1}{2} \iff 2OPT > m-1 \Longrightarrow 2OPT \ge m$$

 \uparrow because m and OPT are integer

Conclusion Greedy Algorithms I

What we have seen so far:

- two problems where a greedy algorithm was optimal
 - money change
 - minimum spanning tree (Kruskal's algorithm)
- but also: greedy not always optimal
 - see the example of bin packing
 - this is true in particular for so-called NP-hard problems

Obvious Question: when is greedy good?

Answer: if the problem is a matroid (not covered here)

From Wikipedia: [...] a matroid is a structure that captures and generalizes the notion of linear independence in vector spaces. There are many equivalent ways to define a matroid, the most significant being in terms of independent sets, bases, circuits, closed sets or flats, closure operators, and rank functions. I hope it became clear...

...what a greedy algorithm is ...that it not always results in the optimal solution ...but that it does if and only if the problem is a matroid

Dynamic Programming

Wikipedia:

"[...] **dynamic programming** is a method for solving a complex problem by breaking it down into a collection of simpler subproblems."

But that's not all:

- dynamic programming also makes sure that the subproblems are not solved too often but only once by keeping the solutions of simpler subproblems in memory ("trading space vs. time")
- it is an exact method, i.e. in comparison to the greedy approach, it always solves a problem to optimality

Note:

the reason why the approach is called "dynamic programming" is historical: at the time of invention by Richard Bellman, no computer "program" existed

Optimal Substructure

A solution can be constructed efficiently from optimal solutions of sub-problems

Overlapping Subproblems

Wikipedia: "[...] a problem is said to have **overlapping subproblems** if the problem can be broken down into subproblems which are reused several times or a recursive algorithm for the problem solves the same subproblem over and over rather than always generating new subproblems."

Note: in case of optimal substructure but independent subproblems, often greedy algorithms are a good choice; in this case, dynamic programming is often called "divide and conquer" instead

Main Idea Behind Dynamic Programming

Main idea: solve larger subproblems by breaking them down to smaller, easier subproblems in a recursive manner

Typical Algorithm Design:

- decompose the problem into subproblems and think about how to solve a larger problem with the solutions of its subproblems
- Specify how you compute the value of a larger problem recursively with the help of the optimal values of its subproblems ("Bellman equation")
- Bottom-up solving of the subproblems (i.e. computing their optimal value), starting from the smallest by using the Bellman equality and a table structure to store the optimal values (top-down approach also possible, but less common)
- eventually construct the final solution (can be omitted if only the value of an optimal solution is sought)

Introduction to Optimization Lecture 7: Discrete Optimization

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Remaining Lecture Overview

- O-notation
- graphs
- greedy algorithms
- dynamic programming
- branch and bound
- randomized search heuristics

Reminder Dynamic Programming (DP)

Dynamic Programming

- exact algorithm
- solve problem via solutions of subproblems ("optimal substructure")
- not solving overlapping subproblems twice, but store solutions

Lecture Outline Dynamic Programming (DP)

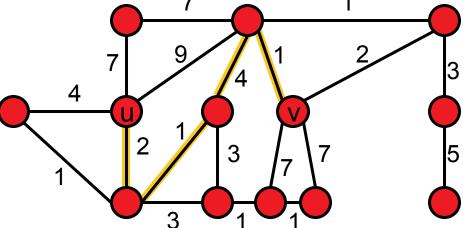
What we will see:

- Example 1: The All-Pairs Shortest Path Problem
- Example 2: The knapsack problem
- Example 3: An approximation algorithm for the knapsack problem

Example 1: The Shortest Path Problem

Shortest Path problem:

Given a graph G=(V,E) with edge weights w_i for each edge e_i . Find the shortest path from a vertex v to a vertex u, i.e., the path (v, $e_1 = \{v, v_1\}, v_1, ..., v_k, e_k = \{v_k, u\}, u$) such that $w_1 + ... + w_k$ is minimized.



Obvious Applications

Google maps

Autonomous cars

Finding routes for packages in a computer network

. . .

Example 1: The Shortest Path Problem

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Note:

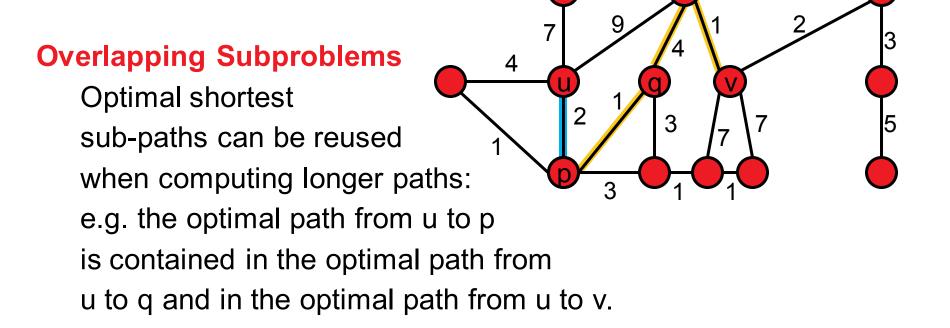
We can often assume that the edge weights are stored in a distance matrix D of dimension |E|x|E| where an entry D_{i,j} gives the weight between nodes i and j and "nonedges" are assigned a value of ∞

Why important? ⇒ determines input size

Opt. Substructure and Overlapping Subproblems

Optimal Substructure

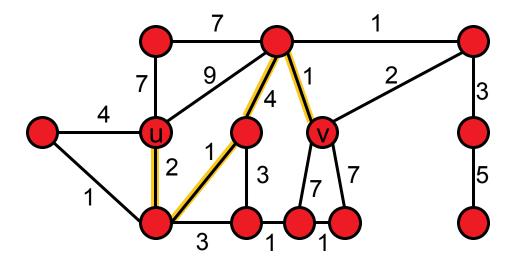
The optimal path from u to v, if it contains another vertex p can be constructed by simply joining the optimal path from u to p with the optimal path from p to v.



The All Pairs Shortest Paths Problem

All Pairs Shortest Path problem:

Given a graph G=(V,E) with edge weights w_i for each edge e_i . Find the shortest path from each source vertex v to each other target vertex u, i.e., the paths (v, e_1 ={v, v_1 }, v_1 , ..., v_k , e_k ={v_k,u}, u) such that w_1 + ... + w_k is minimized for all pairs (u,v) in V².



The Algorithm of Robert Floyd (1962)

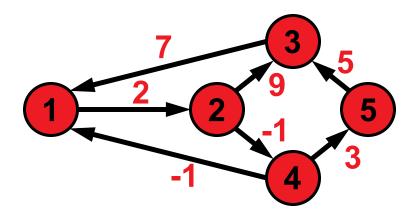
Idea:

- if we knew that the shortest path between source and target goes through node v, we would be able to construct the optimal path from the shorter paths "source→v" and "v→target"
- subproblem P(k): compute all shortest paths where the intermediate nodes can be chosen from v₁, ..., v_k

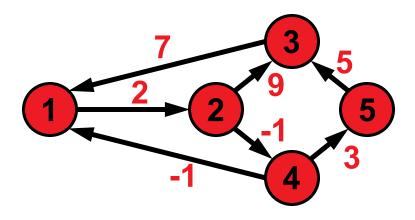
AllPairsShortestPathFloyd(G, D)

- Init: for all $1 \le i,j \le |V|$: dist(i,j) = D_{i,j}
- For k = 1 to |V| # solve subproblems P(k)
 - for all pairs of nodes (i.e. $1 \le i,j \le |V|$):
 - dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }

Note: Bernard Roy in 1959 and Stephen Warshall in 1962 essentially proposed the same algorithm independently.

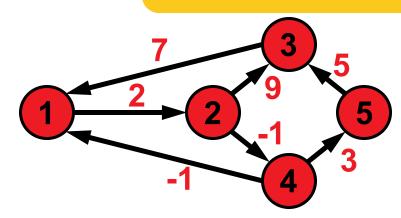


k=0	1	2	3	4	5
1					
2					
3					
4					
5					



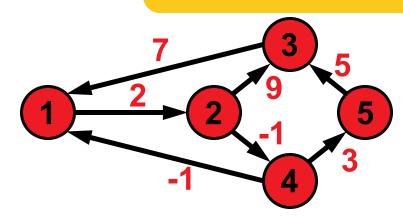
k=0	1	2	3	4	5
1	∞	2	∞	∞	∞
2	∞	∞	9	-1	∞
3	7	∞	∞	∞	∞
4	-1	∞	∞	∞	3
5	∞	∞	5	∞	∞

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



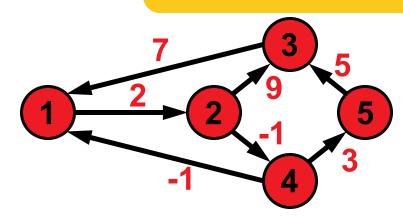
k=0	1	2	3	4	5	k=1	1	2	3	4	5
1	∞	2	∞	∞	∞	1					
2	∞	∞	9	-1	∞	2					
3	7	∞	∞	∞	∞	3					
4	-1	∞	∞	∞	3	4					
5	∞	∞	5	∞	∞	5					

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



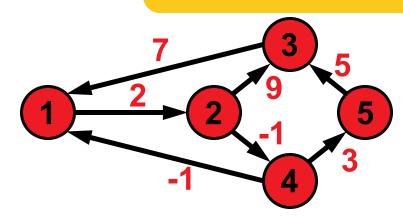
									-		
k=0	1	2	3	4	5	k=1	1	2	3	4	5
1	~	2	∞	∞	∞	1					
2	∞	∞	9	-1	∞	2					
3		∞	∞	∞	∞	3					
4		∞	∞	∞	3	4					
5	∞	∞	5	∞	∞	5					

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



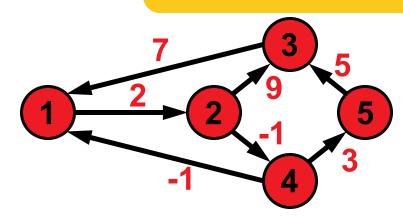
									-		
k=0	1	2	3	4	5	k=1	1	2	3	4	5
1	~	2	∞	∞	∞	1					
2	∞	∞	9	-1	∞	2					
3		∞	∞	∞	∞	3					
4		∞	∞	∞	3	4					
5	∞	∞	5	∞	∞	5					

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



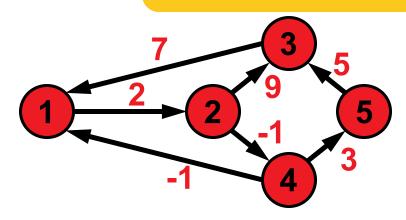
k=0	1	2	3	4	5	k=1	1	2	3	4	5
1	∞	2	∞	∞	∞	1					
2	∞	∞	9	-1	∞	2					
3		∞	∞	∞	∞	3		9			
4		∞	∞	∞	3	4		1			
5	∞	∞	5	∞	∞	5					

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



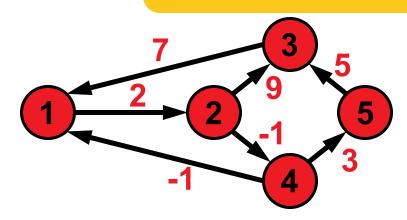
			-								
k=0	1	2	3	4	5	k=1	1	2	3	4	5
1	~ (2	∞	∞	∞	1	∞	2	∞	∞	∞
2	∞	∞	9	-1	∞	2	∞	∞	9	-1	∞
3		∞	∞	∞	∞	3	7	9	∞	∞	∞
4		∞	∞	∞	3	4	-1	1	∞	∞	3
5	∞	∞	5	∞	∞	5	∞	∞	5	∞	∞

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



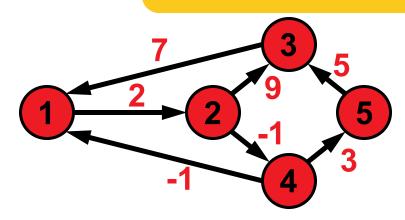
k=1	1	2	3	4	5	k=2	1	2	3	4	5
1	∞	2	∞	∞	∞	1	∞	2	∞	∞	∞
2	∞	∞	9	-1	∞	2	∞	∞	9	-1	∞
3	7	9	∞	∞	∞	3	7	9	∞	∞	∞
4	-1	1	∞	∞	3	4	-1	1	∞	∞	3
5	∞	∞	5	∞	∞	5	∞	∞	5	∞	∞

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



k=1	1	2	3	4	5	k=2	1	2	3	4	5
1 -		2	∞	∞	∞	1	∞	2	∞	∞	∞
2	∞	~		<u>→</u> 1	∞	2	∞	∞	9	-1	∞
3	7	9	∞	∞	∞	3	7	9	∞	∞	∞
4 _	1	1	∞	∞	3	4	-1	1	∞	∞	3
5	∞	∞	5	∞	∞	5	∞	∞	5	∞	∞

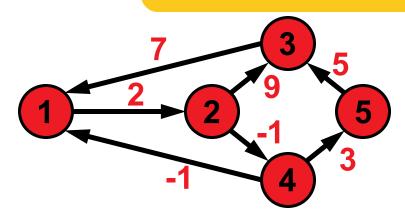
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow 1 & 2 as intermediate nodes

k=1	1	2	3	4	5	k=2	1	2	3	4	5
1-		-2	∞	∞	∞	1	∞	2	11	1	∞
2	∞	∞		-1	∞	2	∞	∞	9	-1	∞
3	7	9	∞	∞	∞	3	7	9	18	8	∞
4 _	1		∞	∞	3	4	-1	1	10	0	3
5	∞	∞	5	∞	∞	5	∞	∞	5	∞	∞

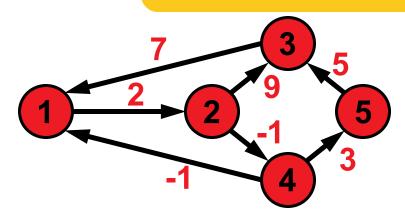
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow {1,2,3} as intermediate nodes

k=2	1	2	3	4	5	k=3	1	2	3	4	5
1	∞	2	11	1	∞	1	∞	2	11	1	∞
2	∞	∞	9	-1	∞	2	∞	∞	9	-1	∞
3	7	9	18	8	∞	3	7	9	18	8	∞
4	-1	1	10	0	3	4	-1	1	10	0	3
5	∞	∞	5	∞	∞	5	∞	∞	5	∞	∞

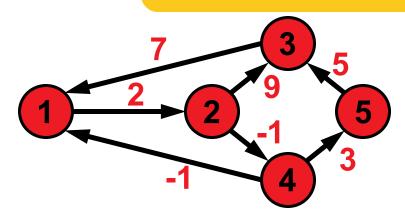
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow {1,2,3} as intermediate nodes

k=2	1	2	3	4	5	k=3	1	2	3	4	5
1	∞	2	11	1	∞	1			11		∞
2	∞	∞	9	-1	∞	2			9		∞
3	7	9	18	8	∞	3	7	9	18	8	∞
4	-1	1	10	0	3	4			10		3
5	∞	∞	5	∞	∞	5			5		∞

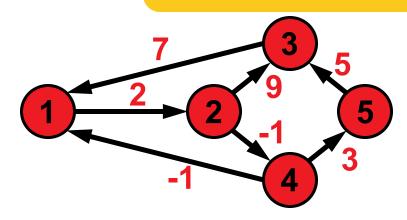
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow {1,2,3} as intermediate nodes

k=2	1	2	3	4	5	k=3	1	2	3	4	5
1	∞	2	11	1	∞	1	18	2	11	1	∞
2	∞	∞	9	-1	∞	2	16	18	9	-1	∞
3	7	9	18	8	∞	3	7	9	18	8	∞
4	-1	1	10	0	3	4	-1	1	10	0	3
5	∞	∞	5	∞	∞	5	12	14	5	13	∞

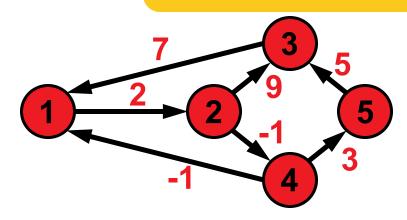
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow {1,2,3,4} as intermediate nodes

k=3	1	2	3	4	5	k=4	1	2	3	4	5
1	18	2	11	1	∞	1	18	2	11	1	∞
2	16	18	9	-1	∞	2	16	18	9	-1	∞
3	7	9	18	8	∞	3	7	9	18	8	∞
4	-1	1	10	0	3	4	-1	1	10	0	3
5	12	14	5	13	∞	5	12	14	5	13	∞

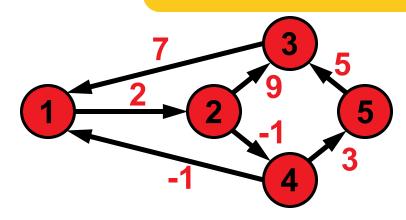
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow {1,2,3,4} as intermediate nodes

k=3	1	2	3	4	5	k=4	1	2	3	4	5
1	18	2	11	1	∞	1				1	
2	16	18	9	-1	∞	2				-1	
3	7	9	18	8	∞	3				8	
4	-1	1	10	0	3	4	-1	1	10	0	3
5	12	14	5	13	∞	5				13	

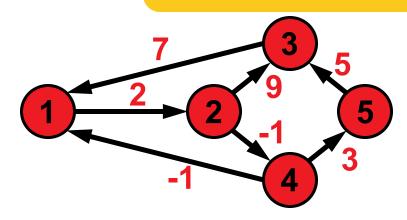
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow {1,2,3,4} as intermediate nodes

k=3	1	2	3	4	5	k=4	1	2	3	4	5
1	18	2	11	1	∞	1	0	2	11	1	4
2	16	18	9	-1	∞	2	-2	0	9	-1	2
3	7	9	18	8	∞	3	7	9	18	8	11
4	-1	1	10	0	3	4	-1	1	10	0	3
5	12	14	5	13	∞	5	12	14	5	13	16

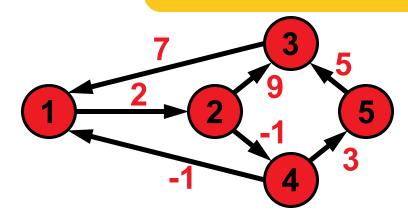
for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow all nodes as intermediate nodes

		-										
k=4	1	2	3	4	5	k=5	1	2	3	4	5	
1	0	2	11	1	4	1	0	2	11	1	4	
2	-2	0	9	-1	2	2	-2	0	9	-1	2	
3	7	9	18	8	11	3	7	9	18	8	11	
4	-1	1	10	0	3	4	-1	1	10	0	3	
5	12	14	5	13	16	5	12	14	5	13	16	

for all pairs of nodes (i.e. $1 \le i,j \le |V|$): dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }



allow all nodes as intermediate nodes

		-									
k=4	1	2	3	4	5	k=5	1	2	3	4	5
1	0	2	11	1	4	1	0	2	9	1	4
2	-2	0	9	-1	2	2	-2	0	7	-1	2
3	7	9	18	8	11	3	7	9	16	8	11
4	-1	1	10	0	3	4	-1	1	8	0	3
5	12	14	5	13	16	5	12	14	5	13	16

Runtime Considerations and Correctness

O(|V|³) easy to show

O(|V|²) many distances need to be updated O(|V|) times

Correctness

given by the Bellman equation

dist(i,j) = min { dist(i,j), dist(i,k) + dist(k,j) }

 only correct if cycles do not have negative total weight (can be checked in final distance matrix if diagonal elements are negative)

But How Can We Actually Construct the Paths?

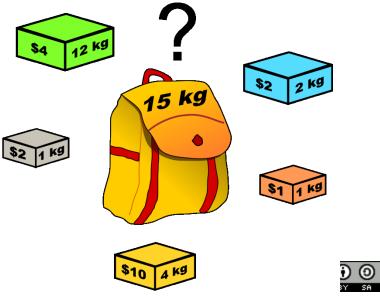
- Construct matrix of predecessors *P* alongside distance matrix
- $P_{i,j}(k)$ = predecessor of node j on path from i to j (at algo. step k)
- no extra costs (asymptotically)

$$P_{i,j}(0) = \begin{cases} 0 & \text{if } i = j \text{ or } d_{i,j} = \infty \\ i & \text{in all other cases} \end{cases}$$

$$P_{i,j}(k) = \begin{cases} P_{i,j}(k-1) & \text{if } \operatorname{dist}(i,j) \leq \operatorname{dist}(i,k) + \operatorname{dist}(k,j) \\ P_{k,j}(k-1) & \text{if } \operatorname{dist}(i,j) > \operatorname{dist}(i,k) + \operatorname{dist}(k,j) \end{cases}$$

Example 2: The Knapsack Problem (KP)

Knapsack Problem
max.
$$\sum_{j=1}^{n} p_j x_j$$
 with $x_j \in \{0, 1\}$
s.t. $\sum_{j=1}^{n} w_j x_j \le W$



Dake

Opt. Substructure and Overlapping Subproblems

Consider the following subproblem:

P(i, j): optimal profit when packing the first *i* items into a knapsack of size *j*

Optimal Substructure

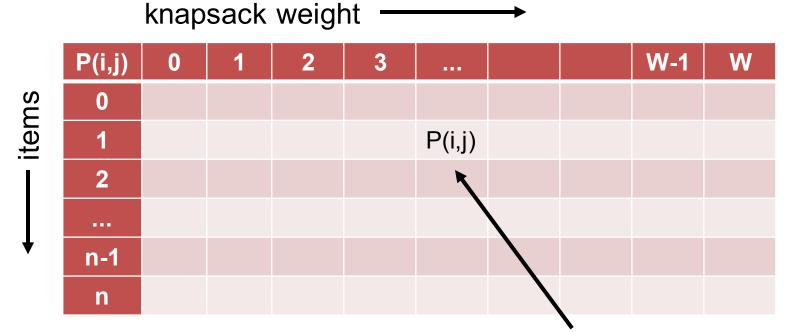
The optimal choice of whether taking item *i* or not can be made easily for a knapsack of weight *j* if we know the optimal choice for items $1 \dots i - 1$:

$$P(i,j) = \begin{cases} 0 & \text{if } i = 0 \text{ or } j = 0\\ P(i-1,j) & \text{if } w_i > j\\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

Overlapping Subproblems

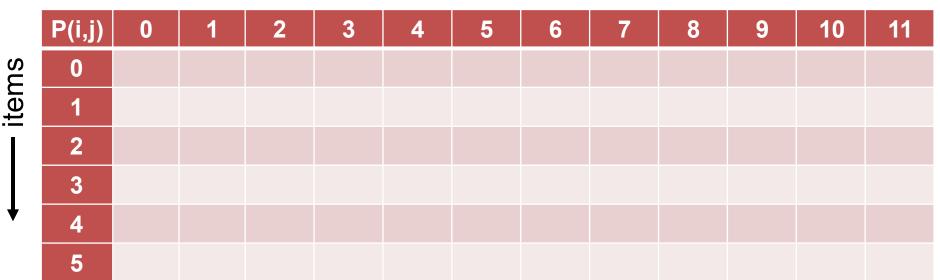
a recursive implementation of the Bellman equation is simple, but the P(i, j) might need to be computed more than once!

To circumvent computing the subproblems more than once, we can store their results (in a matrix for example)...



best achievable profit with items 1...i and a knapsack of size j

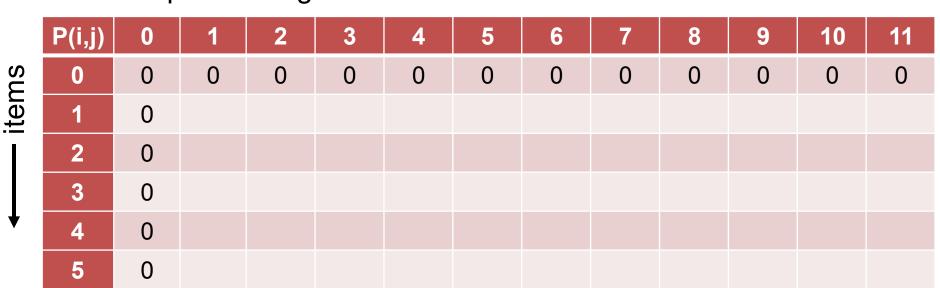
Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W=11.



knapsack weight

initialization: P(i,j) = 0 if i = 0 or j = 0

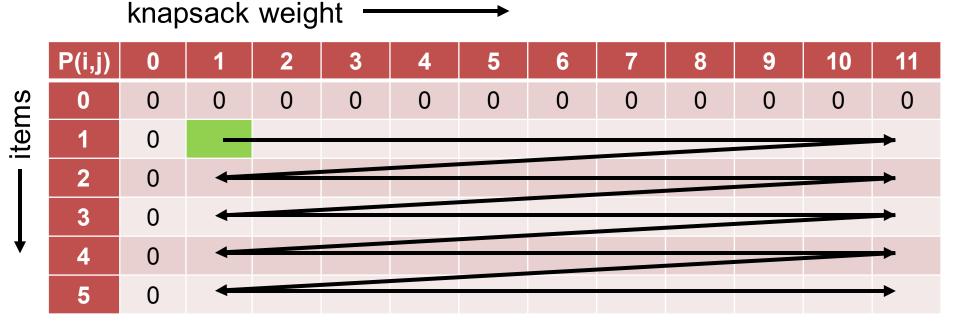
Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W=11.



knapsack weight -----

initialization: P(i,j) = 0 if i = 0 or j = 0

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.



for
$$i = 1$$
 to n :
for $j = 1$ to W :

$$P(i,j) = \begin{cases} P(i-1,j) & \text{if } w_i > j \\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

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Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

		•		U									
	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0	0										
	2	0											
	3	0											
♦	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

		•		0									
	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0	0	0									
	2	0											
	3	0											
♦	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

		•		0									
	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0	0	0	0	0							
	2	0											
	3	0											
♦	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

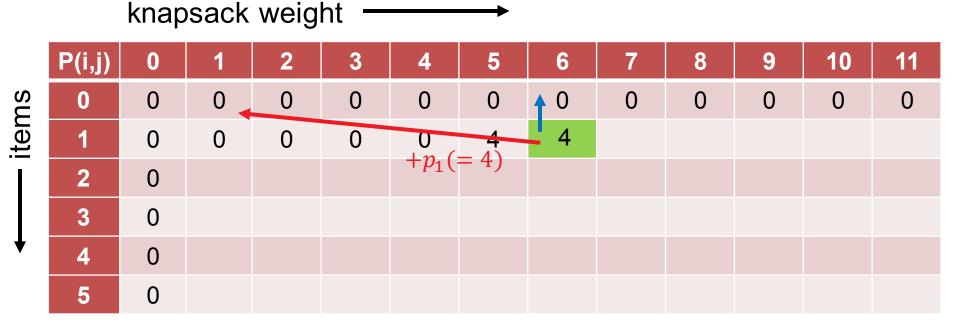
		•		Ŭ									
	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
item	1	0	0	0	0	0	4						
	2	0			$+p_{1}($	= 4)							
	3	0											
♥	4	0											
	5	0											

knapsack weight -----

for
$$i = 1$$
 to n :
for $j = 1$ to W :

$$P(i,j) = \begin{cases} P(i-1,j) & \text{if } w_i > j \\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.



for
$$i = 1$$
 to n :
for $j = 1$ to W :

$$P(i,j) = \begin{cases} P(i-1,j) & \text{if } w_i > j \\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

		•		•									
	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0											
	3	0											
♦	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ms	0	0	0	0	0	0	0	0	0	0	0	0	0
iter	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0	0	0	0	0	4	4					
	3	0											
¥	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0 🔺	0	0	0	0	4	4	↑ 4	4	4	4	4
	2	0	0	0	0	0	4	10)	- 10				
	3	0					$+p_{2}($	= 10)					
♥	4	0											
	5	0											

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ms	0	0	0	0	0	0	0	0	0	0	0	0	0
iter	1	0	0	0	0	0	4	4	4	4	4	4	4
Ι	2	0	0	0	0	0	4	4	10	10	10	10	10
	3	0											
¥	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ms	0	0	0	0	0	0	0	0	0	0	0	0	0
iter	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0	0	0	0	0	4	4	10	10	10	10	10
	3	0	0	3	3	3							
¥	4	0											
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0	0	0	0	0	4	4	4	4	4	4	4
Ι	2	0	0	0	0	0	↑ 4	4	10	10	10	10	10
	3	0	0	3	3	3	4						
♦	4	0			$+p_3($	- 5)							
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ns	0	0	0	0	0	0	0	0	0	0	0	0	0
items	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0	0	0	0	0	4	↑ 4	10	10	10	10	10
	3	0	0	3	3	3	4	4					
•	4	0				$\pm p_3$	(- 3)						
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
ms	0	0	0	0	0	0	0	0	0	0	0	0	0
iter	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0	0	0	0	0	4	4	1 0	10	10	10	10
	3	0	0	3	3	3	4	4	10	etc.			
•	4	0					$+p_3$	(- 5)					
	5	0											

knapsack weight

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
← items	0	0	0	0	0	0	0	0	0	0	0	0	0
	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0	0	0	0	0	4	4	10	10	10	10	10
	3	0	0	3	3	3	4	4	10	10	13	13	13
	4	0	0	3	3	5	5	8	10	10	13	13	15
	5	0	0	3	3	5	6	8	10	10	13	13	15

knapsack weight

for i = 1 to n:

for
$$j = 1$$
 to W :

$$P(i,j) = \begin{cases} P(i-1,j) & \text{if } w_i > j \\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

Example instance with 5 items with weights and profits (5,4), (7,10), (2,3), (4,5), and (3,3). Weight restriction is W = 11.

	P(i,j)	0	1	2	3	4	5	6	7	8	9	10	11
	0	0	0	0	0	0	0	0	0	0	0	0	0
	1	0	0	0	0	0	4	4	4	4	4	4	4
	2	0	0	0	0	0	4	4	10	10	10	10	10
	3	0	0	3	3	3	4	4	10	10	13	13	13
¥	4	0	0	3	3	5	5	8	10	10	13	13	15
	5	0	0	3	3	5	6	8	10	10	13	13	15

knapsack weight

for i = 1 to n:

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$$P(i,j) = \begin{cases} P(i-1,j) & \text{if } w_i > j \\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

Question: How to obtain the actual packing?

Answer: we just need to remember where the max came from!

P(i,j) $(\mathbf{0})$ items x_1 $x_2 = 1$ $x_3^{10} = 0^{10}$ 13 x4 <u>1</u>3₁ x_5 = 0

knapsack weight

for i = 1 to n:

for
$$j = 1$$
 to W :

$$P(i,j) = \begin{cases} P(i-1,j) & \text{if } w_i > j \\ \max\{P(i-1,j), p_i + P(i-1,j-w_i)\} & \text{if } w_i \le j \end{cases}$$

Example 3: Approximating the Knapsack Problem

- Simple to prove runtime of the previous algorithm as O(nW)
- In practice, i.e. for large n and W, computing the exact optimum for the knapsack problem might be too costly.

no polynomial-time algo exists under famous P≠NP hypothesis

 If we want to have a polynomial-time algorithm, we have to trade for approximation quality.

Approximations, PTAS, and FPTAS

An algorithm is a ρ -approximation algorithm for problem Π if, for each problem instance of Π , it outputs a feasible solution which function value is within a ratio ρ of the true optimum for that instance.

remember the 2-approximation algo for bin packing?

- An algorithm A is an *approximation scheme* for a *maximization* problem Π if for any instance I of Π and a parameter $\varepsilon > 0$, it outputs a solution s with f $_{\Pi}(I,s) \ge (1-\epsilon) \cdot OPT$.
- An approximation scheme is called *polynomial time* approximation scheme (PTAS) if for a fixed $\varepsilon > 0$, its running time is polynomially bounded in the size of the instance I.

note: runtime might be exponential in $1/\epsilon$ actually!

An approximation scheme is a *fully polynomial time* approximation scheme (FPTAS) if its running time is bounded polynomially in both the size of the input I and in $1/\epsilon$.

An FPTAS for the Knapsack Problem

Similar to the previous dynamic programming algorithm, we can design a dynamic programming algorithm for which

- a subproblem is restricting the items to {1, ..., k} and searches for the lightest packing with prefixed profit P
- runs in $O(n^2 P_{\text{max}})$ [idea: fill matrix of size $n \times nP_{\text{max}}$]

What strange runtime is $O(n^2 P_{\text{max}})$?

Answer: pseudo-polynomial (polynomial if P_{max} would be polynomial in input size)

Idea behind FPTAS:

- scale all profits p_i smartly to $\left[\frac{p_i n}{\varepsilon P_{\max}}\right]$ to make P_{\max} polynomially bounded
- prove that dynamic programming approach computes profit of at least (1-ε)·OPT (not shown here)

Branch and Bound

Branch and Bound: General Ideas

- Systematic enumeration of candidate solutions in terms of a rooted tree
- Each tree node corresponds to a set of solutions; the whole search space on the root
- At each tree node, the corresponding subset of the search space is split into (non-overlapping) sub-subsets:
 - the optimum of the larger problem must be contained in at least one of its subproblems
- If tree nodes correspond to small enough subproblems, they are solved exhaustively.
- The smart part of the algorithm is the estimation of upper and lower bounds on the optimal function value achieved by solutions in the tree nodes
 - the exploration of a tree node is stopped ("pruning the tree") if a node's upper bound is already lower than the lower bound of an already explored node (assuming maximization)

Applying Branch and Bound

Needed for successful application of branch and bound:

- optimization problem
- finite (or at least countable) set of solutions
- clear idea of how to split problem into smaller subproblems
- efficient calculation of the following modules:
 - upper bound calculation
 - lower bound calculation

Computing Bounds (Maximization Problems)

Assume w.l.o.g. maximization of f(x) here

Lower Bounds

- any actual feasible solution will give a lower bound (which will be exact if the solution is the optimal one for the subproblem)
- hence, sampling a (feasible) solution can be one strategy
- using a heuristic to solve the subproblem another one

Upper Bounds

 upper bounds can be achieved by solving a relaxed version of the problem formulations (i.e. by either loosening or removing constraints)

Note: the better/tighter the bounds, the quicker the branch and bound tree can be pruned

Properties of Branch and Bound Algorithms

- Exact, global solver
- Can be slow; only exponential worst-case runtime
 - due to the exhaustive search behavior if no pruning of the search tree is possible
- but might work well in some cases

Advantages:

- can be stopped if lower and upper bound are "close enough" in practice (not necessarily exact anymore then)
- can be combined with other techniques, e.g. "branch and cut" (not covered here)

Example Branching Decisions

0-1 problems:

- choose unfixed variable x_i
- one subproblem defined by setting x_i to 0
- one subproblem defined by setting x_i to 1

General integer problem:

- choose unfixed variable x_i
- choose a value c that x_i can take
- one subproblem defined by restricting $x_i \le c$
- one subproblem defined by restricting x_i > c

Combinatorial Problems:

 branching on certain discrete choices, e.g. an edge/vertex is chosen or not chosen

The branching decisions are then induced as additional constraints when defining the subproblems.

Which Tree Node to Branch on?

Several strategies (again assuming maximization):

- choose the subproblem with highest upper bound
 - gain the most in reducing overall upper bound
 - if upper bound not the optimal value, this problem needs to be branched upon anyway sooner or later
- choose the subproblem with lowest lower bound
- simple depth-first search or breadth-first search

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see for example
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https://en.wikipedia.org/wiki/Depth-first_search
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https://en.wikipedia.org/wiki/Breadth-first_search

problem-specific approach most likely to be a good choice

4 Steps Towards a Branch and Bound Algorithm

Concrete steps when designing a branch and bound algorithm:

- How to split a problem into subproblems ("branching")?
- How to compute upper bounds (assuming maximization)?
- Optional: how to compute lower bounds?
- How to decide which next tree node to split?

Example: Application to ILPs

 $\begin{array}{ll} \text{maximize} & c^T x\\ \text{subject to} & Ax \leq b\\ & x \geq 0\\ & \text{and} & x \in \mathbb{Z}^n \end{array}$

The ILP formalization covers many problems such as

- Traveling Salesperson Person (TSP)
- Vertex Cover and other covering problems
- Set packing and other packing problems
- Boolean satisfiability (SAT)

Possible Ways to Solve an ILP

- Do not restrict the solutions to integers and round the solution found of the relaxed problem (=remove the integer constraints) by a continuous solver (i.e. solving the so-called *LP relaxation*)
 → no guarantee to be exact
- Exploiting the instance property of A being total unimodular:
 - feasible solutions are guaranteed to be integer in this case
 - algorithms for continuous relaxation can be used (e.g. the simplex algorithm)
- Using heuristic methods (typically without any quality guarantee)
 - we'll see these type of algorithms later in the lecture
- Using exact algorithms such as branch and bound

Here, we just give an idea instead of a concrete algorithm...

- How to split a problem into subproblems ("branching")?
- How to compute upper bounds (assuming maximization)?
- Optional: how to compute lower bounds?
- How to decide which next tree node to split?

Here, we just give an idea instead of a concrete algorithm...

- How to compute upper bounds (assuming maximization)?
- How to split a problem into subproblems ("branching")?
- Optional: how to compute lower bounds?
- How to decide which next tree node to split?

Branch and Bound for ILPs

How to compute upper bounds (assuming maximization)?

- drop the integer constraints and solve the so-called LPrelaxation
- can be done by standard LP algorithms such as scipy.optimize.linprog or Matlab's linprog

What's then?

- The LP has no feasible solution. Fine. Prune.
- We found an integer solution. Fine as well. Might give us a new lower bound to the overall problem.
- The LP problem has an optimal solution which is worse than the highest lower bound over all already explored subproblems. Fine. Prune.
- Otherwise: Branch on this subproblem: e.g. if optimal solution has x_i=2.7865, use x_i≤2 and x_i≥3 as new constraints

Branch and Bound for ILPs

How to split a problem into subproblems ("branching")?

- mainly needed if the solution of the LP-relaxation is not integer
- branch on a variable which is rational

Not discussed here in depth due to time:

- Optional: how to compute lower bounds?
- How to decide which next tree node to split?
 - seems to be good choice: subproblem with largest upper bound of LP-relaxation

I hope it became clear...

...what the algorithm design ideas of dynamic programming and branch and bound are

...and for which problem types they are supposed to be suitable

(Randomized) Search Heuristics

Motivation General Search Heuristics

- often, problem complicated and not much time available to develop a problem-specific algorithm
- search heuristics are a good choice:
 - relatively easy to implement
 - easy to adapt/change/improve
 - e.g. when the problem formulation changes in an early product design phase
 - or when slightly different problems need to be solved over time
- randomized/stochastic algorithms are a good choice because they are robust to noise

Lecture Outline Randomized Search Heuristics

Which algorithms will we touch?

- Randomized Local Search (RLS)
- Variable Neighborhood Search (VNS)
- Tabu Search (TS)
- Evolutionary Algorithms (EAs)

Neighborhoods

For most (stochastic) search heuristics, we need to define a *neighborhood structure*

which search points are close to each other?

Example: k-bit flip / Hamming distance k neighborhood

- search space: bitstrings of length n (Ω={0,1}ⁿ)
- two search points are neighbors if their Hamming distance is k
- in other words: x and y are neighbors if we can flip exactly k bits in x to obtain y
- 0001001101 is neighbor of 0001000101 for k=1 0101000101 for k=2 1101000101 for k=3

Neighborhoods II

Example: possible neighborhoods for the knapsack problem

- search space again bitstrings of length n ($\Omega = \{0, 1\}^n$)
- Hamming distance 1 neighborhood:
 - add an item or remove it from the packing
- replacing 2 items neighborhood:
 - replace one chosen item with an unchosen one
 - makes only sense in combination with other neighborhoods because the number of items stays constant
- Hamming distance 2 neighborhood on the contrary:
 - allows to change 2 arbitrary items, e.g.
 - add 2 new items
 - remove 2 chosen items
 - or replace one chosen item with an unchosen one

Randomized Local Search (RLS)

Idea behind (Randomized) Local Search:

explore the local neighborhood of the current solution (randomly)

Pure Random Search:

go to randomly chosen neighbor

First Improvement Local Search:

go to first (randomly) chosen neighbor which is better

Best Improvement strategy:

- always go to the best neighbor
- not random anymore
- computationally expensive if neighborhood large

Variable Neighborhood Search

Main Idea: [Mladenovic and P. Hansen, 1997]

- change the neighborhood from time to time
 - local optima are not the same for different neighborhood operators
 - but often close to each other
 - global optimum is local optimum for all neighborhoods
- rather a framework than a concrete algorithm
 - e.g. deterministic and stochastic neighborhood changes
- typically combined with (i) first improvement, (ii) a random order in which the neighbors are visited and (iii) restarts

N. Mladenovic and P. Hansen (1997). "Variable neighborhood search". Computers and Operations Research 24 (11): 1097–1100.

Disadvantages of local searches (with or without varying neighborhoods)

- they get stuck in local optima
- have problems to traverse large plateaus of equal objective function value ("random walk")

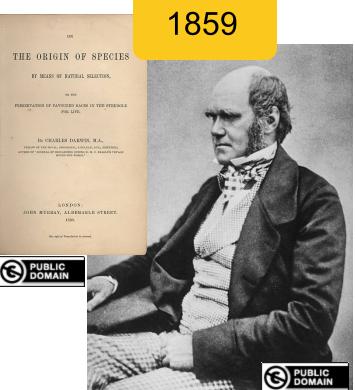
Tabu search addresses these by

- allowing worsening moves if all neighbors are explored
- introducing a tabu list of temporarily not allowed moves
- those restricted moves are
 - problem-specific and
 - can be specific solutions or not permitted "search directions" such as "don't include this edge anymore" or "do not flip this specific bit"
- the tabu list is typically restricted in size and after a while, restricted moves are permitted again

Stochastic Optimization Algorithms

One class of (bio-inspired) stochastic optimization algorithms: Evolutionary Algorithms (EAs)

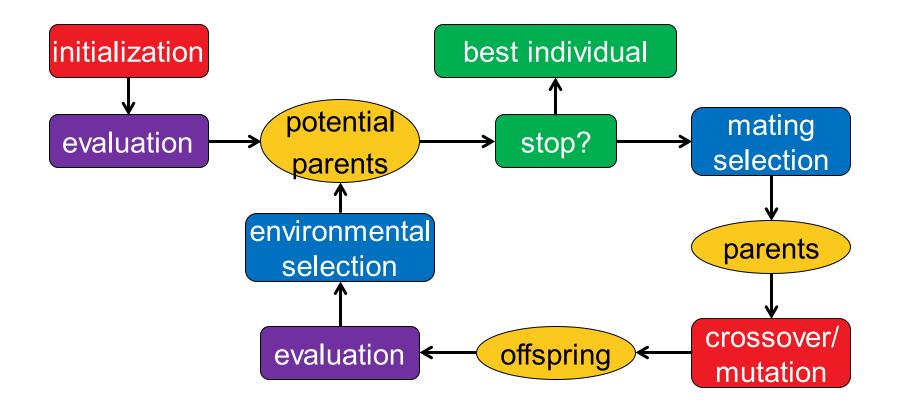
- Class of optimization algorithms originally inspired by the idea of biological evolution
- selection, mutation, recombination



Metaphors

Classical Optimization	Evolutionary Computation
variables or parameters	variables or chromosomes
candidate solution vector of decision variables / design variables / object variables	individual, offspring, parent
set of candidate solutions	population
objective function loss function cost function error function	fitness function
iteration	generation

Generic Framework of an EA



stochastic operators

"Darwinism"

stopping criteria

Important: representation (search space)

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The Historic Roots of EAs

Genetic Algorithms (GA)

J. Holland 1975 and D. Goldberg (USA) $\Omega = \{0,1\}^n$

Evolution Strategies (ES)

I. Rechenberg and H.P. Schwefel, 1965 (Berlin) $\Omega = \mathbb{R}^{n}$

Evolutionary Programming (EP)

L.J. Fogel 1966 (USA)

Genetic Programming (GP)

 $\Omega = \text{space of all programs}$

nowadays one umbrella term: evolutionary algorithms

Genotype – Phenotype mapping

The genotype – phenotype mapping

- related to the question: how to come up with a fitness ("quality") of each individual from the representation?
- related to DNA vs. actual animal (which then has a fitness)

fitness of an individual not always = f(x)

- include constraints
- include diversity
- others
- but needed: always a total order on the solutions

Handling Constraints

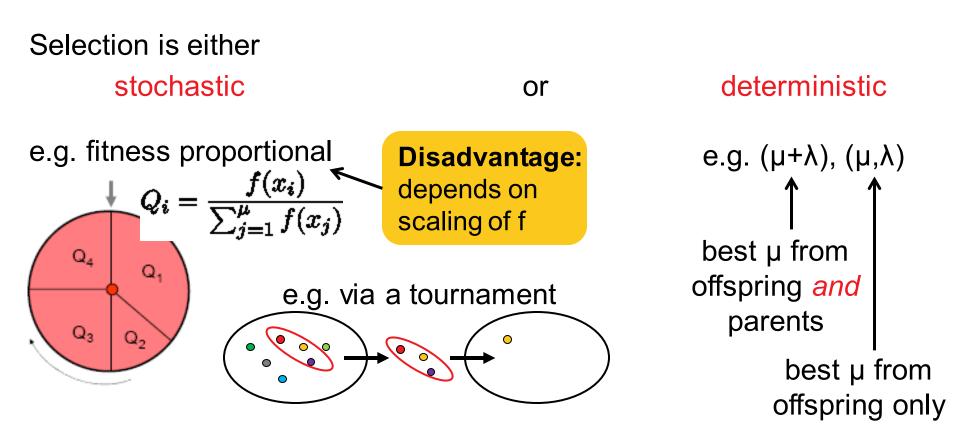
Several possible ways to handle constraints, e.g.:

- resampling until a new feasible point is found ("often bad idea")
- penalty function approach: add constraint violation term (potentially scaled)
- repair approach: after generation of a new point, repair it (e.g. with a heuristic) to become feasible again if infeasible
 - continue to use repaired solution in the population or
 - use repaired solution only for the evaluation?
- multiobjective approach: keep objective function and constraint functions separate and try to optimize all of them in parallel

Examples for some EA parts

Selection

Selection is the major determinant for specifying the trade-off between exploitation and exploration



Mating selection (selection for variation): usually stochastic Environmental selection (selection for survival): often deterministic

Variation Operators

Variation aims at generating new individuals on the basis of those individuals selected for mating

Variation = Mutation and Recombination/Crossover

mutation: mut: $\Omega \to \Omega$ recombination: recomb: $\Omega^r \to \Omega^s$ where $r \ge 2$ and $s \ge 1$

- choice always depends on the problem and the chosen representation
- however, there are some operators that are applicable to a wide range of problems and tailored to standard representations such as vectors, permutations, trees, etc.

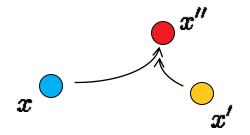
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Variation Operators: Guidelines

Two desirable properties for mutation operators:

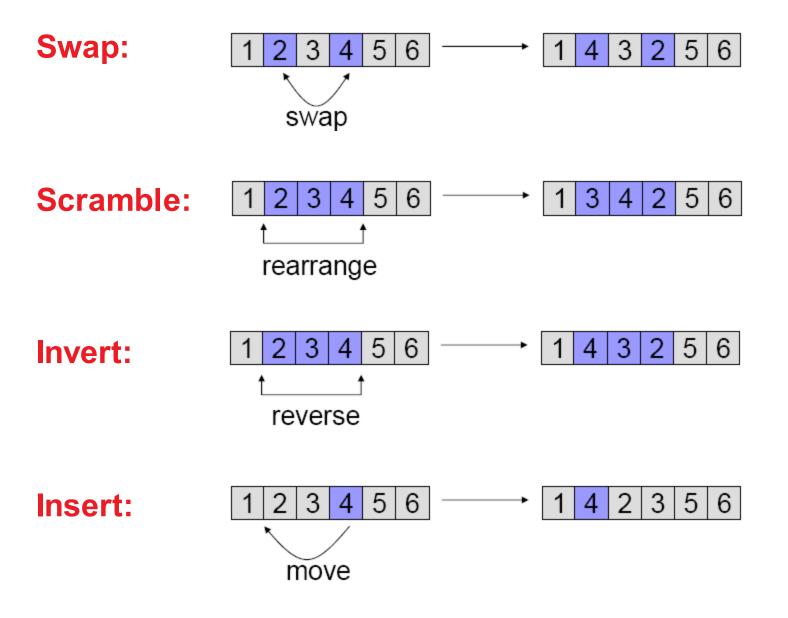
- every solution can be generation from every other with a probability greater than 0 ("exhaustiveness")
- $d(x, x') < d(x, x'') \Longrightarrow Prob(mut(x) = x') > Prob(mut(x) = x'')$ ("locality")

Desirable property of recombination operators ("in-between-ness"): $x'' = \operatorname{recomb}(x, x') \Rightarrow d(x'', x) \leq d(x, x') \land d(x'', x') \leq d(x, x')$



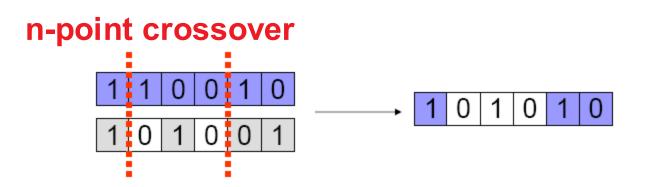
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Examples of Mutation Operators on Permutations

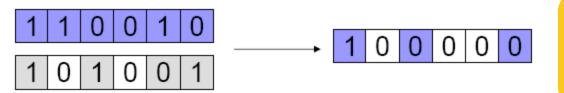


Examples of Recombination Operators: {0,1}ⁿ

1-point crossover 1 1 0 0 1 0 1 0 1 0 0 1 → 1 1 0 0 0 1



uniform crossover



choose each bit independently from one parent or another

- binary search space, maximization
- uniform initialization
- generational cycle: of the population
 - evaluation of solutions
 - mating selection (e.g. roulette wheel)
 - crossover (e.g. 1-point)
 - environmental selection (e.g. plus-selection)

Conclusions

- EAs are generic algorithms (randomized search heuristics, meta-heuristics, ...) for black box optimization
 no or almost no assumptions on the objective function
- They are typically less efficient than problem-specific (exact) algorithms (in terms of #funevals)
 less differences in the continuous case (as we have seen)
- Allow for an easy and rapid implementation and therefore to find good solutions fast

easy to incorporate (and recommended!) to incorporate problem-specific knowledge to improve the algorithm

I hope it became clear...

...that heuristics is what we typically can afford in practice (no guarantees and no proofs) ...what are the main ideas behind evolutionary algorithms ...and that evolutionary algorithms and genetic algorithms are no synonyms