

# Introduction to Optimization

## Approximation Algorithms and Heuristics

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# Exercise:

## The Knapsack Problem and Dynamic Programming

`http://researchers.lille.inria.fr/  
~brockhof/optimizationSaclay/`

# Conclusions

I hope it became clear...

- ...what the algorithm design idea of **dynamic programming** is
- ...for which problem types is is supposed to be **suitable**
- ...and how to **apply** the idea to the **knapsack problem**

# Course Overview

Date		Topic
Mon, 21.9.2015		Introduction
Mon, 28.9.2015	D	Basic Flavors of Complexity Theory
Mon, 5.10.2015	D	Greedy algorithms
Mon, 12.10.2015	D	Branch and bound (switched w/ dynamic programming)
Mon, 2.11.2015	D	Dynamic programming [ <i>salle Proto</i> ]
<b>Fri, 6.11.2015</b>	<b>D</b>	<b>Approximation algorithms and heuristics [S205/S207]</b>
Mon, 9.11.2015	C	Introduction to Continuous Optimization I [ <i>S118</i> ]
Fri, 13.11.2015	C	Introduction to Continuous Optimization II <i>[from here onwards always: S205/S207]</i>
Fri, 20.11.2015	C	Gradient-based Algorithms
Fri, 27.11.2015	C	End of Gradient-based Algorithms + Linear Programming
Fri, 4.12.2015	C	Stochastic Optimization and Derivative Free Optimization
<b>Tue, 15.12.2015</b>		Exam

all classes + exam last 3 hours (incl. a 15min break)

# Overview of Today's Lecture

## Approximation Algorithms

- a greedy approximation algorithm for bin packing
- an FPTAS for the KP

## Overview of (Randomized) Search Heuristics

- randomized local search
- variable neighborhood search
- tabu search
- evolutionary algorithms
- [exercise: simple randomized algorithms for the knapsack problem]

# Approximation Algorithms

# 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 (now)

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

## Heuristics (later today)

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

# Approximations, PTAS, and FPTAS

- An algorithm is a  *$\rho$ -approximation algorithm* for problem  $\Pi$  if, for each problem instance of  $\Pi$ , it outputs a feasible solution which function value is within a ratio  $\rho$  of the true optimum for that instance.
- An algorithm  $A$  is an *approximation scheme* for a *minimization\** problem  $\Pi$  if for any instance  $I$  of  $\Pi$  and a parameter  $\varepsilon > 0$ , it outputs a solution  $s$  with  $f_{\Pi}(I, s) \leq (1 + \varepsilon) \cdot \text{OPT}$ .
- 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  $1/\varepsilon$  actually!
- An approximation scheme is a *fully polynomial time approximation scheme (FPTAS)* if its runtime is bounded polynomially in both the size of the input  $I$  and in  $1/\varepsilon$ .

# Today's Lecture

- only example algorithm(s)
- no detailed proofs here due to the time restrictions of the class
- rather spend more time on the heuristics part and the exercise

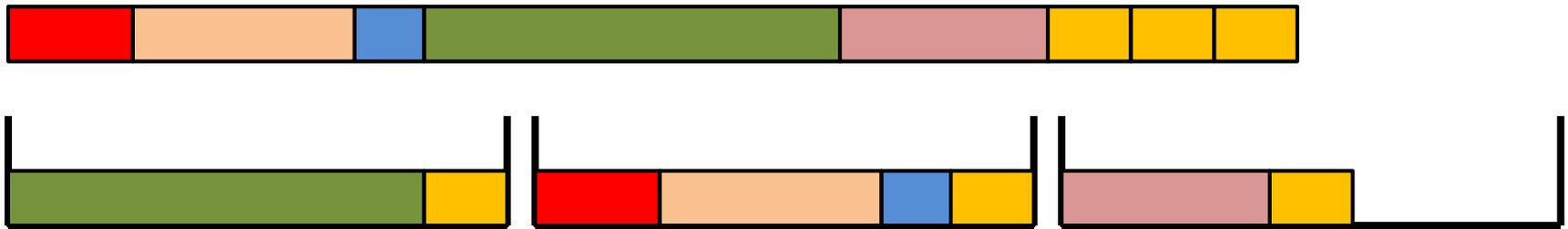
## Actually Two Examples:

- a greedy approximation algorithm for bin packing
- an FPTAS for the KP

# Bin Packing (BP)

## Bin Packing Problem

Given a set of  $n$  items with sizes  $a_1, a_2, \dots, a_n$ . 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  $\leq V$ .



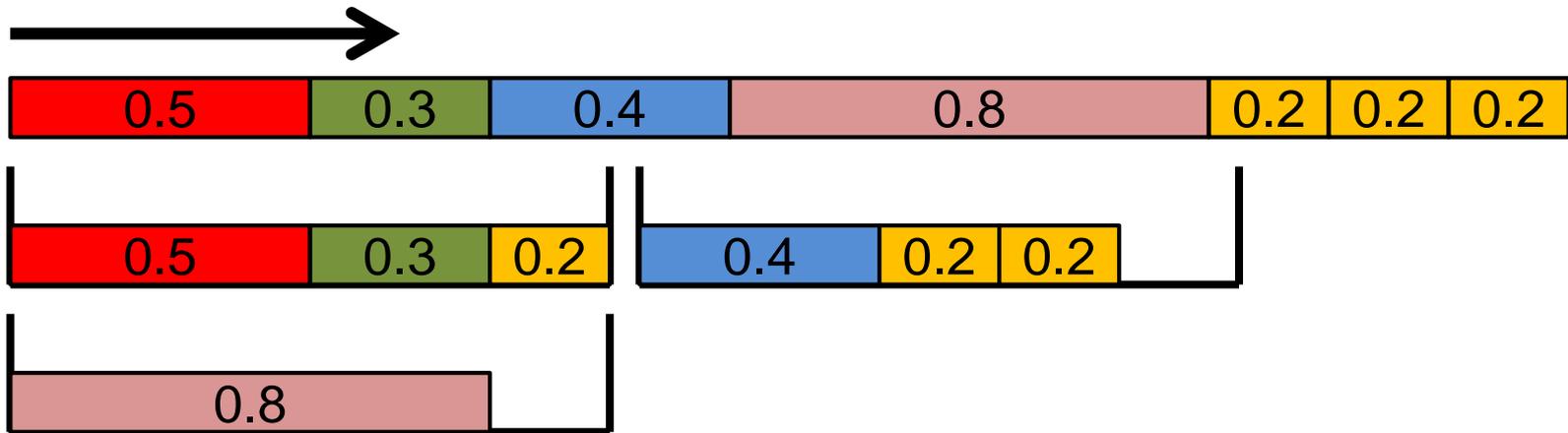
## Known Facts

- 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  $\rho$ -ratio of 2

# First-Fit Approach

## 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).

$$\text{OPT} > \frac{m-1}{2} \iff 2\text{OPT} > m-1 \implies 2\text{OPT} \geq m$$

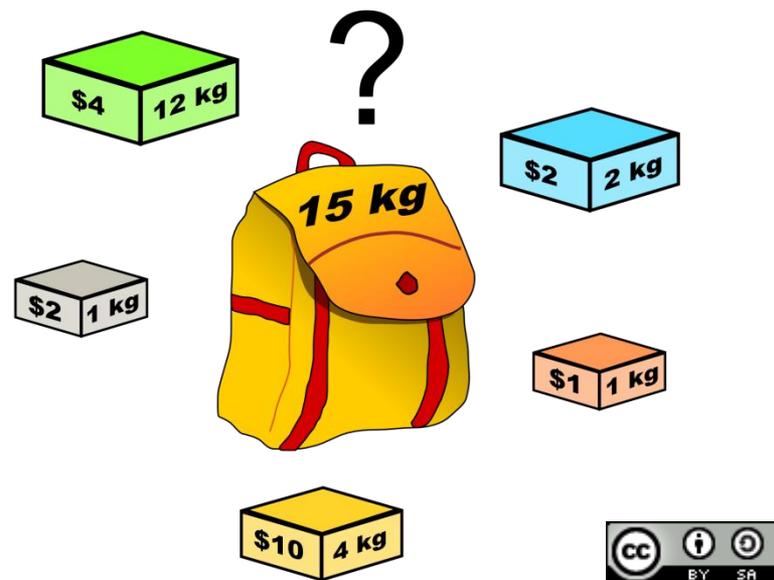
↑ because  $m$  and  $\text{OPT}$  are integer

# An FPTAS for the Knapsack Problem

## Knapsack Problem

$$\max. \sum_{j=1}^n p_j x_j \text{ with } x_j \in \{0, 1\}$$

$$\text{s.t. } \sum_{j=1}^n w_j x_j \leq W$$



# An FPTAS for the Knapsack Problem

Similar to this week's exercise, we can design a dynamic programming algorithm for which

- a subproblem is restricting the items to  $\{1, \dots, k\}$  and searches for the lightest packing with prefixed profit  $P$
- runs in  $O(n^2 P_{\max})$

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

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

**Idea behind FPTAS:**

- scale the profits smartly to  $\left\lfloor \frac{p_i n}{\varepsilon P_{\max}} \right\rfloor$  to make  $P_{\max}$  polynomially bounded
- prove that dynamic programming approach computes profit of at least  $(1-\varepsilon) \cdot \text{OPT}$  (not shown here)

# **(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 Overview

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

# Neighborhoods

For most (stochastic) search heuristics in discrete domain, 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 ( $\Omega = \{0,1\}^n$ )
- 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

# 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 (not dependent on obj. function)

## First Improvement Local Search, Randomized Local Search (RLS):

- 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

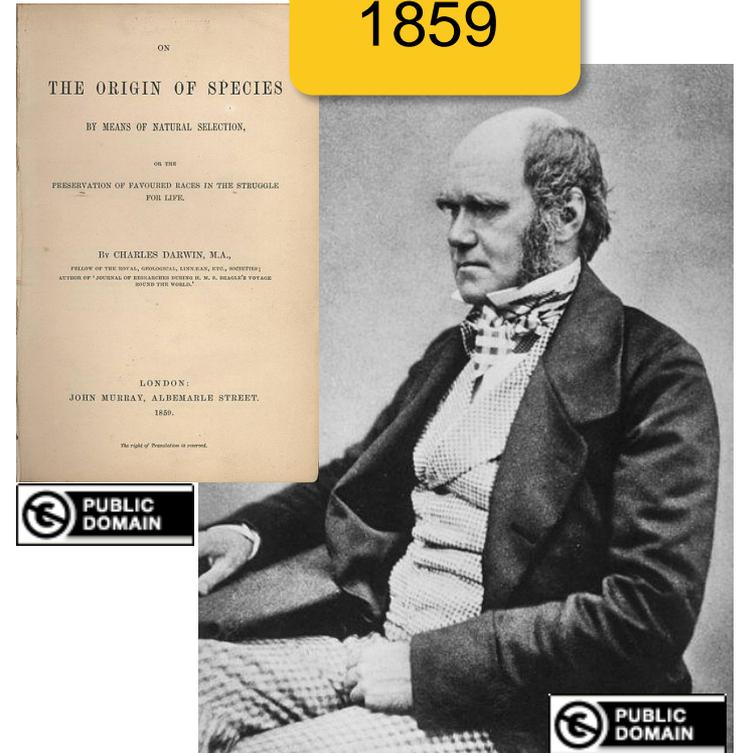
- allowing worsening moves if all neighbors are explored
- introducing a tabu list of temporarily not allowed moves
- those restricted moves are
  - problem-specific and
  - can be specific solutions or not permitted “search directions” such as “don’t include this edge anymore” or “do not flip this specific 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

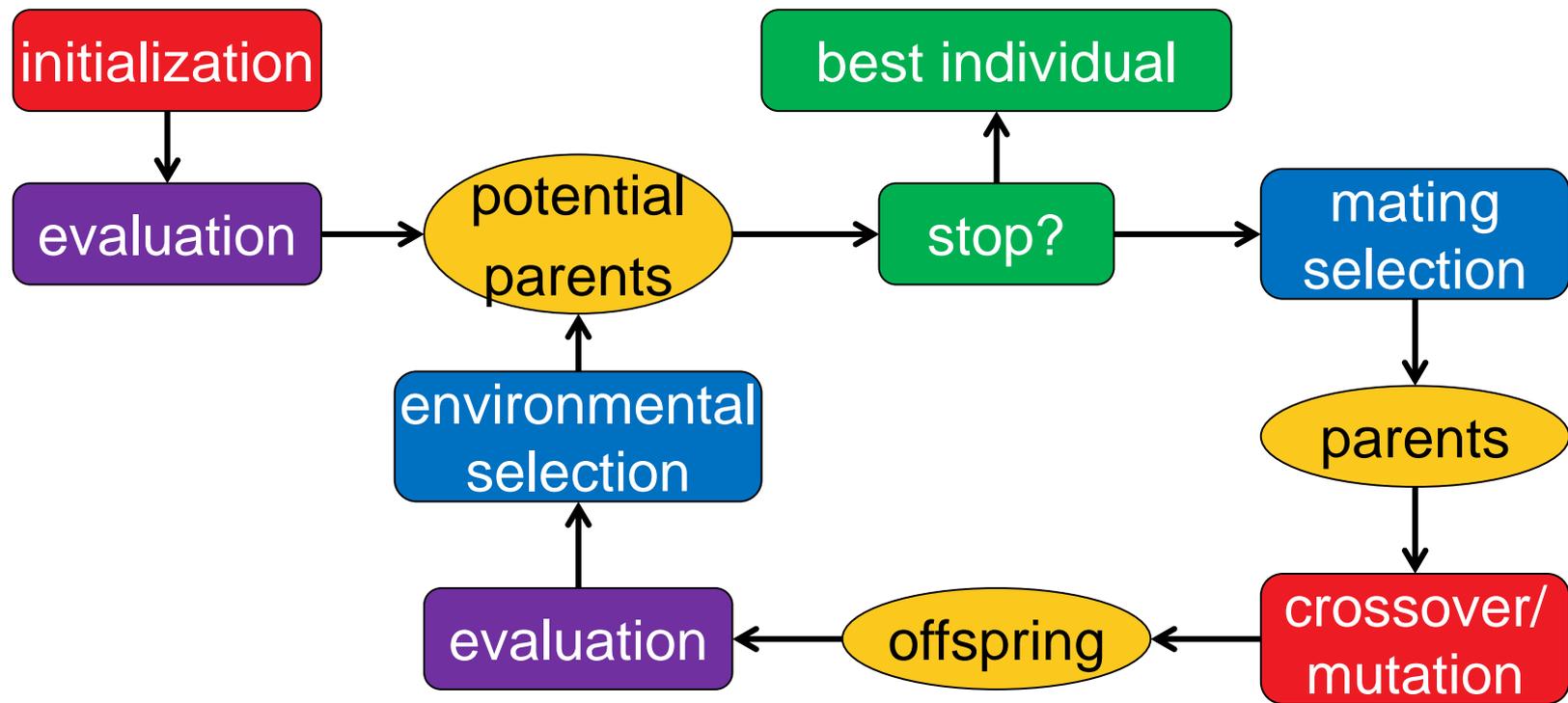
1859



# 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)

# 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)*

$$\Omega = \mathbb{R}^n$$

## Genetic Programming (GP)

*J. Koza 1990 (USA)*

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

nowadays one umbrella term: **evolutionary algorithms**

# Genotype – Phenotype mapping

## The genotype – phenotype mapping

- related to the question: how to come up with a fitness 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, see also the Lagrangian in the continuous part of the lecture)
- **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
- some more...

# **Examples for some EA parts**

## **(for discrete search spaces)**

# Selection

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

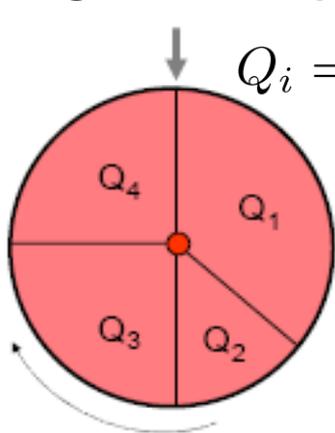
Selection is either

**stochastic**

or

**deterministic**

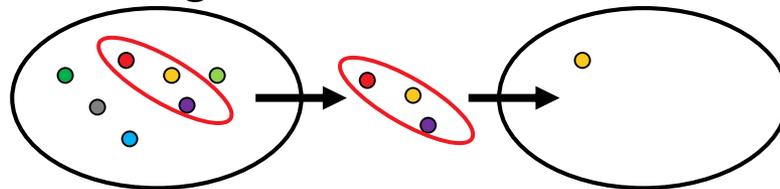
e.g. fitness proportional



$$Q_i = \frac{f(x_i)}{\sum_{j=1}^{\mu} f(x_j)}$$

**Disadvantage:**  
depends on  
scaling of  $f$

e.g. via a tournament



e.g.  $(\mu+\lambda)$ ,  $(\mu, \lambda)$



**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:**  $mut: \Omega \rightarrow \Omega$

**recombination:**  $recomb: \Omega^r \rightarrow \Omega^s$  where  $r \geq 2$  and  $s \geq 1$

- 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.

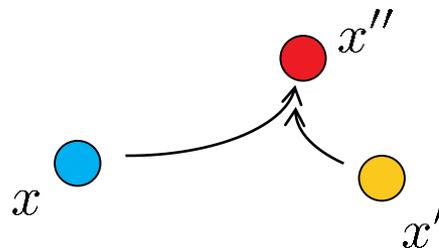
# Variation Operators: Guidelines

Two desirable properties for **mutation** operators:

- every solution can be generation from every other with a probability greater than 0 (“exhaustiveness”)
- $d(x, x') < d(x, x'') \Rightarrow Prob(\text{mut}(x) = x') > Prob(\text{mut}(x) = x'')$  (“locality”)

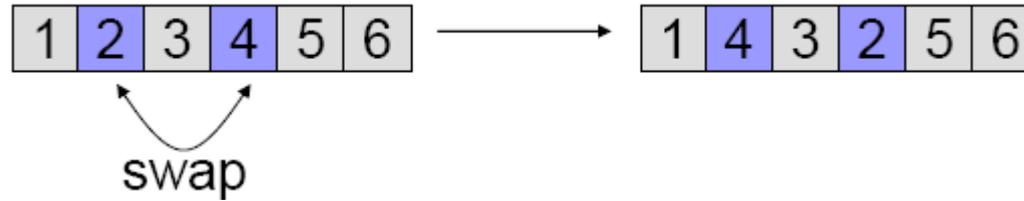
Desirable property of **recombination** operators (“in-between-ness”):

$$x'' = \text{recomb}(x, x') \Rightarrow d(x'', x) \leq d(x, x') \wedge d(x'', x') \leq d(x, x')$$

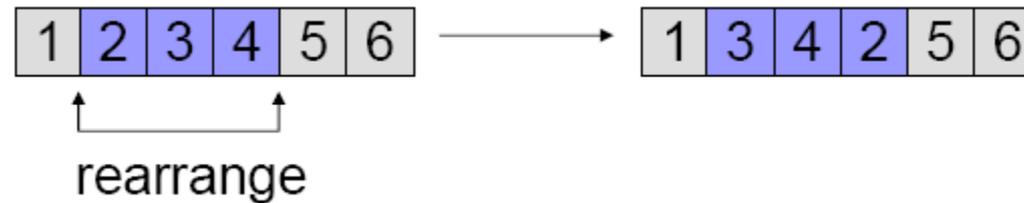


# Examples of Mutation Operators on Permutations

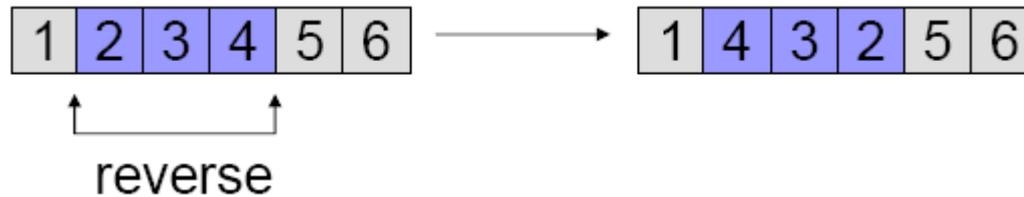
**Swap:**



**Scramble:**



**Invert:**

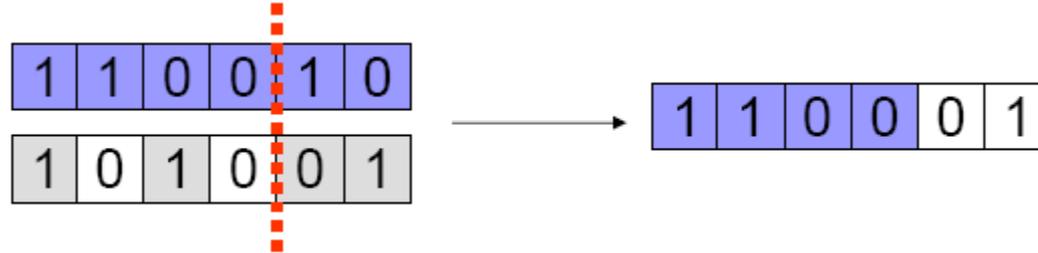


**Insert:**

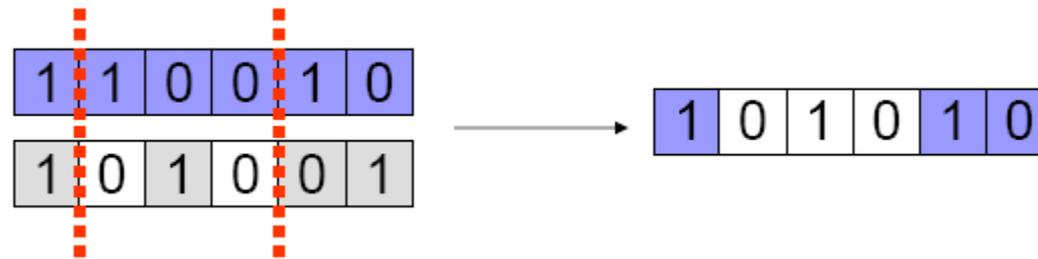


# Examples of Recombination Operators: $\{0,1\}^n$

## 1-point crossover



## n-point crossover



## uniform crossover



choose each bit independently from one parent or another

# A Canonical Genetic Algorithm

- 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)  
*not the case in the continuous case (we will see later)*
- Allow for an easy and rapid implementation and therefore to find good solutions fast  
*easy to incorporate problem-specific knowledge to improve the algorithm*

# Conclusions

I hope it became clear...

...that **approximation algorithms** are often what we can hope for in practice (might be difficult to achieve guarantees though)

...that **heuristics** is what we typically can afford in practice (no guarantees and no proofs)