

# Parameterized Complexity Analysis of Evolutionary Algorithms

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## Introduction

### Our task

- Given a **function**  $f: X \rightarrow \mathbb{R}$
- and a set  $D \subseteq X$  of **feasible solutions**,
- find  $\arg \max_{x \in D} f(x)$ .

We are interested in **general purpose algorithms** that can be applied without problem knowledge

Parameterized Complexity Analysis of EAs

## Introduction

### Why General Purpose Algorithms?

- **Algorithms** are the **heart** of every **nontrivial computer application**.
- For **many problems** we know good or **optimal algorithms**.
  - Sorting
  - Shortest paths
  - Minimum spanning trees
- What about **new or complex problems**?
- Often there are **no good problem specific algorithms**.

Parameterized Complexity Analysis of EAs

## Introduction

### Points that may rule out problem specific algorithms

- Problems that are **rarely understood**.
- **Quality of solutions** is determined by **simulations**.
- Problems that fall into the **black box scenario**.
- **Not enough resources** such as time, money, knowledge.

General purpose algorithms are often a good choice.

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## Introduction

**General purpose algorithms** for optimizing a function  $f: X \rightarrow \mathbb{R}$

1. Choose a **representation** for the elements in  $X$ .
2. Fix a **function** to evaluate the quality (might be different from  $f$ ).
3. Define **operators** that produce new elements.

## Evolutionary Algorithms

Evolutionary algorithms are **general purpose algorithms**.

Follow Darwin's principle (**survival of the fittest**).

Work with a set of solutions called **population**.

Parent population produces offspring population by **variation operators** (mutation, crossover).

**Select** individuals from the parents and children to create a new parent population.

**Iterate** the process until a "**good solution**" has been found.

## Simple Evolutionary Algorithm

### (1+1) EA

```
 $x \leftarrow$  an element of  $\{0, 1\}^n$  uniformly at random.  
repeat forever  
  Produce  $y$  by flipping each bit of  $x$  with probability  $1/n$ .  
  if  $f(y) \geq f(x)$  then  $x \leftarrow y$ 
```

## Theory of Evolutionary Algorithms

Evolutionary algorithms are **successful** for many complex optimization problems.

Rely on **random decisions**  $\Rightarrow$  **randomized algorithms**.

Goal: understand **how** and **why** they work.

Study the **computational complexity** of these algorithms on prominent examples.

## Runtime analysis

### Black box scenario

- Measure the runtime  $T$  by the number of fitness evaluations.
- Studies consider time in dependence of the input to reach
  - An optimal solution
  - A good approximation

### Rigorous estimates

- Expected number of fitness evaluations  $E(T)$
- Tail bounds, e.g., useful bounds on  $\Pr(T \leq g(n))$  where  $n$  measures the size of the problem instance

## Motivation

We want to tackle the analysis of randomized search heuristics applied to NP-hard problems

- Reducing the base of the exponent
  - Conflict-directed walk (Schöning, 1999)  $O(1.334^n)$  for 3-SAT
- Polynomial-time approximation
  - Partition (Witt, 2005)
  - Vertex Cover (Friedrich et al., 2007, Oliveto et al., 2007)
  - Set Cover (Friedrich et al., 2007)
  - Intersection of  $p \geq 3$  matroids (Reichel & Skutella, 2010)
- Average-case analysis
  - Partition (Witt, 2005)

Classical view: understand runtime as a function of problem size alone.

**Real world problems:** inputs are often structured or restricted in some way.

## Motivation

### Type-checking in ML

- No explicit typing in ML: compiler must infer types/check for consistency: complete for EXPTIME
- Let  $k$  be the nesting depth of a type declaration, there is an exact algorithm that solves the problem in  $O(2^k n)$
- In most real-world problems:  $k \leq 10$

## Motivation

### Reconfigurable computing

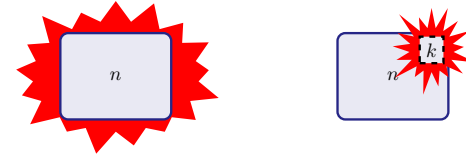
- Given:
  - an  $n \times n$  memory array  $\mathbf{A}$  with some defective elements
  - $k_1$  extra rows of spare memory,  $k_2$  extra columns of spare memory
- A defective element can be *repaired* by replacing the row/column that contains it with a spare row/column
- Determine if there is a replacement arrangement that repairs all defective elements in  $\mathbf{A}$
- Reduces to: constrained minimum vertex cover in bipartite graphs: NP-complete
- Chen & Kanj (2003):  $O(1.26^k n)$  algorithm where  $k = k_1 + k_2$
- In the real world, due to hardware constraints,  $k \leq 40$

## Motivation

Many heuristics are successful in practice because they can take advantage of problem structure...  
...want analyses to capture that

## Parameterized complexity

Find a hardness parameter  $k$  that isolates the source of exponential complexity.



Let  $L$  be a language over a finite alphabet  $\Sigma$ .

A parameterization of  $L$  is a mapping  $\kappa : \Sigma^* \rightarrow \mathbb{N}$

Corresponding parameterized problem is given by  $(L, \kappa)$ .

For a string  $x \in \Sigma^*$ , let  $k = \kappa(x)$  and  $n = |x|$ .

An algorithm deciding  $x \in L$  in the time bounded by  $f(k) \cdot \text{poly}(n)$  is called a fixed-parameter tractable (FPT) algorithm for the parameterization  $\kappa$ .

## Parameterized complexity for EAs

Monte-Carlo FPT algorithm: in FPT-time, accept with probability at least  $1/2$  if  $x \in L$ , with probability 0 if  $x \notin L$ .

### Definition

An evolutionary algorithm is called *fixed-parameter tractable* (FPT) if it finds an optimal solution in expected time  $O(f(k) \cdot \text{poly}(n))$ .

- vertex cover (Kratsch and Neumann, 2013)
- maximum leaf spanning tree (Kratsch et al., 2010)
- MAX-2-SAT (Sutton, Day, Neumann, GECCO 2012)
- Makespan scheduling (Sutton and Neumann, PPSN 2012)
- Euclidean TSP (Nallaperuma et al., CEC 2013)
- Bilevel optimization (Corus, Lehre, and Neumann, GECCO 2013)
- Hypervolume indicator (Bringmann and Friedrich, GECCO 2013)

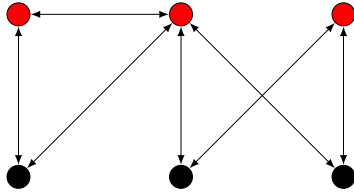
## The Minimum Vertex Problem

Friedrich, He, Hebbinghaus, Neumann and Witt (ECJ 2010)  
Kratsch and Neumann (Algorithmica 2013)

## The Problem

The Vertex Cover Problem:

Given an undirected graph  $G = (V, E)$



Find a minimum subset of vertices such that each edge is covered at least once.

NP-hard, several 2-approximation algorithms

Simple, single-objective evolutionary algorithms fail!!!

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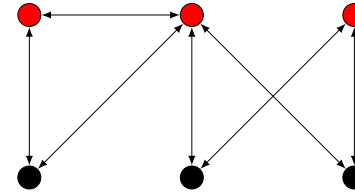
## The Problem

**Integer Linear Program (ILP)**

$$\begin{aligned} \min & \sum_{i=1}^n x_i \\ \text{s.t. } & x_i + x_j \geq 1 \quad \forall \{i, j\} \in E \\ & x_i \in \{0, 1\} \end{aligned}$$

**Linear Program (LP)**

$$\begin{aligned} \min & \sum_{i=1}^n x_i \\ \text{s.t. } & x_i + x_j \geq 1 \quad \forall \{i, j\} \in E \\ & x_i \in [0, 1] \end{aligned}$$



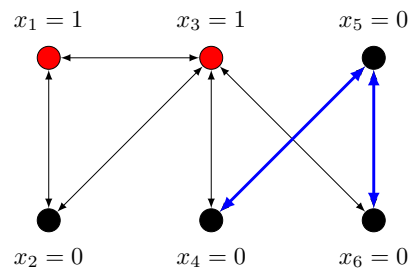
**Decision problem:** is there a set of vertices of size at most  $k$  covering all the edges?

**Our parameter:** value of an optimal solution (OPT)

Parameterized Complexity Analysis of EAs

## Evolutionary Algorithm

**Representation:** bitstrings of length  $n$



**Minimize fitness function:**

$$\begin{aligned} f_1(x) &= (|x|_1, |U(x)|) \\ f_1(x) &= (2, 2) \end{aligned}$$

$$\begin{aligned} f_2(x) &= (|x|_1, LP(x)) \\ f_2(x) &= (2, 1) \end{aligned}$$

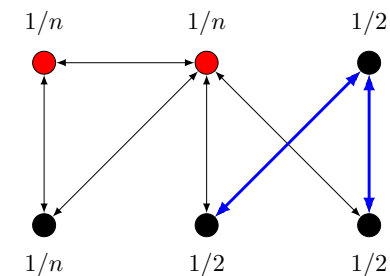
$U(x) :=$  edges not covered by  $x$

$G(x) := G[U(x)]$

$LP(x) :=$  the value of LP applied to  $G[U(x)]$

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## Evolutionary Algorithm



**Two mutation operations:**

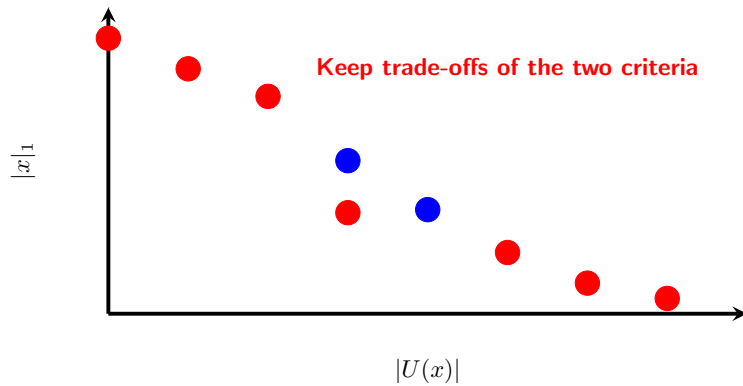
1. Standard bit mutation with probability  $1/n$
2. Mutation probability  $1/2$  for vertices adjacent to edges of  $U(x)$ . Otherwise, mutation probability  $1/n$ .

**Decide uniformly at random which operator to use in each iteration**

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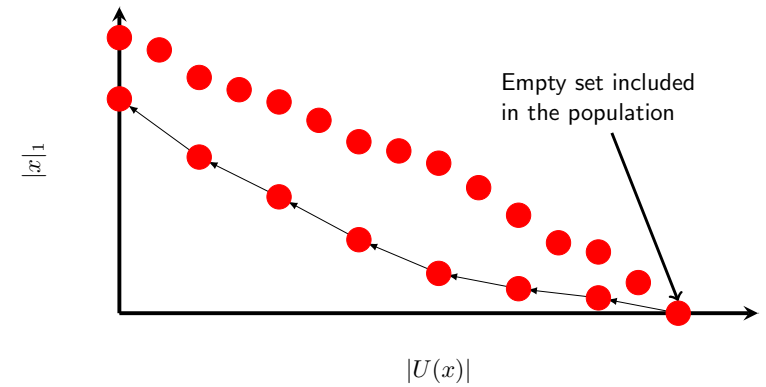
## Multi-objective approach

Treat the different objectives in the same way



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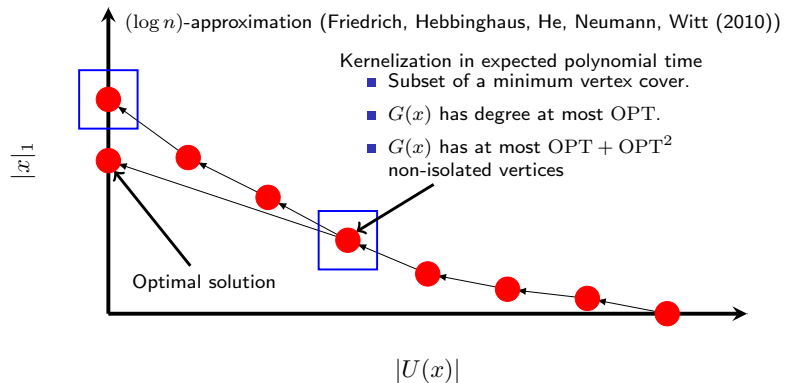
## Multi-objective approach



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## Multi-objective approach

What can we say about these solutions?

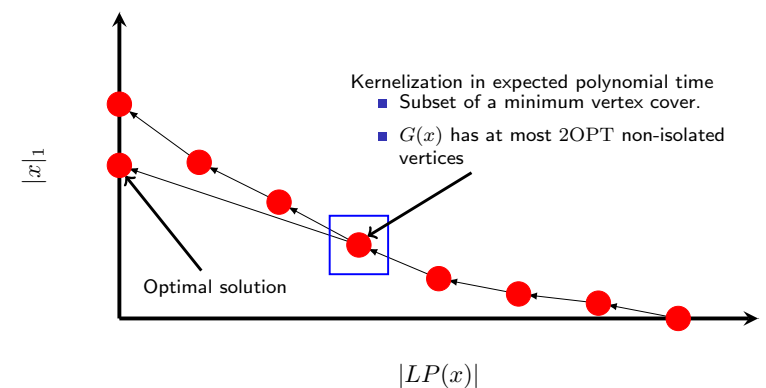


Expected time  $g(\text{OPT}) \cdot \text{poly}(n)$

**Fixed-parameter evolutionary algorithm**

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## Multi-objective approach



Expected time  $4^{\text{OPT}} \cdot \text{poly}(n)$

**Fixed-parameter evolutionary algorithm**

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## Linear programming

### Combination with linear programming

LP-relaxation is half integral, i.e.,

$$x_i \in \{0, 1/2, 1\}, \quad 1 \leq i \leq n.$$

### Theorem (Nemhause, Trotter (1975)):

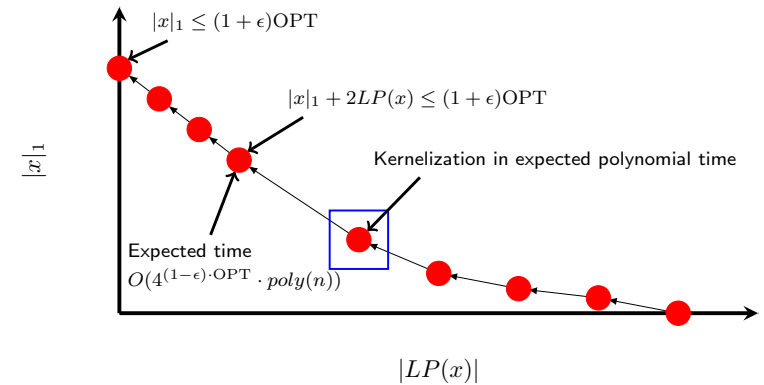
Let  $x^*$  be an optimal solution of the LP. Then there is a minimum vertex cover that contains all vertices  $v_i$  where  $x_i^* = 1$ .

### Lemma:

All search points  $x$  with  $LP(x) = LP(0^n) - |x|_1$  are Pareto optimal. They can be extended to minimum vertex cover by selecting additional vertices.

Can we also say something about approximations?

## Approximations



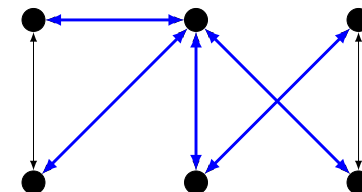
## Maximum Leaf Spanning Trees

Kratsch, Lehre, Neumann and Oliveto (PPSN 2010)

## The Problem

The Maximum Leaf Spanning Tree Problem:

Given an undirected connected graph  $G = (V, E)$ ,



find a spanning tree with a maximum number of leaves.

NP-hard, different classical FPT-studies.

## Two Evolutionary Algorithms

### Algorithm 1: Generic (1+1) EA

Choose a spanning tree  $T$  of  $G$  uniformly at random  
**repeat** forever  
     Produce  $T'$  by swapping each edge of  $T$  indep. w/ prob.  $1/m$   
     **if**  $T'$  is a tree and  $\ell(T') \geq \ell(T)$  **then**  $T \leftarrow T'$

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## Two Evolutionary Algorithms

### Algorithm 2: Tree-based (1+1) EA

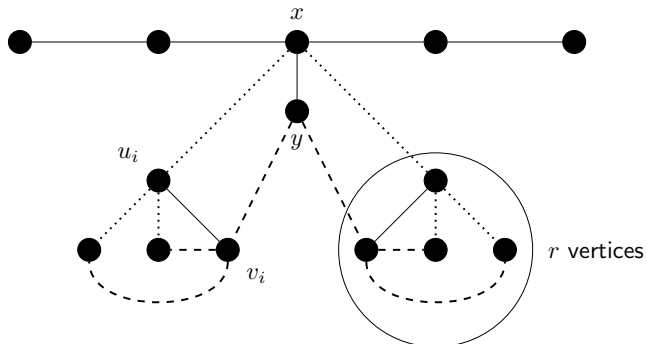
Choose a spanning tree  $T$  of  $G$  uniformly at random.  
**repeat** forever  
     Choose  $s \sim \text{Pois}(\lambda = 1)$   
     Produce  $T'$  by performing sequentially  $s$  random edge-exchange operations.  
     **if**  $\ell(T') \geq \ell(T)$  **then**  $T \leftarrow T'$

A random exchange operation applied to a spanning tree  $\tilde{T}$  chooses an edge  $e \in E \setminus \tilde{T}$  uniformly at random. The edge  $e$  is inserted and one randomly chosen edge of the cycle in  $\tilde{T} \cup \{e\}$  is deleted.

**Does the mutation operator make the difference between FPT and non-FPT runtime?**

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## Local Optimum



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## Lower Bounds

### Theorem 1.

The expected optimization time of Generic (1+1) EA on  $G_{loc}$  is lower bounded by  $(\frac{m}{c})^{2(r-2)}$  where  $c$  is an appropriate constant.

### Theorem 2.

The expected optimization time of Tree-Based (1+1) EA on  $G_{loc}$  is lower bounded by  $(\frac{r-2}{c})^{r-2}$  where  $c$  is an appropriate constant.

### Idea for lower bounds:

- Both algorithms may get stuck in local optimum.
- For the Generic (1+1) EA it is less likely to escape the local optimum as it often flips edges on the path.

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## Structural insights

Similar to Fellows, Lokshatanov, Misra, Mnich, Rosamond, Saurabh (2009)

### Lemma 2.

Any connected graph  $G$  on  $n$  nodes and with a maximum number of  $k$  leaves in any spanning tree has at most  $n + 5k^2 - 7k$  edges and at most  $10k - 14$  nodes of degree at least three.

#### Proof idea:

- Let  $T$  be a maximum leaf spanning tree with  $k$  leaves
- Let  $P_0$  be the set of all leaves and all nodes of degree at least three in  $T$
- Let  $P$  be the set of nodes that are of distance at most 2 (w.r.t.  $T$ ) to any node in  $P_0$  and let  $Q$  be the set of remaining nodes.
- **Show:** all nodes of  $Q$  have degree 2 in  $G$ .
- **Implies:** number of nodes in  $P$  is at most  $10k - 14$ .
- **No node has degree greater than  $k$** , which implies bound on number of edges.

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## Upper Bound

### Theorem 3.

If the maximal number of leaf nodes in any spanning tree of  $G$  is  $k$ , then Tree-Based (1+1) EA finds an optimal solution in expected time  $O(2^{15k^2 \log k})$ .

#### Proof idea:

- We call an edge **distinguished** if it is adjacent to at least one node of degree at least 3 in  $G$ .
- **Number of distinguished edges** on any cycle is at most  $20k - 28$ .
- Total number of edges in  $G$ :  $m \leq n + 5k^2 - 7k$ .
- Probability to introduce a specific non-chosen distinguished edge is at least  $1/(m - (n - 1)) \geq 1/5k^2$ .
- **Show:** length of created cycle is at most  $20k$ .
- Probability to remove edge of the cycle that does not belong to optimal solution is at least  $1/20k$ .

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## Proof of Upper Bound (continued)

- Probability to obtain a specific spanning tree that can be obtained by an edge-swap is at least  $1/(20k \cdot 5k^2)$ .
- Probability to produce optimal spanning tree, which has distance  $r \leq 5k^2$ , is at least

$$r! \cdot \frac{1}{er!} \left( \frac{1}{5k^2} \cdot \frac{1}{20k} \right)^r \geq \frac{1}{e} \left( \frac{1}{100k^3} \right)^{5k^2} \geq \frac{1}{2} \left( \frac{1}{100} \right)^{5k^2} \left( \frac{1}{k} \right)^{3 \cdot 5k^2},$$

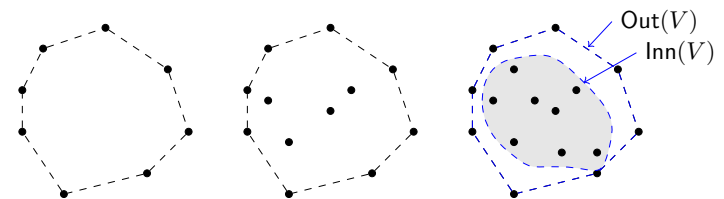
- Implies that the expected time to get the maximum leaf spanning tree is at most  $O(2^{15k^2 \log k})$ .  $\square$

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## Euclidean Planar TSP

Given a set  $V$  of  $n$  points in the plane, find a Hamiltonian cycle of minimum length (NP-hard, Papadimitriou, 1977)

Deĭneko et al. (2006): dynamic programming (simple polygon,  $k$  inner points)



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## TSP parameterization

### How does this structure affect evolutionary algorithms?

For  $n$  points in the plane with  $|\text{Inn}(V)| = k$  interior to the convex hull, what is the runtime of an EA in terms of  $n$  and  $k$ ?

Each tour is represented by a permutation  $\pi : V \rightarrow V$ .

$$v_{\pi(1)} \Rightarrow v_{\pi(2)} \Rightarrow \dots \Rightarrow v_{\pi(n)} \Rightarrow v_{\pi(1)}$$

### Fitness function

$$f(\pi) = \left( \sum_{i=1}^{n-1} d(v_{\pi(i)}, v_{\pi(i+1)}) \right) + d(v_{\pi(n)}, v_{\pi(1)})$$

where  $d(u, v)$  is the distance between points  $u$  and  $v$ .

## TSP parameterization

**Main structural idea:** An optimal tour does not intersect itself.

### Lemma

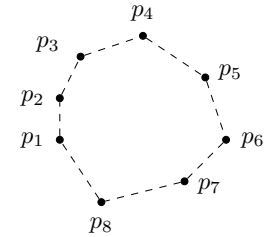
Suppose  $\pi^*$  is a permutation that minimizes  $f$ . Then the elements of  $\text{Out}(V)$  appear in  $\pi^*$  in the **same** order they appear on the hull.

### Definition

We define  $\gamma$  as a linear order on  $\text{Out}(V)$

$$\gamma = (p_1, p_2, \dots, p_{n-k})$$

such that for all  $i \in \{1, \dots, n-k\}$ ,  $p_i$  and  $p_{i+1}$  are adjacent on the boundary of the convex hull of  $V$ .



**For any  $V$ ,  $\gamma$  can be computed in  $O(n \log n)$  time**

## TSP parameterization

### Definition

A permutation  $\pi$  on a subset  $S$  of  $V$  is  $\gamma$ -**respecting** if and only if, for any  $p_i, p_j \in \gamma \cap S$ ,

$$\pi^{-1}(p_i) < \pi^{-1}(p_j) \implies i < j.$$

where  $\gamma \cap S$  means the restriction of  $\gamma$  to  $S$ .

### Some examples...

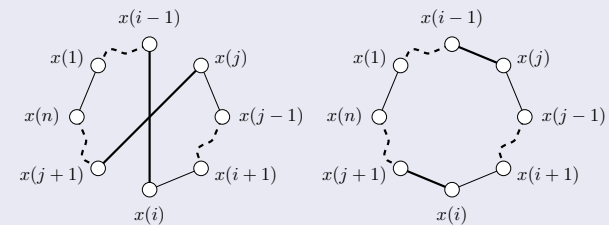
$$(p_1, v_4, v_6, p_2, v_1, v_3, p_3, p_4, v_2, p_5, v_7, p_6, p_7, v_5)$$

$$(v_7, v_5, p_1, v_4, p_2, v_2, v_6, p_3, p_4, p_5, v_1, v_3, p_6, p_7)$$

## (1+1) EA in the black-box setting

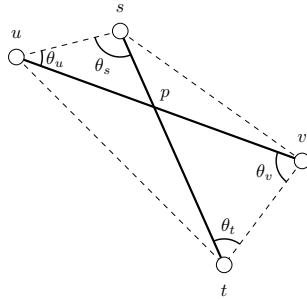
We start in the **black-box** setting (EA has no access to instance structure)

No crossover, mutation is by **edge-exchange** operations, e.g., 2-opt:



## (1+1) EA in the black-box setting

Improvement in fitness if sum of edge lengths of new edges is strictly less than sum of edge lengths of old edges.



**Main challenge with edge-exchange operations:** if angles can be arbitrarily close, fitness improvements can be arbitrarily small.

**Idea:** assume the angles are **bounded** (or embedded on an  $m \times m$  grid).

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## (1+1) EA in the black-box setting

### Theorem

Given a set of (angle bounded) points, a (1+1) EA solves the Euclidean TSP with  $k$  inner points in expected time  $O(n^{4k}(2k-1)!)$ .

### Proof idea.

If a tour has edges that cross, an improving move is possible.

With appropriate angle bounds, EA spends  $\text{poly}(n)$  time on such tours (independent of  $k$ ).

If the tour has no edges that cross, then it is  $\gamma$ -respecting.

$\gamma$ -respecting tours are *closer* to optimal tours: the EA only must operate on the inner points to find a solution.

Time to fix inner points:  $O(n^{4k}(2k-1)!)$ .

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## $(\mu+1)$ EA

### FPT evolutionary algorithms (we leave the black-box setting)

FPT  $(\mu+1)$  EA: based on exact  $(\mu+1)$  EA for TSP by Theile (2009)

Population of permutations on *subsets* of  $V$  with *special structure*

### Ground set

- An integer  $i \in \{1, \dots, n-k\}$
- A set  $S \subseteq \text{Inn}(V)$
- A vertex  $r \in S \cup \{p_i\}$

we identify  $(i, S, r)$  with the set  $S \cup \{p_1, \dots, p_i\}$  distinguished by  $r$

An individual  $\pi = \pi_{(i, S, r)}$  is a permutation on the ground set  $S \cup \{p_1, p_2, \dots, p_i\}$  and a “tail” vertex  $r$  where

- $\pi(1) = p_1$  and  $\pi(|S| + i) = r$ ,
- $\pi$  is  $\gamma$ -respecting that is,  $p_1, p_2, \dots, p_i$  appear in order.

Parameterized Complexity Analysis of EAs

## $(\mu+1)$ EA

The subtour defined by a permutation  $\pi_{(i, S, r)}$ :

$$p_1 \Rightarrow v_{\pi(2)} \Rightarrow \dots \Rightarrow v_{\pi(|S|+i-1)} \Rightarrow r \Rightarrow p_1$$

- starts at  $p_1$ ,
- runs over all nodes in  $(S \cup \{p_2, \dots, p_i\}) \setminus r$  (respecting  $\gamma$ )
- finally visits  $r$  before returning to  $p_1$ .

Full population consists of an individual for every  $i \in \{1, \dots, n-k\}$ , for every  $S \subseteq \text{Inn}(V)$ , and every possible tail vertex  $r$ , given  $S$  and  $i$ .

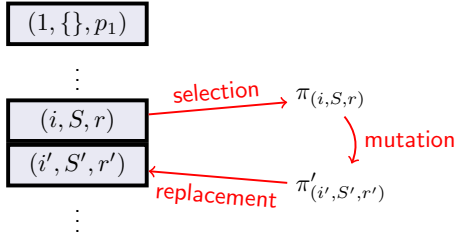
The fitness of an individual is the cost of the corresponding subtour

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## $(\mu+1)$ EA

### $(\mu+1)$ EA

- Maintain a population  $P$  such that each ground-set/tail vertex combination  $(i, S, r)$  is represented exactly once.
- While optimal tour is not in  $P$ 
  - Select an individual  $\pi_{(i, S, r)} \in P$  uniformly at random
  - Mutate  $\pi_{(i, S, r)}$  to produce  $\pi_{(i', S', r')}$  (mutation extends the ground set, and only creates  $\gamma$ -respecting permutations).
  - If the fitness of the mutant  $\pi_{(i', S', r')}$  is better than the current individual representing  $(i', S', r')$ , then replace that individual with the mutant.



Parameterized Complexity Analysis of EAs

## $(\mu+1)$ EA

### Mutation

To mutate a single individual  $\pi = \pi_{(i, S, r)}$ ,

- choose  $v$  uniformly at random from  $(\text{Inn}(V) \setminus S) \cup \{p_{i+1}\}$
- concatenate  $v$  to the linear order described by  $\pi$ . For  $j \in \{1, \dots, |S| + i + 1\}$ ,

$$\pi'(j) = \begin{cases} v & \text{if } j = |S| + i + 1; \\ \pi(j) & \text{otherwise.} \end{cases}$$

Thus  $\pi'$  is defined on a different (slightly larger) ground set than  $\pi$  using  $v$  as the new tail vertex.

$$\pi' = \begin{cases} \pi'_{(i, S \cup \{v\}, v)} & \text{if } v \in \text{Inn}(V); \\ \pi'_{(i+1, S, v)} & \text{if } v = p_{i+1}. \end{cases}$$

When  $i = n - k$  and  $S = \text{Inn}(V)$  no effect.

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## Runtime of the $(\mu+1)$ EA

### Lemma

The population size  $\mu$  is bounded by  $O(2^k kn)$ .

**Proof.** For every ground set / tail vertex combination, there is exactly one individual (invariant).

- There are  $\binom{k}{|S|}$  ways to choose a distinct set  $S \subseteq \text{Inn}(V)$
- There are  $(n - k)$  ways to choose a distinct set of  $\gamma$ -respecting outer points
- There are  $|S| + 1$  ways of choosing the tail vertex  $r \in S \cup \{p_i\}$ .

$$(n - k) \sum_{s=0}^k \binom{k}{s} (s + 1) = O(2^k kn).$$

□

Parameterized Complexity Analysis of EAs

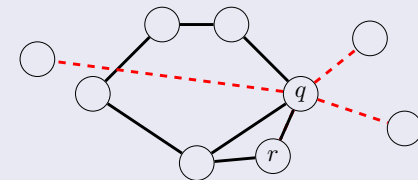
## $(\mu+1)$ EA

### Optimal substructure property

If there is an optimal permutation  $\pi_{(i, S, r)}$  in  $P$ , there exists some correct mutation that can construct a slightly larger subtour that is also optimal.

$F(i, S, r) :=$  length of optimal tour for  $(i, S, r)$  – can be defined recursively using the Bellman Principle

$$F(i, S, r) = \begin{cases} \min_{q \in S \cup \{p_{i-1}\}} F(i - 1, S, q) + d(q, r) & \text{if } r \in \text{Out}(V); \\ \min_{q \in (S \setminus \{r\}) \cup \{p_i\}} F(i, S \setminus \{r\}, q) + d(q, r) & \text{if } r \in \text{Inn}(V). \end{cases}$$



Parameterized Complexity Analysis of EAs

## $(\mu+1)$ EA

### Theorem

Let  $V$  be a set of  $n$  points in the Euclidean plane with  $|\text{Inn}(V)| = k$ . After  $O(2^k k^2 n^2)$  generations, the  $(\mu+1)$  EA has solved the TSP on  $V$  to optimality in expectation and with probability  $1 - e^{-\Omega(n)}$ .

**Proof.** Suppose  $\exists \pi = \pi_{(i,S,r)} \in P$  with  $f(\pi_{(i,S,r)}) = F(i, S, r)$ .

- with probability  $1/\mu$ ,  $\pi$  is selected for mutation
- with probability at least  $1/(k+1)$ ,  $\pi$  is extended optimally

Probability of extending an optimal path of length  $m$  is at least  $\Omega(1/(\mu(k+1)))$  (**Bernoulli trial**).

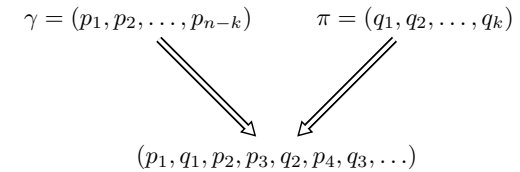
We can use induction on  $m$  since the permutation corresponding to  $(1, \{ \}, p_1)$  is already optimal.

Since optimal paths for a given  $(i, S, r)$  are never lost, the expected time until the optimal path of length  $n$  exists is  $O(n\mu(k+1)) = O(2^k k^2 n^2)$ .  $\square$

## $(1+1)$ EA

We already know that every optimal permutation must be  $\gamma$ -respecting. Suppose we only search the space of orderings on  $k$  inner points. Given a permutation  $\pi$  on  $\text{Inn}(V)$ , need to find where to insert the outer points into  $\pi$  so that the resulting permutation

1. respects  $\gamma$  and  $\pi$
2. corresponds to the shortest tour of all permutations that respect  $\gamma$  and  $\pi$



Exhaustive search  $O(n^k)$ , but we can be more clever...

## $(1+1)$ EA

### Direct dynamic programming

Maintain  $F[i, j, m]$ , a  $(n-k) \times (k+1) \times 2$  array where  $i \in \{1, \dots, n-k\}$ ,  $j \in \{0, 1, \dots, k\}$  and  $m \in \{\text{Inn}, \text{Out}\}$ .

$F[i, j, m]$  stores fitness of optimal permutation through points  $p_1, \dots, p_i$  and  $q_1, \dots, q_j$  ending on an outer point ( $m = \text{Out}$ ) or an inner point ( $m = \text{Inn}$ ).

### Fitness of $\pi$

$$\text{Dyn}(\pi) = \min\{F[n-k, k, \text{Out}] + d(p_{n-k}, p_1), F[n-k, k, \text{Inn}] + d(q_k, p_1)\}$$

Starting with  $F[1, 0, \text{Out}] = 0$ , use dynamic programming to fill out  $F$ .

$$F[i, j, \text{Inn}] = \min\{F[i, j-1, \text{Out}] + d[p_i, q_j], F[i, j-1, \text{Inn}] + d[q_{j-1}, q_j]\}$$

**Cost of computing**  $\text{Dyn}(\pi)$  is  $O(kn)$ .

## $(1+1)$ EA

### Search the space of permutations on $\text{Inn}(V)$

#### $(1+1)$ EA

- Choose uniformly at random a permutation  $x = (q_1, \dots, q_k)$  on the inner points
- While optimum not found
  - Construct  $x'$  from  $x$  by applying  $s+1$  random inversions where  $s$  is chosen according to  $\text{Pois}(1)$
  - If  $\text{Dyn}(x') \leq \text{Dyn}(x)$  then  $x \leftarrow x'$ .

Inversion mutation (pick a subsequence of the permutation and invert it)

$$(2, \mathbf{1}, \mathbf{6}, \mathbf{7}, \mathbf{4}, \mathbf{5}, 3) \Rightarrow (2, \mathbf{4}, \mathbf{7}, \mathbf{6}, \mathbf{1}, \mathbf{5}, 3)$$

This corresponds to the common 2-opt operation for the TSP.

## Runtime analysis of the (1+1) EA

### Theorem

The (1+1) EA solves the TSP with  $k$  inner points in  $O((k-1)!k^{2k-2})$  expected calls to the fitness function.

**Proof.** The probability that a mutation operation for a specific sequence of  $\ell$  basic operations is at least

$$\frac{1}{e(\ell-1)!} \cdot \frac{1}{k^{2\ell}}.$$

Expected waiting time for such a mutation operation is

$$\left( \frac{1}{e(\ell-1)!} \cdot \frac{1}{k^{2\ell}} \right)^{-1} = O(\ell!k^{2\ell}).$$

Need at most  $(k-1)$  inversions to transform arbitrary permutation on  $k$  points to another.  $\square$

Fitness function costs  $O(kn) \implies$  the (1+1) EA is FPT.

## Makespan Scheduling

- Given a set of  $n$  jobs to be scheduled on two machines
- Job  $j$  time  $p_j$  on either machine.
- A schedule is a decision vector  $x \in \{0, 1\}^n$
- The *load* of a machine is the sum of processing times assigned to it
- The *makespan* is the maximum load over both machines:

$$f : \{0, 1\}^n \rightarrow \mathbb{N} := x \mapsto \max \left\{ \sum_{j=1}^n x_j p_j, \sum_{j=1}^n (1 - x_j) p_j \right\}.$$

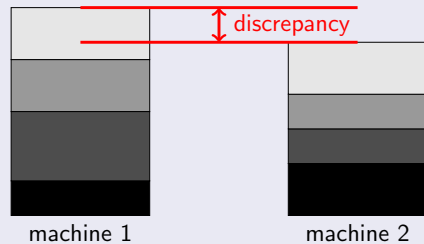
Objective is to find the schedule with the minimum makespan.

- $P = \sum_{j=1}^n p_j$ .
- $P/2 \leq f(x) \leq P$ .
- WLOG,  $p_1 \geq \dots \geq p_n$ .

## Makespan Scheduling – parameterization

### Definition

The *discrepancy*  $\Delta(x) = 2f(x) - P$  is the difference in load across machines.



Given an instance of makespan scheduling and an integer  $k$ , is  $p_k \geq \Delta^* \geq p_{k+1}$ ?

- $\Delta^* \geq 0$  discrepancy of optimal schedule
- $p_{n+1} = 0$

## Makespan Scheduling

Let  $\ell(n)$  denote the run length.

### $k$ -biased RLS

```

 $x \leftarrow$  an element of  $\{0, 1\}^n$  uniformly at random.
for  $i \leftarrow 1$  to  $\ell(n)$ 
   $y \leftarrow x$ 
  Choose  $0 \leq r \leq 1$  uniformly at random.
  if  $r < 1/n$ , then Choose  $j \in \{1, \dots, k\}$  u.a.r.
  else Choose  $j \in \{k+1, \dots, n\}$  u.a.r.
   $y_j \leftarrow 1 - y_j$ 
  if  $f(y) \leq f(x)$  then  $x \leftarrow y$ 
    
```

Same as traditional RLS, but prefers not to flip “large” jobs

### Lemma.

Let  $k$  be such that  $p_{k+1} \leq \Delta^*$ . Let  $x'$  be a decision vector such that the contribution of jobs  $1, \dots, k$  is minimal. Then starting from  $x'$ ,  $k$ -biased RLS with a run length of  $\ell(n) = \lceil 2n(\ln n + 1) \rceil$  solves the problem with probability bounded below by  $\Omega(n^{-2})$ .

### Proof sketch.

- Jobs 1 through  $k$  are already correct.
- Need to move any small jobs (index  $> k$ ) off of the fuller machine.
- Always possible since  $\Delta(x)$  is always larger than a small job, ELSE it is optimal.
- Coupon collector and Markov inequality:  $\lceil 2n(\ln n + 1) \rceil$  probability  $\Omega(1)$  as long as  $k$  large jobs aren't moved.
- Prob. large jobs aren't touched in  $\lceil 2n(\ln n + 1) \rceil$  steps:  $(1 - 1/n)^{\lceil 2n(\ln n + 1) \rceil} = \Omega(n^{-2})$ .

### Theorem.

A multi-start  $k$ -biased RLS procedure using a run length of  $\lceil 2n(\ln n + 1) \rceil$  solves the problem after  $O(2^k n^3 \log n)$  steps with probability at least  $1 - 1/e$ .

### Proof sketch.

- Probability that run starts with the first  $k$  jobs correctly placed is at least  $2^{-k}$ . Let  $q(n)$  be the probability that such a run is successful.
- Failure probability of  $t$  consecutive runs is at most

$$\left(1 - \frac{1}{2^k q(n)^{-1}}\right)^t$$

- Setting  $t = 2^k q(n)^{-1}$  makes the failure probability at most  $1/e$
- By the previous lemma  $q(n) = \Omega(n^{-2})$  so  $t = O(2^k n^2)$
- Run length is  $O(n \log n)$

Thank you

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