

Bioinspired computation in combinatorial optimization - Algorithms and their computational complexity

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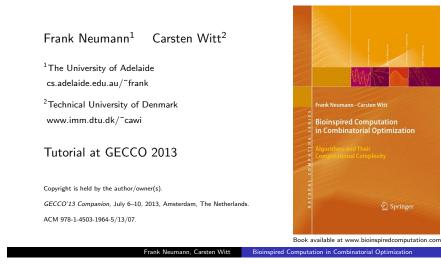
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Bioinspired Computation in Combinatorial Optimization – Algorithms and Their Computational Complexity



Why Do We Consider Randomized Search Heuristics?

- Not enough resources (time, money, knowledge) for a tailored algorithm
- Black Box Scenario –*

rules out problem-specific algorithms

- We like the simplicity, robustness, ... of Randomized Search Heuristics
- They are surprisingly successful.

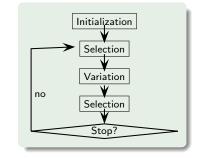
Point of view

Want a solid theory to understand how (and when) they work.

Evolutionary Algorithms and Other Search Heuristics

Most famous search heuristic: Evolutionary Algorithms (EAs)

- a bio-inspired heuristic
- paradigm: evolution in nature, "survival of the fittest"
- actually it's only an algorithm, a randomized search heuristic (RSH)



- Goal: optimization
- Here: discrete search spaces, combinatorial optimization, in particular pseudo-boolean functions

Optimize $f: \{0,1\}^n \to \mathbb{R}$

What RSHs Do We Consider?

Theoretically considered RSHs

- (1+1) EA
- $(1+\lambda)$ EA (offspring population)
- $(\mu+1)$ EA (parent population)
- $(\mu+1)$ GA (parent population and crossover)
- SEMO, DEMO, FEMO, ... (multi-objective)
- Randomized Local Search (RLS)
- Metropolis Algorithm/Simulated Annealing (MA/SA)
- Ant Colony Optimization (ACO)
- Particle Swarm Optimization (PSO)
- . . .

First of all: define the simple ones

f(*x*)

(1+1) EA and RLS for maximization problems

(1+1) EA

- Choose $x_0 \in \{0,1\}^n$ uniformly at random.
- **2** For $t := 0, \ldots, \infty$
 - Create y by flipping each bit of x_t indep. with probab. 1/n. • If $f(y) \ge f(x_t)$ set $x_{t+1} := y$ else $x_{t+1} := x_t$.

RLS

- Choose $x_0 \in \{0,1\}^n$ uniformly at random.
- **2** For $t := 0, ..., \infty$
 - Create y by flipping one bit of x_t uniformly.
 - If $f(y) \ge f(x_t)$ set $x_{t+1} := y$ else $x_{t+1} := x_t$.

How Do We Obtain Results?

We use (rarely in their pure form):

- Coupon Collector's Theorem
- Concentration inequalities: Markov, Chebyshev, Chernoff, Hoeffding, ... bounds
- Markov chain theory: waiting times, first hitting times
- Rapidly Mixing Markov Chains
- Random Walks: Gambler's Ruin, drift analysis, martingale theory, electrical networks
- Random graphs (esp. random trees)
- Identifying typical events and failure events
- Potential functions and amortized analysis
- . . .

Adapt tools from the analysis of randomized algorithms; understanding the stochastic process is often the hardest task.

What Kind of Theory Are We Interested in?

- Not studied here: convergence, local progress, models of EAs (e.g., infinite populations), ...
- Treat RSHs as randomized algorithm!
- Analyze their "runtime" (computational complexity) on selected problems

Definition

Let RSH *A* optimize *f*. Each *f*-evaluation is counted as a time step. The *runtime* $T_{A,f}$ of *A* is the random first point of time such that *A* has sampled an optimal search point.

- Often considered: expected runtime, distribution of $T_{A,f}$
- Asymptotical results w.r.t. n

Early Results

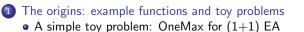
Analysis of RSHs already in the 1980s:

- Sasaki/Hajek (1988): SA and Maximum Matchings
- Sorkin (1991): SA vs. MA
- Jerrum (1992): SA and Cliques
- Jerrum/Sorkin (1993, 1998): SA/MA for Graph Bisection
- . . .

High-quality results, but limited to SA/MA (nothing about EAs) and hard to generalize.

Since the early 1990s

Systematic approach for the analysis of RSHs, building up a completely new research area



• A simple toy problem. One wax for (1+1)

2 Combinatorial optimization problems

- Minimum spanning trees
- Maximum matchings
- Shortest paths
- Makespan scheduling
- Covering problems
- Traveling salesman problem



4 References

Simple example functions (test functions)

- OneMax $(x_1, \ldots, x_n) = x_1 + \cdots + x_n$
- LeadingOnes $(x_1, \ldots, x_n) = \sum_{i=1}^n \prod_{j=1}^i x_j$
- BinVal $(x_1, ..., x_n) = \sum_{i=1}^n 2^{n-i} x_i$
- polynomials of fixed degree
- Goal: derive first runtime bounds and methods

Artificially designed functions

- with sometimes really horrible definitions
- but for the first time these allow rigorous statements
- Goal: prove benefits and harm of RSH components,
 - e.g., crossover, mutation strength, population size ...

Agenda

The origins: example functions and toy problems A simple toy problem: OneMax for (1+1) EA

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Example: OneMax

Theorem (e.g., Droste/Jansen/Wegener, 1998)

The expected runtime of the RLS, (1+1) EA, (μ +1) EA, (1+ λ) EA on ONEMAX is $\Omega(n \log n)$.

Proof by modifications of Coupon Collector's Theorem.

Theorem (e.g., Mühlenbein, 1992)

The expected runtime of RLS and the (1+1) EA on ONEMAX is $O(n \log n)$.

Holds also for population-based $(\mu+1)$ EA and for $(1+\lambda)$ EA with small populations.

- Fitness levels: $L_i := \{x \in \{0,1\}^n \mid \text{ONEMAX}(x) = i\}$
- (1+1) EA never decreases its current fitness level.
- From i to some higher-level set with prob. at least

$$\underbrace{\binom{n-i}{1}}_{1} \cdot \underbrace{\binom{1}{n}}_{r} \cdot \underbrace{\binom{1}{n}}_{r} \cdot \underbrace{\binom{1-\frac{1}{n}}_{r}}_{r} \geq \frac{n-i}{en}$$

choose a 0-bit flip this bit keep the other bits

- Expected time to reach a higher-level set is at most $\frac{en}{n-i}$.
- Expected runtime is at most

$$\sum_{i=0}^{n-1} \frac{en}{n-i} = O(n \log n).$$

Later Results Using Toy Problems

- Find the theoretically optimal mutation strength (1/n for OneMax!).
- Bound the optimization time for linear functions (O(n log n)).
- optimal population size (often 1!)
- \bullet crossover vs. no crossover \rightarrow Real Royal Road Functions
- multistarts vs. populations
- frequent restarts vs. long runs
- dynamic schedules
- . . .

RSHs for Combinatorial Optimization

- Analysis of runtime and approximation quality on well-known combinatorial optimization problems, e.g.,
 - sorting problems (is this an optimization problem?),
 - covering problems,
 - cutting problems,
 - subsequence problems,
 - traveling salesman problem,
 - Eulerian cycles,
 - minimum spanning trees,
 - maximum matchings,
 - scheduling problems,
 - shortest paths,
 - ...
- We do not hope: to be better than the best problem-specific algorithms
- Instead: maybe reasonable polynomial running times
- In the following no fine-tuning of the results

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Minimum Spanning Trees:

- Given: Undirected connected graph G = (V, E) with n vertices and m edges with positive integer weights.
- Find: Edge set E' ⊆ E with minimal weight connecting all vertices.
- Search space {0,1}^m
- Edge e_i is chosen iff x_i=1
- Consider (1+1) EA

Fitness function:

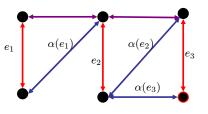
- Decrease number of connected components, find minimum spanning tree.
- f (s) := (c(s),w(s)).
 Minimization of f with respect to the lexicographic order.

First goal: Obtain a connected subgraph of G.

How long does it take?

Connected graph in expected time O(mlog n) (fitness-based partitions)

Bijection for minimum spanning trees:



$$\begin{split} &k := |\textbf{E}(\textbf{T}^*) \setminus \textbf{E}(\textbf{T}) | \\ & \text{Bijection } \alpha : \textbf{E}(\textbf{T}^*) \setminus \textbf{E}(\textbf{T}) \rightarrow \textbf{E}(\textbf{T}) \setminus \textbf{E}(\textbf{T}^*) \\ & \alpha(e_i) \text{ on the cycle of } \textbf{E}(\textbf{T}) \cup \{e_i\} \\ & w(e_i) \leq w(\alpha(e_i)) \\ & \Rightarrow k \text{ accepted } 2\text{-bit flips that turn } \textbf{T} \text{ into } \textbf{T}^* \end{split}$$

Upper Bound

Theorem:

The expected time until (1+1) EA constructs a minimum spanning tree is bounded by O(m²(log n + log w_{max})).

Sketch of proof:

- w(s) weight current solution s.
- w_{opt} weight minimum spanning tree T*
- set of m + 1 operations to reach T*
- m' = m (n 1) 1-bit flips concerning non-T* edges
 ⇒ spanning tree T
- k 2-bit flips defined by bijection
- n k non accepted 2-bit flips
- \Rightarrow average distance decrease (w(s) w_{opt})/(m + 1)

Proof

1-step (larger total weight decrease of 1-bit flips)2-step (larger total weight decrease of 2-bit flips)

Consider 2-steps:

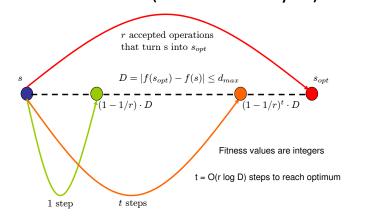
- Expected weight decrease by a factor 1 (1/(2n))
- Probability (n/m²) for a good 2-bit flip
- Expected time until q 2-steps O(qm²/n)

Consider 1-steps:

- Expected weight decrease by a factor 1 (1/(2m'))
- Probability (m'/m) for a good 1-bit flip
- Expected time until q 1-steps O(qm/m')

1-steps faster \Rightarrow show bound for 2-steps.

Expected Multiplicative Distance Decrease (aka Drift Analysis)

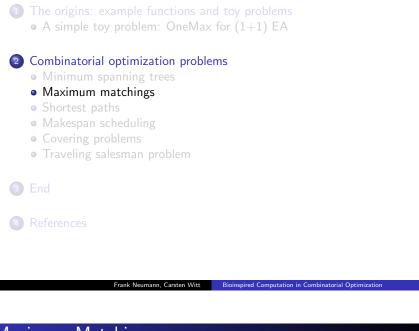


Maximum distance: $w(s) - w_{opt} \le D := m \cdot wmax$

1 step: Expected distance at most (1 - 1/(2n))(w(s) w_{opt})

t steps: Expected distance at most $(1 - 1/(2n))^t(w(s) - w_{opt})$ t := $[2 \cdot (ln 2)n(log D + 1)]: (1 - 1/(2n))^t(w(s) - w_{opt}) \le 1/2$ Expected number of 2-steps 2t = O(n(log n + log w_{max}))(Markov)

Expected optimization time $O(tm^2/n) = O(m^2(\log n + \log w_{max})).$



Maximum Matchings

A matching in an undirected graph is a subset of pairwise disjoint edges; aim: find a maximum matching (solvable in poly-time)

Simple example: path of odd length



Maximum matching with more than half of edges

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Maximum Matchings

A matching in an undirected graph is a subset of pairwise disjoint edges; aim: find a maximum matching (solvable in poly-time)

Simple example: path of odd length



Suboptimal matching

Concept: augmenting path

- Alternating between edges being inside and outside the matching
- Starting and ending at "free" nodes not incident on matching
- Flipping all choices along the path improves matching

Example: whole graph is augmenting path

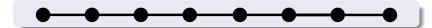
Interesting: how simple EAs find augmenting paths

Maximum Matchings: Upper Bound

Fitness function $f: \{0,1\}^{\# \text{ edges}} \to \mathbb{R}$:

- one bit for each edge, value 1 iff edge chosen
- value for legal matchings: size of matching
- otherwise penalty leading to empty matching

Example: path with n + 1 nodes, n edges: bit string selects edges



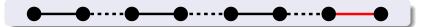
Theorem

The expected time until (1+1) EA finds a maximum matching on a path of n edges is $O(n^4)$.

Maximum Matchings: Upper Bound (Ctnd.)

Proof idea for $O(n^4)$ bound

- Consider the level of second-best matchings.
- Fitness value does not change (walk on plateau).
- If "free" edge: chance to flip one bit! \rightarrow probability $\Theta(1/n)$.
- Else steps flipping two bits \rightarrow probability $\Theta(1/n^2)$.
- Shorten or lengthen augmenting path
- At length 1, chance to flip the free edge!

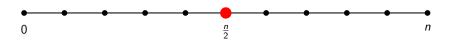


• Length changes according to a fair random walk \rightarrow equal probability for lengthenings and shortenings

Fair Random Walk

Scenario: fair random walk

- Initially, player A and B both have $\frac{n}{2}$ USD
- Repeat: flip a coin
- If heads: A pays 1 USD to B, tails: other way round
- Until one of the players is ruined.



How long does the game take in expectation?

Theorem:

Fair random walk on $\{0, \ldots, n\}$ takes in expectation $O(n^2)$ steps.

Maximum Matchings: Upper Bound (Ctnd.)

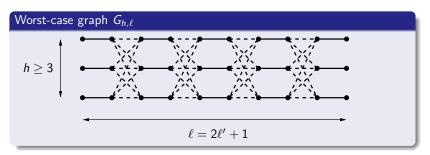
Proof idea for $O(n^4)$ bound

- Consider the level of second-best matchings.
- Fitness value does not change (walk on *plateau*).
- If "free" edge: chance to flip one bit! ightarrow probability $\Theta(1/n)$.
- Else steps flipping two bits \rightarrow probability $\Theta(1/n^2)$.
- Shorten or lengthen augmenting path
- At length 1, chance to flip the free edge!



Length changes according to a fair random walk, expected $O(n^2)$ two-bit flips suffice, expected optimization time $O(n^2) \cdot O(n^2) = O(n^4)$.

Maximum Matchings: Lower Bound



Augmenting path can get shorter but is more likely to get longer. (unfair random walk)

Theorem

For $h \ge 3$, (1+1) EA has exponential expected optimization time $2^{\Omega(\ell)}$ on $G_{h,\ell}$.

Proof requires analysis of negative drift (simplified drift theorem).

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Maximum Matching: Approximations

Insight: do not hope for exact solutions but for approximations

For maximization problems: solution with value *a* is called $(1 + \varepsilon)$ -approximation if $\frac{\mathsf{OPT}}{a} \leq 1 + \varepsilon$, where OPT optimal value.

Theorem

For $\varepsilon > 0$, (1+1) EA finds a (1 + ε)-approximation of a maximum matching in expected time $O(m^{2/\varepsilon+2})$ (m number of edges).

Proof idea: If current solution worse than $(1 + \varepsilon)$ -approximate, there is a "short" augmenting path (length $\leq 2/\varepsilon + 1$); flip it in one go.

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All-pairs-shortest-path (APSP) problem

Given: Connected directed graph G = (V, E), |V| = n and |E| = m, and a function $w : E \to N$ which assigns positive integer weights to the edges.

Compute from each vertex $v_i \in V$ a shortest path (path of minimal weight) to every other vertex $v_j \in V \setminus \{v_i\}$

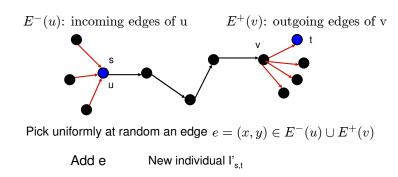
Representation:

Individuals are paths between two particular vertices v_i and v_i

Initial Population: $P := \{I_{u,v} = (u,v) | (u,v) \in E\}$

Mutation-based EA

Mutation:



Pick individual $I_{u,v}$ uniformly at random

Steady State EA

- 1. Set $P = \{I_{u,v} = (u,v) \mid (u,v) \in E\}.$
- 2. Choose an individual $I_{x,y} \in P$ uniformly at random.
- 3. Mutate $I_{x,y}$ to obtain an individual $I'_{s,t}$.
- 4. If there is no individual $I_{s,t} \in P$, $P = P \cup \{I'_{s,t}\}$, else if $f(I'_{s,t}) \leq f(I_{s,t})$, $P = (P \cup \{I'_{s,t}\}) \setminus \{I_{s,t}\}$
- 5. Repeat Steps 2–4 forever.

Lemma:

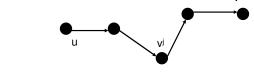
Let $\ell \ge \log n$. The expected time until has found all shortest paths with at most ℓ edges is $O(n^3 \ell)$.

Proof idea:

Consider two vertices u and $v, u \neq v$.

Let $\gamma := (v^1 = u, v^2, \dots, v^{\ell'+1} = v)$ be a shortest path from u to v consisting of $\ell', \ell' \leq \ell$, edges in G

the sub-path $\gamma' = (v^1 = u, v^2, \dots, v^j)$ is a shortest path from u to v^j .



Population size is upper bounded n² (for each pair of vertices at most one path)

- Pick shortest path from u to v_j and append edge (v_i , v_{i+1})
- Shortest path from u to v_{i+1}
- Probability to pick $I_{u,vj}$ is at least $1/n^2$
- Probability to append right edge is at least 1/(2n)
- Success with probability at least p = 1/(2n³)
- At most I successes needed to obtain shortest path from u to v

Consider typical run consisting of T=cn³l steps.

What is the probability that the shortest path from u to v has been obtained?

We need at most I successes, where a success happens in each step with probability at least $p = 1/(2n^3)$

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Define for each step i a random variable X_i .

 $X_i = 1$ if step *i* is a success

 $X_i = 0$ if step *i* is not a success

Analysis

 $\begin{array}{ll} Prob(X_i=1) \geq p = 1/(2n^3) \quad X = \sum_{i=1}^T X_i \qquad X \geq \ell \; ???\\ \text{Expected number of successes } E(X) \geq T/(2n^3) = \frac{cn^3\ell}{2n^3} = \frac{c\ell}{2}\\ \text{Chernoff:} \quad Prob(X < (1-\delta)E(x)) \leq e^{-E(X)\delta^2/2}\\ \delta = \frac{1}{2}\\ Prob(X < (1-\frac{1}{2})E(x)) \leq e^{-E(X)/8} \leq e^{-T/(16n^3)} = e^{-cn^3\ell/(16n^3)} = e^{-c\ell/(16)}\\ \text{Probability for failure of at least one pair of vertices at most:} \;\; n^2 \cdot e^{-c\ell/16}\\ c \; \text{large enough and} \; \ell \geq \log n \text{:}\\ \text{No failure in any path with probability at least } \alpha = 1 - n^2 \cdot e^{-c\ell/16} = 1 - o(1)\\ \text{Holds for any phase of T steps}\\ & \text{Expected time upper bound by } \; T/\alpha = O(n^3\ell) \end{array}$

Shortest paths have length at most n-1.

Set l = n-1

Theorem

The expected optimization time of Steady State EA for the APSP problem is $O(n^4)$.

Remark:

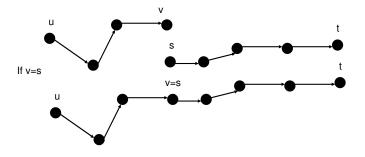
There are instances where the expected optimization of $(\mu+1)\text{-}\mathrm{EA}$ is $\Omega(n^4)$

Question:

Can crossover help to achieve a better expected optimization time?

Crossover

Pick two individuals $I_{u,v}$ and $I_{s,t}$ from population uniformly at random.



Steady State GA

- 1. Set $P = \{I_{u,v} = (u,v) \mid (u,v) \in E\}.$
- 2. Choose $r \in [0, 1]$ uniformly at random.
- 3. If $r \leq p_c$, choose two individuals $I_{x,y} \in P$ and $I_{x',y'} \in P$ uniformly at random and perform crossover to obtain an individual $I'_{s,t}$, else choose an individual $I_{x,y} \in P$ uniformly at random and mutate $I_{x,y}$ to obtain an individual $I'_{s,t}$.
- 4. If $I'_{s,t}$ is a path from s to t then \star If there is no individual $I_{s,t} \in P, P = P \cup \{I'_{s,t}\},$ \star else if $f(I'_{s,t}) \leq f(I_{s,t}), P = (P \cup \{I'_{s,t}\}) \setminus \{I_{s,t}\}.$
- 5. Repeat Steps 2–4 forever.

 p_c is a constant

Theorem:

The expected optimization time of Steady State GA is $O(n^{3.5}\sqrt{\log n})$.

Mutation and $\ell^* := \sqrt{n \log n}$

All shortest path of length at most I* edges are obtained

Show: Longer paths are obtained by crossover within the stated time bound.

Analysis Crossover

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Long paths by crossover:

Assumption: All shortest paths with at most I* edges have already been obtained.

Assume that all shortest paths of length $k \le l^*$ have been obtained.

What is the expected time to obtain all shortest paths of length at most 3k/2?

Analysis Crossover

Consider pair of vertices x and y for which a shortest path of r, $k < r \le 3k/2$, edges exists.

There are 2k-r pairs of shortest paths of length at most k that can be joined to obtain shortest path from x to y.

Probability for one specific pair: at least 1/n⁴

At least 2k+1-r possible pairs: probability at least $(2k+1-r)/n^4 \ge k/(2n^4)$

At most n² shortest paths of length r, $k < r \le 3k/2$

Time to collect all paths O(n⁴ log n/ k) (similar to Coupon Collectors Theorem)

Analysis Crossover

Sum up over the different values of k, namely

 $\sqrt{n\log n}, c \cdot \sqrt{n\log n}, c^2 \cdot \sqrt{n\log n}, \dots, c^{\log_c(n/\sqrt{n\log n})} \cdot \sqrt{n\log n},$

where c = 3/2.

Expected Optimization

$$\sum_{s=0}^{\log_c(n/\sqrt{n\log n})} \left(O\left(\frac{n^4\log n}{\sqrt{n\log n}}\right)c^{-s}\right) = O(n^{3.5}\sqrt{\log n})\sum_{s=0}^{\infty}c^{-s} = O(n^{3.5}\sqrt{\log n})$$

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Makespan Scheduling

What about NP-hard problems? \rightarrow Study approximation quality

Makespan scheduling on 2 machines:

- *n* objects with weights/processing times w_1, \ldots, w_n
- 2 machines (bins)
- Minimize the total weight of fuller bin = makespan.

Formally, find $I \subseteq \{1, \ldots, n\}$ minimizing

$$\max\left\{\sum_{i\in I}w_i,\sum_{i\notin I}w_i\right\}.$$



Sometimes also called the Partition problem.

Frank Neumann, Carsten Witt

This is an "easy" NP-hard problem, good approximations possible

Fitness Function

- Problem encoding: bit string x_1, \ldots, x_n reserves a bit for each object, put object *i* in bin $x_i + 1$.
- Fitness function

$$f(x_1,...,x_n) := \max\left\{\sum_{i=1}^n w_i x_i, \sum_{i=1}^n w_i (1-x_i)\right\}$$

to be minimized.

Consider (1+1) EA and RLS.

- Worst-case results
- Success probabilities and approximations
- An average-case analysis
- A parameterized analysis

Sufficient Conditions for Progress

Abbreviate $S := w_1 + \cdots + w_n \Rightarrow$ perfect partition has cost $\frac{S}{2}$.

Suppose we know

- $s^* = \text{size of smallest object in the fuller bin}$,
- $f(x) > \frac{S}{2} + \frac{s^*}{2}$ for the current search point x

then the solution is improvable by a single-bit flip.



If $f(x) < \frac{s}{2} + \frac{s^*}{2}$, no improvements can be guaranteed.

Lemma

If smallest object in fuller bin is always bounded by s^* then (1+1) EA and RLS reach f-value $\leq \frac{5}{2} + \frac{s^*}{2}$ in expected $O(n^2)$ steps.

Sufficient Conditions for Progress

Abbreviate $S := w_1 + \cdots + w_n \Rightarrow$ perfect partition has cost $\frac{S}{2}$.

Suppose we know

- $s^* =$ size of smallest object in the fuller bin,
- $f(x) > \frac{S}{2} + \frac{s^*}{2}$ for the current search point x

then the solution is improvable by a single-bit flip.



If $f(x) < \frac{5}{2} + \frac{s^*}{2}$, no improvements can be guaranteed.

Lemma

If smallest object in fuller bin is always bounded by s^* then (1+1) EA and RLS reach f-value $\leq \frac{5}{2} + \frac{s^*}{2}$ in expected $O(n^2)$ steps.

Sufficient Conditions for Progress

Abbreviate $S := w_1 + \cdots + w_n \Rightarrow$ perfect partition has cost $\frac{S}{2}$.

Suppose we know

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- $f(x) > \frac{5}{2} + \frac{s^*}{2}$ for the current search point x

then the solution is improvable by a single-bit flip.



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Bioinspired Computation in Combina

Worst-Case Results

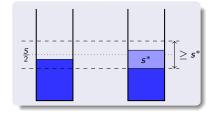
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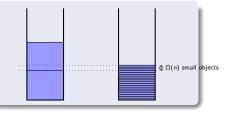
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Bioinspired Computation in Combinatorial Opt

Worst-Case Instance

Instance $W_{\varepsilon}^* = \{w_1, \ldots, w_n\}$ is defined by $w_1 := w_2 := \frac{1}{3} - \frac{\varepsilon}{4}$ (big objects) and $w_i := \frac{1/3 + \varepsilon/2}{n-2}$ for $3 \le i \le n$, ε very small constant; n even Sum is 1; there is a perfect partition.

But if one bin with big and one bin with small objects: value $\frac{2}{3} - \frac{\varepsilon}{2}$. Move a big object in the emptier bin \Rightarrow value $(\frac{1}{3} + \frac{\varepsilon}{2}) + (\frac{1}{3} - \frac{\varepsilon}{4}) = \frac{2}{3} + \frac{\varepsilon}{4}!$ Need to move $> \varepsilon n$ small objects at once for improvement: very unlikely.



With constant probability in this situation, $n^{\Omega(n)}$ needed to escape.

On any instance to the makespan scheduling problem, the (1+1) EA and RLS reach a solution with approximation ratio $\frac{4}{3}$ in expected time $O(n^2)$.

Use study of object sizes and previous lemma.

Theorem

There is an instance W_{ε}^* such that the (1+1) EA and RLS need with prob. $\Omega(1)$ at least $n^{\Omega(\check{n})}$ steps to find a solution with a better ratio than $4/3 - \varepsilon$.

Worst Case – PRAS by Parallelism

Previous result shows: success dependent on big objects

Theorem

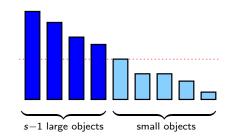
On any instance, the (1+1) EA and RLS with prob. $\geq 2^{-c\lceil 1/\varepsilon\rceil} \ln(1/\varepsilon)$ find a $(1 + \varepsilon)$ -approximation within $O(n \ln(1/\varepsilon))$ steps.

- 2^{O([1/ε] ln(1/ε))} parallel runs find a (1 + ε)-approximation with prob. ≥ 3/4 in O(n ln(1/ε)) parallel steps.
- Parallel runs form a polynomial-time randomized approximation scheme (PRAS)!

Worst Case – PRAS by Parallelism (Proof Idea)

Set $s := \left\lceil \frac{2}{\varepsilon} \right\rceil$

Assuming $w_1 \geq \cdots \geq w_n$, we have $w_i \leq \varepsilon \frac{S}{2}$ for $i \geq s$.



analyze probability of distributing

- large objects in an optimal way,
- small objects greedily \Rightarrow error $\leq \varepsilon \frac{S}{2}$,

Random search rediscovers algorithmic idea of early algorithms.

Average-Case Analyses

Makespan Scheduling – Known Averge-Case Results

Models: each weight drawn independently at random, namely

- ① uniformly from the interval [0, 1],
- exponentially distributed with parameter 1 (i. e., Prob(X ≥ t) = e^{-t} for t ≥ 0).

Approximation ratio no longer meaningful, we investigate: discrepancy = absolute difference between weights of bins.

How close to discrepancy 0 do we come?

Deterministic, problem-specific heuristic LPT

Sort weights decreasingly, put every object into currently emptier bin.

Known for both random models: LPT creates a solution with discrepancy $O((\log n)/n)$.

What discrepancy do the (1+1) EA and RLS reach in poly-time?

Average-Case Analysis of the (1+1) EA

Theorem

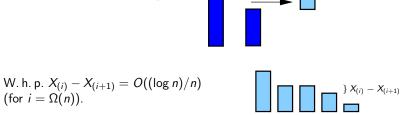
In both models, the (1+1) EA reaches discrepancy $O((\log n)/n)$ after $O(n^{c+4}\log^2 n)$ steps with probability $1 - O(1/n^c)$.

Almost the same result as for LPT!

Proof exploits order statistics:

If $X_{(i)}$ (*i*-th largest) in fuller bin, $X_{(i+1)}$ in emptier one, and discrepancy $> 2(X_{(i)} - X_{(i+1)}) > 0$, then objects can be swapped; discrepancy falls

Consider such "difference objects".



A Parameterized Analysis

Have seen: problem is hard for (1+1) EA/RLS in the worst case, but not so hard on average.

What parameters make the problem hard?

Definition

A problem is *fixed-parameter tractable (FPT)* if there is a problem parameter k such that it can be solved in time $f(k) \cdot poly(n)$, where f(k) does not depend on n.

Intuition: for small k, we have an efficient algorithm.

Considered parameters (Sutton and Neumann, 2012):

- Value of optimal solution
- In No. jobs on fuller machine in optimal solution
- Unbalance of optimal solution

Value of Optimal Solution

Recall approximation result: decent chance to distribute k big jobs optimally if k small.

Since $w_1 \geq \cdots \geq w_n$, already $w_k \leq S/k$.

Consequence: optimal distribution of first k objects \rightarrow can reach makespan S/2 + S/k by greedily treating the other objects.

Theorem

(1+1) EA and RLS find solution of makespan $\leq S/2 + S/k$ with probability $\Omega((2k)^{-ek})$ in time $O(n \log k)$. Multistarts have success probability $\geq 1/2$ after $O(2^{(e+1)k}k^{ek}n \log k)$ evaluations.

 $2^{(e+1)k}k^{ek}\log k$ does not depend on $n \rightarrow$ a randomized FPT-algorithm.

No. Objects on Fuller Machine

Suppose: optimal solution puts only k objects on fuller machine. Notion: k is called *critical path size*.

Intuition:

- Good chance of putting k objects on same machine if k small,
- other objects can be moved greedily.

Theorem

For critical path size k, multistart RLS finds optimum in $O(2^k(en)^{ck} n \log n)$ evaluations with probability $\geq 1/2$.

Due to term n^{ck} , result is somewhat weaker than FPT (a so-called XP-algorithm). Still, for constant k polynomial.

Remark: with (1+1)-EA, get an additional log w_1 -term.

Unbalance of Optimal Solution

Consider discrepancy of optimum $\Delta^* := 2(OPT - S/2)$.

Question/decision problem: Is $w_k \ge \Delta^* \ge w_{k+1}$?

Observation: If $\Delta^* \ge w_{k+1}$, optimal solution will put w_{k+1}, \ldots, w_n on emptier machine. Crucial to distribute first k objects optimally.

Theorem

Multistart RLS with biased mutation (touches objects w_1, \ldots, w_k with prob. 1/(kn) each) solves decision problem in $O(2^k n^3 \log n)$ evaluations with probability $\geq 1/2$.

Again, a randomized FPT-algorithm.

Agenda

The origins: example functions and toy problems
 A simple toy problem: OneMax for (1+1) EA

2 Combinatorial optimization problems

- Minimum spanning trees
- Maximum matchings
- Shortest paths
- Makespan scheduling
- Covering problems
- Traveling salesman problem

3 End

4 References

The Problem

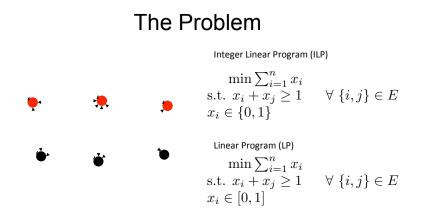
The Vertex Cover Problem: Given an undirected graph G=(V,E).



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

NP-hard, several 2-approximation algorithms.

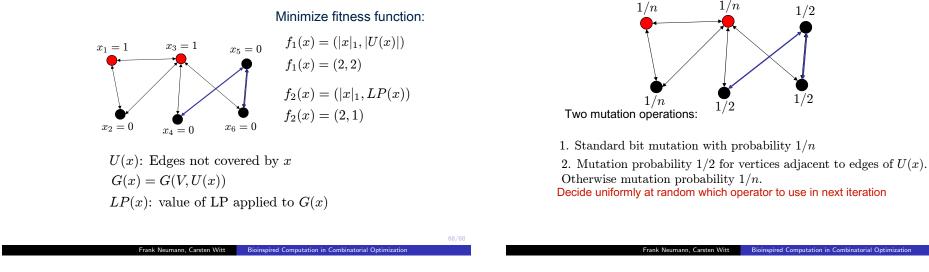
Simple single-objective evolutionary algorithms fail!!!



Decision problem: Is there a set of vertices of size at most k covering all edges? Our parameter: Value of an optimal solution (OPT)

Evolutionary Algorithm

Representation: Bitstrings of length n

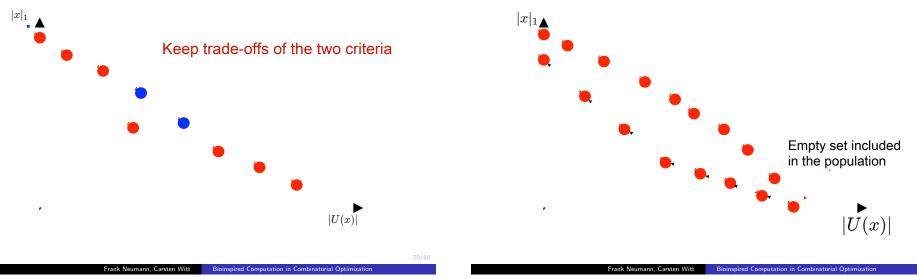


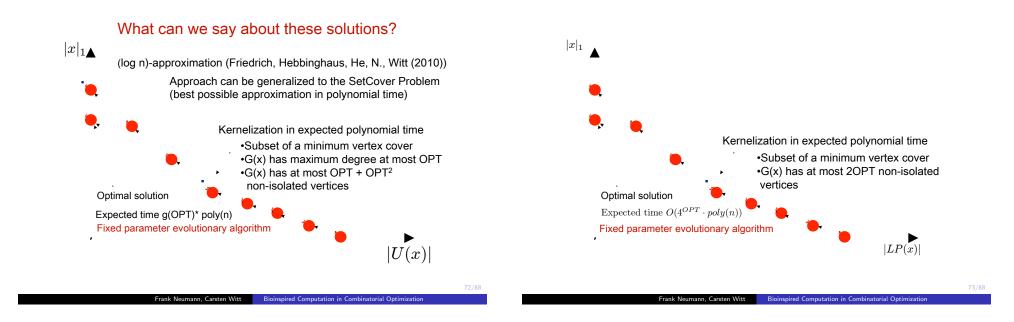
Evolutionary Algorithm

1/n

Multi-Objective Approach:

Treat the different objectives in the same way





Linear Programming

Combination with Linear Programming

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

$x_i \in \{0, 1/2, 1\}, 1 \le i \le n$

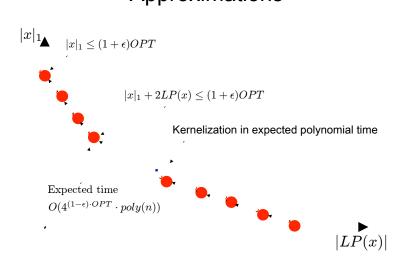
Theorem (Nemhauser, Trotter (1975)):

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?





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References

Euclidean TSP

Given n points in the plane and Euclidean distances between the cities.

Find a shortest tour that visits each city exactly once and return to the origin.

NP-hard, PTAS, FPT when number of inner points is the parameter.

Representation and Mutation

Representation: Permutation of the n cities

For example: (3, 4, 1, 2, 5)

Inversion (inv) as mutation operator:

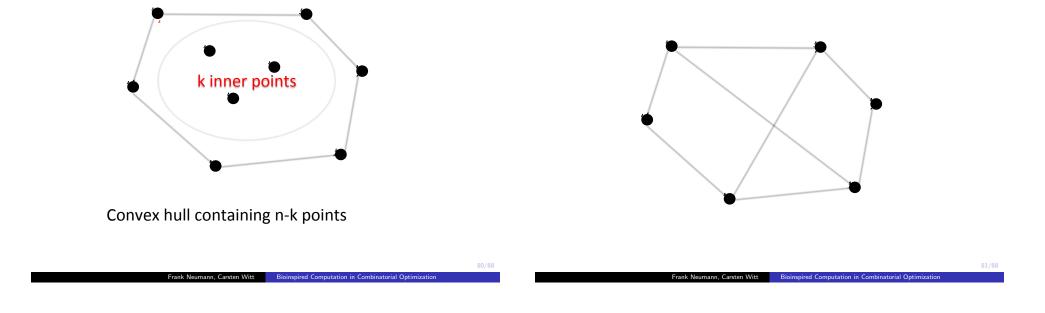
- Select i,j from {1, ...n} uniformly at random and invert the part from position i to position j.
- Inv(2,5) applied to (3, 4, 1, 2, 5) yields (3, 5, 2, 1, 4)

(1+1) EA

 $\begin{array}{l} x \leftarrow \text{a random permutation of } [n].\\ \textbf{repeat} \quad \text{forever}\\ y \leftarrow \text{MUTATE}(x)\\ \textbf{if } f(y) < f(x) \textbf{ then } x \leftarrow y \end{array}$

Mutation:

(1+1) EA: k random inversion, k chosen according to 1+Pois(1)

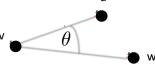


Angle bounded set of points

There may be an exponential number of inversion to end up in a local optimum if points are in arbitrary positions (Englert et al, 2007).

We assume that the set V is angle bounded

V is angle-bounded by $\epsilon > 0$ if for any three points $u, v, w \in V, 0 < \epsilon < \theta < \pi - \epsilon$ where θ denotes the angle formed by the line from u to v and the line from v to w.



If V is angle-bounded then we get a lower bound on an improvement depending on $\boldsymbol{\epsilon}$

Progress

Intersection and Mutation

Assumptions:

 d_{max} : Maximum distance between any two points d_{min} : Minimum distance between any two points V is angle-bounded by ϵ

Whenever the current tour is not intersectionfree, we can guarantee a certain progress

Lemma:

Let x be a permutation such that is not intersection-free. Let y be the permutation constructed from an inversion on x that replaces two intersecting edges with two non-intersecting edges. Then, $f(x) - f(y) > 2d_{min} \left(\frac{1-\cos(\epsilon)}{\cos(\epsilon)}\right)$.

Tours

A tour x is either

- Intersection free
- Non intersection free

Intersection free tour are good. The points on the convex hull are already in the right order (Quintas and Supnick, 1965).

Claim: We do not spend too much time on non intersection free tours.

Time spend on intersecting tours

Lemma:

Let $(x^{(1)}, x^{(2)}, \ldots, x^{(t)}, \ldots)$ denote the sequence of permutations generated by the (1+1)-EA. Let α be an indicator variable defined on permutations of [n] as

 $\alpha(x) = \begin{cases} 1 & x \text{ contains intersections;} \\ 0 & \text{otherwise.} \end{cases}$ Then $E\left(\sum_{t=1}^{\infty} \alpha(x^{(t)})\right) = O\left(n^3 \left(\frac{d_{max}}{d_{min}} - 1\right) \left(\frac{\cos(\epsilon)}{1 - \cos(\epsilon)}\right)\right).$

For an m x m grid:

For points on an $m \times m$ grid this bound becomes $O(n^3m^5)$.

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Parameterized Result

Lemma:

Suppose V has k inner points and x is an intersection-free tour on V. Then there is a sequence of at most 2k inversions that transforms x into an optimal permutation.

Theorem:

Let V be a set of points quantized on an $m \times m$ and k be the number of inner points. Then the expected optimisation time of the (1+1)-EA on V is $O(n^3m^5) + O(n^{4k}(2k-1)!)$.

Summary and Conclusions

- Runtime analysis of RSHs in combinatorial optimization
- Starting from toy problems to real problems
- Insight into working principles using runtime analysis
- General-purpose algorithms successful for wide range of problems
- Interesting, general techniques
- Runtime analysis of new approaches possible
- $\rightarrow\,$ An exciting research direction.

Thank you!

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References

F. Neumann and C. Witt (2010): Bioinspired Computation in Combinatorial Optimization - Algorithms and Their Computational Complexity. Springer A. Auger and B. Doerr (2011): Theory of Randomized Search Heuristics - Foundations and Recent Developments. World Scientific Publishing F. Neumann and I. Wegener (2007): Randomized local search, evolutionary algorithms, and the minimum spanning tree problem. Theoretical Computer Science 378(1):32-40. O. Giel and I. Wegener (2003): Evolutionary algorithms and the maximum matching problem. Proc. of STACS '03, LNCS 2607, 415–426, Springer B. Doerr, E. Happ and C. Klein (2012): Crossover can provably be useful in evolutionary computation. Theoretical Computer Science 425:17-33. C. Witt (2005): Worst-case and average-case approximations by simple randomized search heuristics. Proc. of STACS 2005, LNCS 3404, 44-56, Springer T. Friedrich, J. He, N. Hebbinghaus, F. Neumann and C. Witt (2010): Approximating covering problems by randomized search heuristics using multi-objective models. Evolutionary Computation 18(4):617-633. S. Kratsch and F. Neumann (2009): Fixed-parameter evolutionary algorithms and the vertex cover problem. In Proc. of GECCO 2009, 293-300. ACM. A. M. Sutton and F. Neumann (2012): A parameterized runtime analysis of evolutionary algorithms for the Euclidean traveling salesperson problem. Proc. of AAAI 2012.

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