Computação Bioinspirada - 5955010-2

1. Computação Evolutiva

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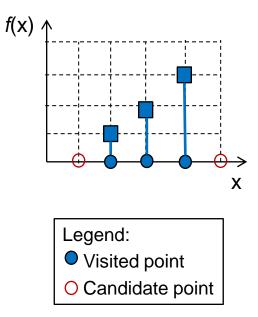
Programa de Pós-Graduação Em Computação Aplicada

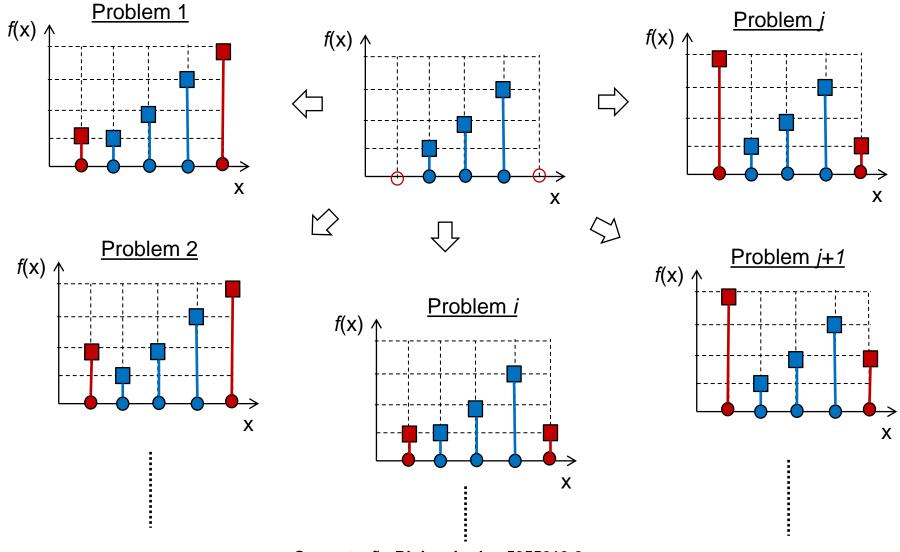
1.6. Aspectos Teóricos*

- 1.6.1. Is interesting to apply Evolutionary Algorithm (EAs) to my problem?
- 1.6.2. Introduction to the theory of EAs
- 1.6.3. Theory of Genetic Algorithms (GAs): Some Approaches
- 1.6.4. Examples
- 1.6.5. Conclusions

* Apresentado anteriormente como um tutorial na Latin American and Brazilian Schools on Computational Intelligence (LASCI & SBIC), Curitiba, October 13th, 2015

- Black box optimization
 - What should be the next point?





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- No Free Lunch Theorem
 - Considering all possible problems:

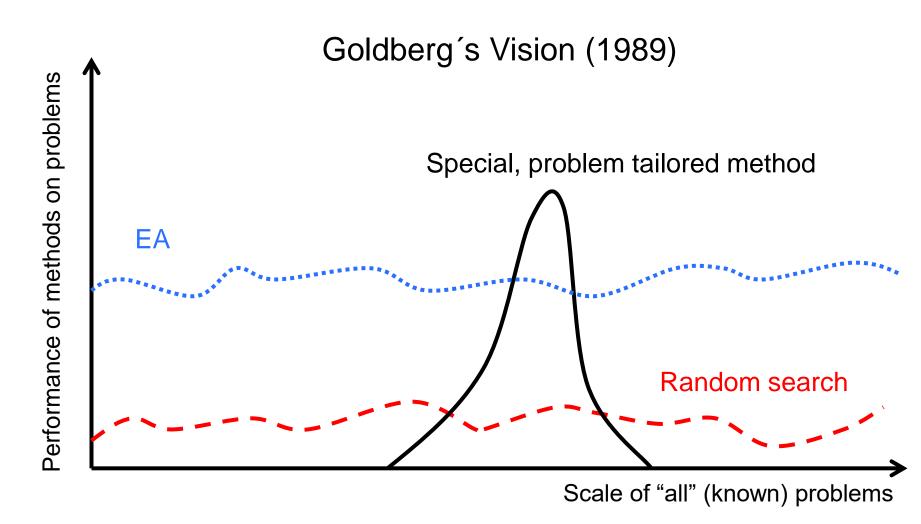
All black box algorithms that do not revisit points will show the same average performance!

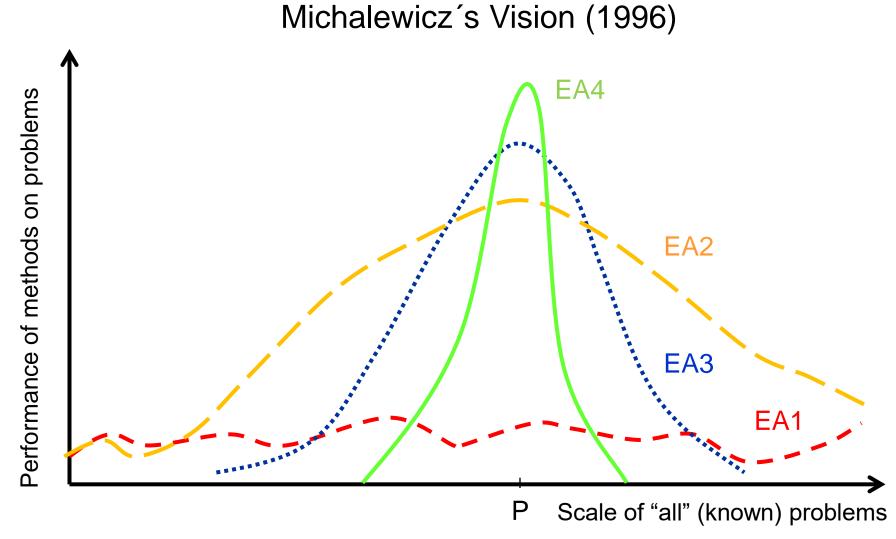
- Implications:
 - New black-box algorithm A is better than old black-box algorithm B (e.g., random walk without revisiting points) in half of the problems (in average)

□ But it is worse in the other half (in average)

> So, why should we bother in creating new algorithms?

- Why are then a lot of people applying EAs and obtaining good results?
 - In fact, <u>most</u> of the "possible" problems are not interesting
 - In general, points in search spaces of real-world problems have continuous (and smooth) neighborhood
 - In such cases, experience has shown that EAs (and other search techniques) perform well





- HOWEVER, BE CAREFUL!
- When EAs (generally) should not be used?
 - Instances of problems that can be solved by exact methods in "reasonable" time
 - <u>Example</u>: The researcher develops a "new" EA and test it in instances of the Travelling Salesman Problem (TSP) with N=100, 200 and 300 cities
 - However, the algorithm Concorde deterministically solves instances of the TSP with hundreds of cities in seconds

- When EAs (generally) should not be used?
 - Problems where it is known that a global optimum can be found in "reasonable" time

Examples

 \Box Many of the problems with polynomial time complexity, i.e., O(n^k)

- When EAs (generally) should not be used?
 - Problems where other optimization algorithms (traditional optimization algorithms, other metaheuristics, heuristics, ...) perform better
 For the given constraints to solve the problem

≻ How do we know this?

i. Experimental comparison

□ traditional method adopted by most of the researches

ii. Theory (when possible)

 Is it possible to "predict" if applying a given EA to my problem will be interesting?

In order to answer this question, we should understand how the EA works from a theoretical point of view

When applicable, theory can

- Provide performance guarantees for the algorithm
 - Examples: runtime analysis
- Help designing new algorithms, operators, or modifications of the known algorithms
- Help understanding the influence of the algorithm's parameters
- Eventually be used to explain phenomena in other areas, e.g., Biology

- Some criticisms to EAs...
 - "There is no guarantee of convergence to global optimum!" (?)
 - "It is not possible to understand how EAs work!" (?)
 - "It is not possible to understand how the parameter's setting influence the performance!" (?)
- In fact, there is a lot of questions that must be answered

We must not rely (only) on the inspiration of evolution by natural selection to justify the use of EAs

- However, some of the criticisms are not exclusive to EAs
 - Example: For all known optimization algorithms, we should be very careful when we speak about convergence for algorithms applied to problems in the NP class

- So, why apparently the theory is much well understood in other algorithms?
- Difficulties with (a) Theory for EAs
 - EAs are vast and complex dynamical systems with many degrees of freedom
 - EAs are generally applied to complex problems with fitness landscapes that are difficult to be properly modeled
 - EAs involve probabilistic operators
 - > We often need statistical tools to analyze them
 - Results are many times described over average behavior
 - Parallel: a predictive model for biological evolution Computação Bioinspirada - 5955010-2

- In Evolutionary Computation, there are more theoretical studies for Evolution Strategies
 - Real codification
 - Most of the papers on Theory deals with well-defined searchspaces

Runtime analysis (time complexity)

Important question: How many iterations until global optima (or very good solutions) found?

http://www.cs.nott.ac.uk/~psxld/seminars/seminar_slides/pkl_seminar.pdf

1.6.3. Theory of GAs: Some Approaches

- Here, we will discuss theory for Genetic Algorithms (GA)
- Some approaches:
 - Schema Theorem
 - ➤GA process investigated as a Markovian Process
 - GA seen as a dynamical systemExact model
 - Mechanical statistics approach
 - Fitness landscapes approach

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• How do GAs work?

One explanation for the operation of GAs is the Building Blocks Hypothesis

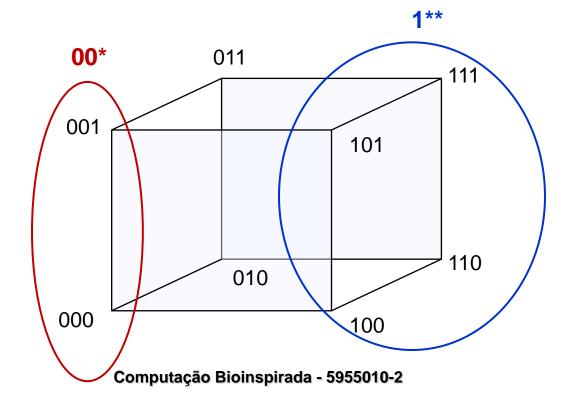
Building block

□ Short low-order **schema** with good fitness

≻Schema

- Template describing a subset of solutions (strings) with similarity in some positions
 - In other words, a schema is a hyperplane in the search space

- Schemata (considering the binary representation)
 - Strings composed by 0, 1, * ("don't care")
 - ➤ Examples



- Why do we use shemata?
 - They represent a subset of solutions (chromosomes) instead of only one solution
 - The analysis of the population becomes easier
 - Some positions of the solutions (genes) may be more important to the optimization process than others
 - Reproduction operators can change a chromosome and not change a schema

Properties of a schema H

Order *o(H)* ➢ Number of fixed positions ➢ Examples: *o*(011*1**) = 4, *o*(0*1*1**) = 3

Defining lenght δ (H)

> Distance between first and last fixed positions > Examples: δ (*011**1*) = 5, δ (0*1***1) = 6

Holland's formulation for the Standard GA

- Standard GA: fitness proportionate parent selection, one point crossover, and bit-flip mutation
- Considering a chromosome of length / that contains a schema H. The probability of disrupting the schema

➢ By crossover is:

$$P_{crossover}(H) = \frac{\delta(H)}{l-1}$$

➢ By mutation is:

$$P_{mutation}(H) = o(H)p_m$$

Holland's formulation for the standard GA

Combining with proportionate selection, we have the equation for the expected number of individuals representing schema *H* in next generation

$$m(H,t+1) \ge m(H,t) \frac{f(H)}{\bar{f}} \left[1 - p_c \frac{\delta(H)}{l-1} - o(H) p_m \right]$$

Schema Theorem

The number of instance within the population of short loworder schemata of above-average fitness will increase exponentially in subsequent generations

Building Blocks Hypothesis

Short low-order schemata of above-average fitness (building blocks) are combined and recombined to form strings with potentially better fitness

- In this way, the complexity of the problem would be reduced
 - Instead of building strings with high fitness directly, a procedure that requires trying all possible combinations of genes, strings each time better would be built by combining the best partial solutions found by various strings with lower order

• The Two-Armed Bandit Problem

Consider a machine with two independent arms
 > One arm pays an average reward of μ₁ (with variance σ₁²) while the other arm pays an average reward of μ₂ (with variance σ₂²)
 > Which arm should we explore?

An "optimal" strategy is to exponentially increase the number of trials in the best observed arm

- The K-Armed Bandit Problem
 - In the GAs, we are not solving the previous problem, but a problem where K hyperplanes (schemata) are explored simultaneously
 - According to Holland, the GA approaches the "optimal" strategy to exponentially increase the number of attempts of the current best hyperplanes (schemata)

- Arguing pro...
 - The building block hypothesis "can" be used to explain why some problems are difficult for GAs

Examples

- Deceptive Problems: when low-order schemata, rather than combining to generate higher-order promising schemata, combine to form schemata that result in suboptimal solutions
- Scaling: when some schemata have very higher fitness when compared to others

- Arguing con...
 - There are criticisms about the schema theorem and the building block hypothesis. Some of them:
 - It does not consider the constructive effects of crossover and mutation
 - > Due to the use of the estimated fitness of a given schema, the theorem says nothing about the future generations from t + 2
 - Problems designed to be easier for the GA according to the building blocks hypothesis (e.g., Royal Road Functions) are sometimes easier for other algorithms, e.g., hill-climbing with random mutation Computação Bioinspirada - 5955010-2

- Arguing pro (again)...
 - Algorithms and operators can be designed to explicitly explore the building blocks

Examples: some Estimation of Distribution Algorithms (EDAs) explicitly identify and recombine building blocks (gene linkage)

Messy GA

Population-based incremental learning (PBIL)

Compact Genetic Algorithm (cGA)

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- Arguing pro (again)...
 - One legacy of the interest in schemata: application of Walsh transforms to binary GAs

Walsh Transforms

- Allow to perform function decomposition for binary representation
 - Parallel: Fourier Transform decomposes a continuous function (signal dependent on time)
 - Parallel: Walsh coefficients similar to Fourier coefficients (frequencies)

□Can be used in theoretical studies

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- Initially proposed by M. Vose
- Simple GA is seen as a discrete dynamical system
 - The Exact Model Approach is also known as Dynamical Systems Approach
 - Dynamical system
 - Mathematical concept in which fixed rules describe the dependence on time of a state in a geometric space (state space)

$$\mathbf{x}(t+1) = \mathbf{f}(\mathbf{x}(t))$$

- GA as a discrete dynamical system
 - Let *n* be the (finite) size of the search space
 ➢ If the chromosome has *l* elements, then *n* = 2^{*l*}
 - In the exact model, all possible candidate solutions are represented in a discrete space with n dimensions
 - Thus, the current population of the GA can be described as an *n*-dimensional vector
 - □ Each element defines the proportion of each candidate solution in the population, i.e., p(k) = v(k) / N, where
 - The k-th element of v indicates the number of copies of the k-th candidate solution in the population of size N

GA as a discrete dynamical system

As the sum of elements in p is equal to 1, the vector population may be described as belonging to a simplex

$$\Lambda = \left\{ \mathbf{p} \in \mathbb{R}^n : p_k \ge 0, \text{ for } k = 0, 1, \dots, n-1 \text{ and } \sum_{k=0}^{n-1} p_k = 1 \right\}$$

Thus, the GA's behavior is seen as a trajectory in a simplex

• GA as a discrete dynamical system

The model considers infinite population (exact model)

 \blacktriangleright However, GAs with finite population can be analyzed

The deviation (relative to the trajectory for the infinite population) is inversely proportional to the population size

Dynamical System of the GA

 $\mathbf{p}(t) = \mathcal{G}(\mathbf{p}(t-1), t)$ $\mathbf{p}(t) = \mathcal{G}^t(\mathbf{p}(0))$

 For the GA with (bit-flip) mutation and proportional selection

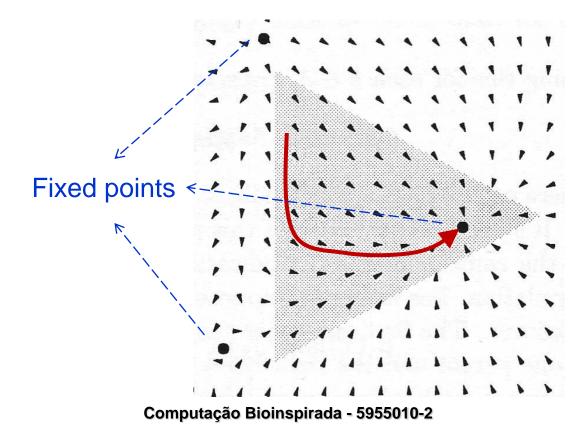
$$\mathcal{G}(\mathbf{p}) = \frac{UF \mathbf{p}}{\mathbf{f}^{\mathrm{T}} \mathbf{p}}$$

 U: nxn matrix representing the transitions due to mutation
 F: nxn diagonal matrix with the fitness of each candidate solution Computação Bioinspirada - 5955010-2

Dynamical System of the GA

Example

Figure adapted from [REEVES & ROWE, 2004]



Some observations

- The existence, location and stability of fixed points and attractors can be defined by the analysis of the generational operator
- For GAs with proportional selection and bit-flip mutation, fixed points and attractors are given by the eigenvectors of UF
- All population trajectories converge to the main fixed point (in which part of the population is in a global optimum)
 In other words: the system is <u>asymptotically stable</u>

1.6.3.2. Theory of GAs: Exact Model

Some observations

The other eigenvectors are related to <u>metastable states</u>

 \blacktriangleright They play very important roles in the evolutionary process

They can change the trajectory in the simplex and trap the population for generations

Local optima

Similar analysis can be made for the case with crossover and with other operators

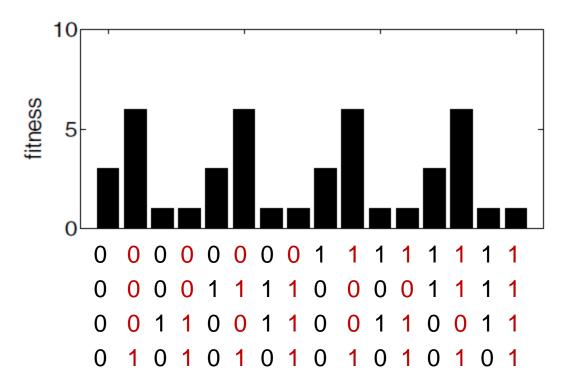
The exact model allow to understand the effects of parameters like population size and mutation rate Computação Bioinspirada - 5955010-2

1.6.3.2. Theory of GAs: Exact Model

Problems

- A very large number of equations to be analyzed for practical problem
 - \blacktriangleright In general, applicable for small solution spaces
- The fitness of all candidate solutions must be known

- Fitness landscape
 - What is a fitness landscape?



- Fitness landscape
 - The landscape observed for a particular function is an artifact of the algorithm used

 \blacktriangleright In other words: of the neighborhood induced by the operators

- Can be defined by the triple (S, n, f) where
 - S: search space
 - \blacktriangleright n(**x**): neighborhood function
 - \succ f(**x**): fitness function
- In this way, when analyzing the fitness landscape, it is essential to analyze the neighborhood structure induced by the operatorsmputação Bioinspirada - 5955010-2

- Fitness landscape
 - How the neighborhood relations are defined?

 \blacktriangleright We can use an adjacency matrix **A**

Using the diagonal matrix, D, containing the degrees of each vertex, we can still define the graph Laplacian

$$\Delta = A-D$$

Elementary landscapes

Fitness landscapes that satisfy for all points s the following equation

 $\Delta f(\mathbf{s}) + (C / m) f(\mathbf{s}) = 0$

 \succ C is a problem-specific parameter

- \succ m is the size of the problem instance
- Several combinatorial optimization problems, e.g., TSP, have elementary landscapes
 - All minima are lower, all maxima are higher than the averaged fitness (f_m) for all candidate solutions in the space
 - Cost to find a local optimum in a maximization problem using neighborhood search (under mild conditions on the nature of the fitness): O($m \log_2(f_{max}/f_m)$)

Dynamic Evolutionary Optimization

EAs applied to Dynamic Optimization Problems (DOPs)

> The fitness landscape changes during the optimization process

Example: Evolutionary Robots

Robots totally or partially designed by EAs

In general, when the fitness of the solutions are experimentally obtained (e.g., when the individual of the EA defines a control law that is tested during a period of time in a real robot), days are required for the optimization process

During this long period, changes often occur:

- In the robot. Examples: Battery charge oscillation, faults, ...
- In the environment. Examples: illumination variation, ...

Problem

 Analysis of the fitness modifications in a dynamic problem with evolutionary robots (simulations)

Problem: simple navigation task

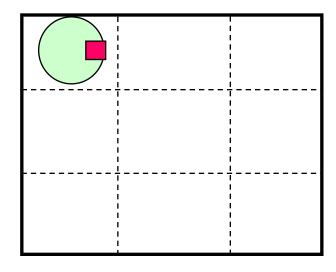
DOP:

Faults occur in the robot during the optimization process

The analysis of the fitness modifications in the problems studied here, and in other problems too, can help the development and analysis of benchmark DOP generators

Problem

- Mobile robot with a frontal sensor
- Controller: *I*-dimensional binary vector (control vector)
 - Indicates the action for each possible state
 - State: input from sensor and memory of last action
- Fitness of the individual (control vector)
 - number of positions occupied by the robot during 10 iterations or until the robot hits a wall



Problem

Model 1 (*I*=4 bits): two actions

Move forward and rotate 90 degrees

- Model 2 (*l*=8 bits): four actions
 - ➢ Move forward, rotate 90 or -90 degrees, wait
- Three faults can occur in the robot
 - Fault 1: sensor inputs always equal to zero
 - ➤ Fault 2: sensor inputs always equal to one
 - Fault 3: wrong sensor inputs
- Each fault represents a different change (DOP)
- The effects of the changes on the fitness vector were analysed and the dynamical system was simulated Computação Bioinspirada - 5955010-2

Simulation: GA with mutation and proportional selection

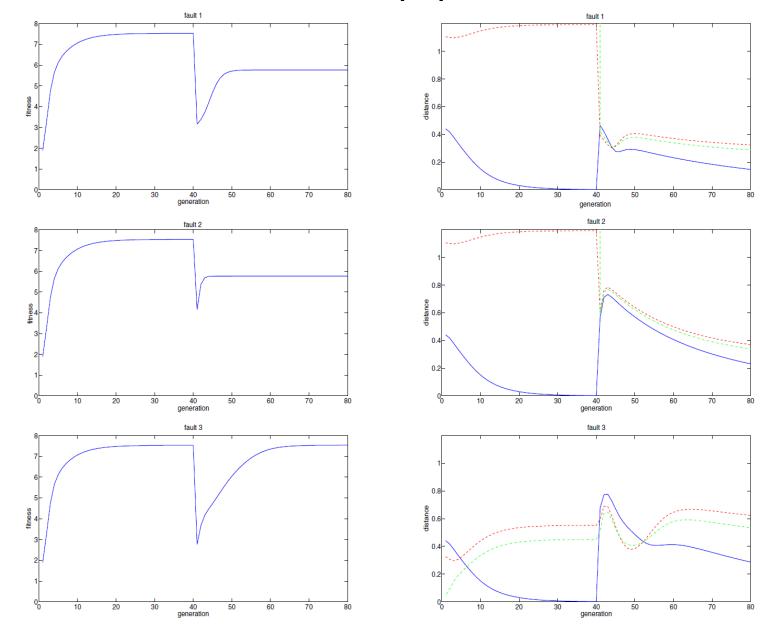


Figure 3: Mean fitness and distances from the population to three metastable states for problem 1 with l = 8. The solid line shows the distance to the main metastable state.

k-bounded pseudo-Boolean optimization

Example: NK Landscape Model

➤Cost function given by

$$f(\mathbf{x}) = \frac{1}{N} \sum_{i=1}^{N} f_i(\mathbf{x}, \mathbf{m}_i)$$

x: solution with size *N*

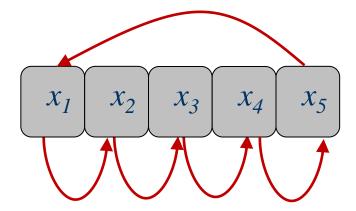
m_{*j*}: binary mask with size *N* and *K*+1 ones

K: integer controlling the epistasis degree

Example

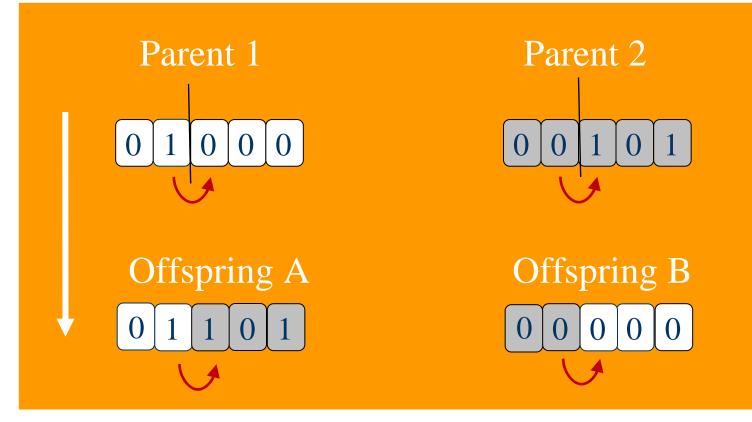
N=5, K=1, adjacent neighborhood

$$f(\mathbf{x}) = \frac{1}{5} (f_1(x_1, x_5) + f_2(x_2, x_1) + f_3(x_3, x_2) + f_4(x_4, x_3) + f_5(x_5, x_4))$$



$$f(\mathbf{x}) = \frac{1}{5} \left(f_1(x_1, x_5) + f_2(x_2, x_1) + f_3(x_3, x_2) + f_4(x_4, x_3) + f_5(x_5, x_4) \right)$$

1-point crossover

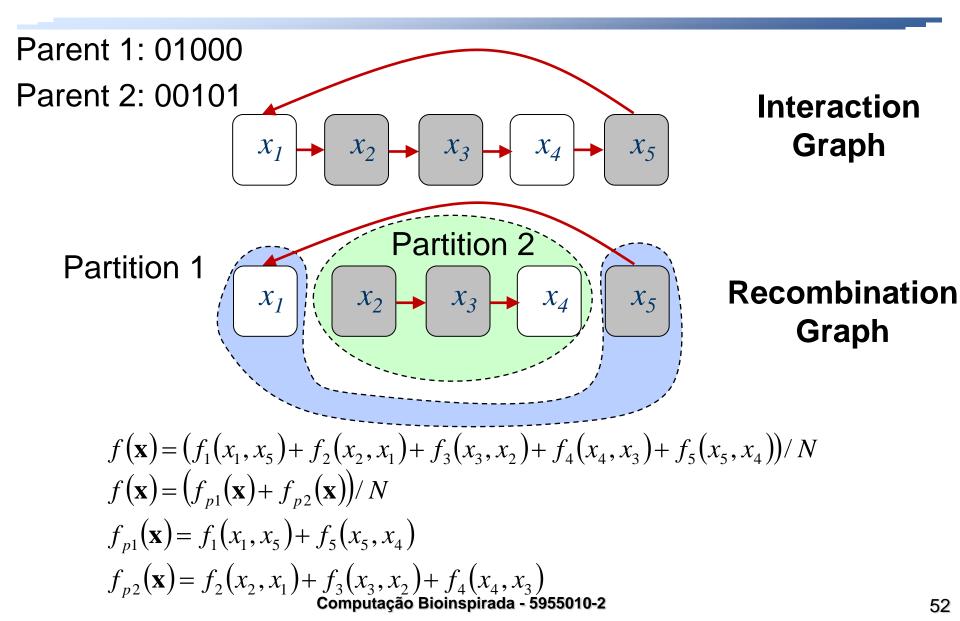


 How do we preserve the interaction of the solution components in order to allow the linear decomposition of the cost function?

$$f(\mathbf{x}) = \frac{1}{5} (f_1(x_1, x_5) + f_2(x_2, x_1) + f_3(x_3, x_2) + f_4(x_4, x_3) + f_5(x_5, x_4))$$

- Solution: we partition the solutions according to the interactions and common characteristics of the parents
 - Recombination by Decomposition
 - ≻ Example:

Partition Crossover



Analysis

q is the number of partitions found

- The decomposition of the evaluation function allows to deterministically find the best among 2^q offspring at the cost of evaluating only two solutions
 - Thus, the cost of finding the best among a number of offspring that grows <u>exponentially</u> with the number of partitions is <u>linear</u> (if the cost of evaluating one solution is linear)

Table 1: Percentage over 50 runs where the global optimum was found (Found) in the experiments of the hybrid GA with the adjacent model. The average percentage difference (% Difference) with respect to the global optimum evaluation is also given.

		2-point crossover		Uniform crossover		PX		PX fit/dist selection	
N	K	Found	% Difference	Found	% Difference	Found	% Difference	Found	% Difference
100	1	100	0.000 ± 0.000	88	0.011 ± 0.051	100	0.000 ± 0.000	100	0.0000 ± 0.0000
100	2	90	0.007 ± 0.029	24	0.294 ± 0.347	100	$0.000\ {\pm}0.000$	100	0.0000 ± 0.0000
100	3	62	$0.086\ {\pm}0.192$	4	$0.657\ {\pm}0.432$	100	$0.000\ \pm 0.000$	100	$0.0000\ \pm 0.0000$
300	1	18	0.090 ± 0.087	0	0.859 ± 0.262	100	0.000 ± 0.000	100	0.0000 ± 0.0000
300	2	0	0.611 ± 0.212	0	2.157 ± 0.431	100	0.000 ± 0.000	100	0.0000 ± 0.0000
300	3	0	1.503 ± 0.402	0	3.464 ± 0.506	80	$0.009\ {\pm}0.023$	98	0.0001 ± 0.0007
500	1	0	0.364 ± 0.153	0	1.371 ± 0.307	100	0.000 ± 0.000	100	0.0000 ± 0.0000
500	2	0	1.398 ± 0.327	0	3.261 ± 0.377	98	$0.001\ {\pm}0.004$	98	$0.0011\ {\pm}0.0078$
500	3	0	$2.791\ {\pm}0.467$	0	$4.851\ {\pm}0.518$	40	$0.029\ {\pm}0.042$	70	$0.0079\ {\pm}0.0183$

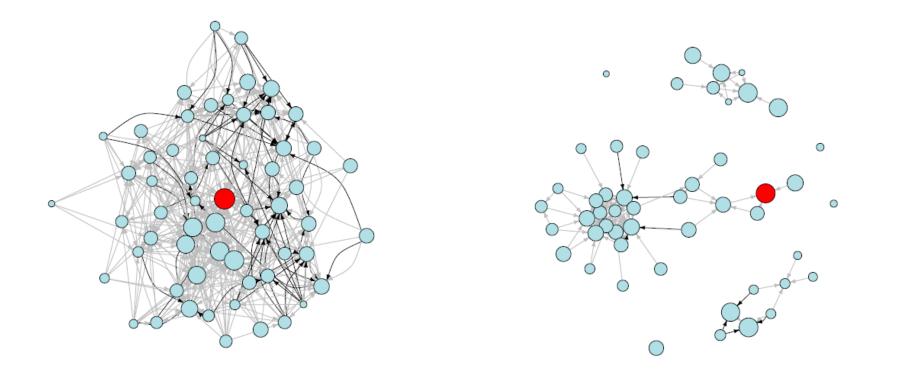


Figure 2: PX local optima networks for two selected instances with values N = 20, K = 2, q = 100. Nodes are local optima and edges connect parents to offspring after partition crossover (black edges indicate PX followed by hill-climbing). Vertex area is proportional to their fitness and the global optimum is highlighted in red (darker color). Left: Adjacent model, the network has 60 nodes and features a single connected component. Right: Random model, the network has 50 nodes (local optima) and features 7 connected components, 4 of which are isolated nodes.

1.6.5. Conclusions

- We must not rely (only) on the inspiration of evolution by natural selection to explain how EAs work
- Theoretical studies are necessary
- There are few theoretical studies in Evolutionary Computation
 - Main difficulties
 - > EAs are vast, non-determistic, complex dynamical systems
 - There is not a general method applicable to all situations
 - □ We do not know completely the fitness landscape in most of the real-word problems

1.6.5. Conclusions

• However, there are several cases of success

Example: runtime analysis for several problems

Theory, when applicable, can

- Provide performance guarantees for the algorithm
- Help designing new algorithms and operators
- Help understanding the influence of the algorithm's parameters
- Eventually be used to explain phenomena in other areas

Anyway, experimental comparison is essential

References

Theory of EAs

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