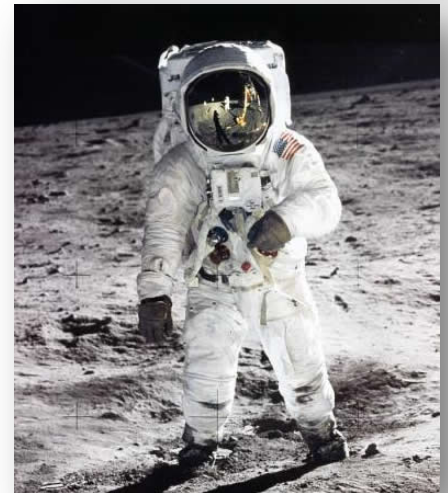


# Advanced Evolutionary Algorithms: An Introduction

Luis Martí Orosa

LIRA/DEE/PUC-Rio

# Why we (must) study AI?



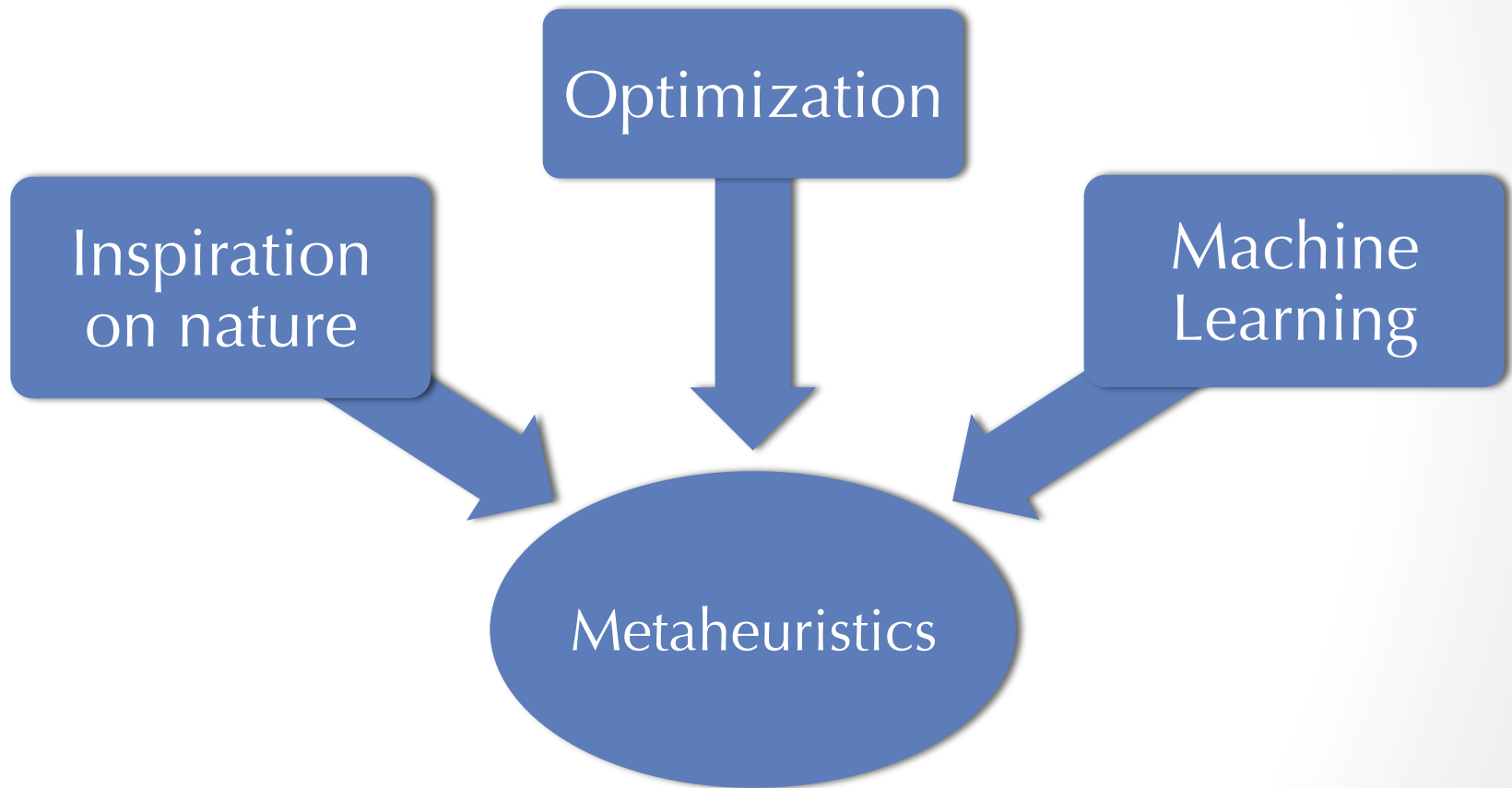
# Intelligence defines us...

...and it would be “great” to have machines that share that feature.

# Inspired by nature

- Neural networks.
- Genetic and evolutionary algorithms.
- Swarm-based approaches.
- Artificial immune systems
- Ants/bees (and other bugs) colonies and emergent behaviors.
- Etc...

# Nature-inspired metaheuristics



# Origin

- Throughout history we have been *fascinated with life*.
- The origin the explanation of life is present in ancient texts.
- In “*western*” countries we have an Abrahamic tradition.
  - Life has been **created** at a given time with a **purpose**.
  - The purpose of life has to do with the human beings.
  - Everything in nature (and therefore life) is ruled by a “supreme entity”.

**... but fossils were discovered.**

# Converging schools of thought

## Jean-Baptiste Lamarck

- Species descend one from other.
- Characteristics acquired during life are passed to offspring.

## Erasmus Darwin

- Organic evolution (similar to Lamarck).
- Introduces the notion of competition for resources and matting.

## Rev. Thomas Robert Malthus.



# Charles Darwin in the Beagle



- During the trip he focused in geology.
  - *Structure and distribution of coral reefs (1842)*



# Conclusions not from the trip

Ornithology



Human selection



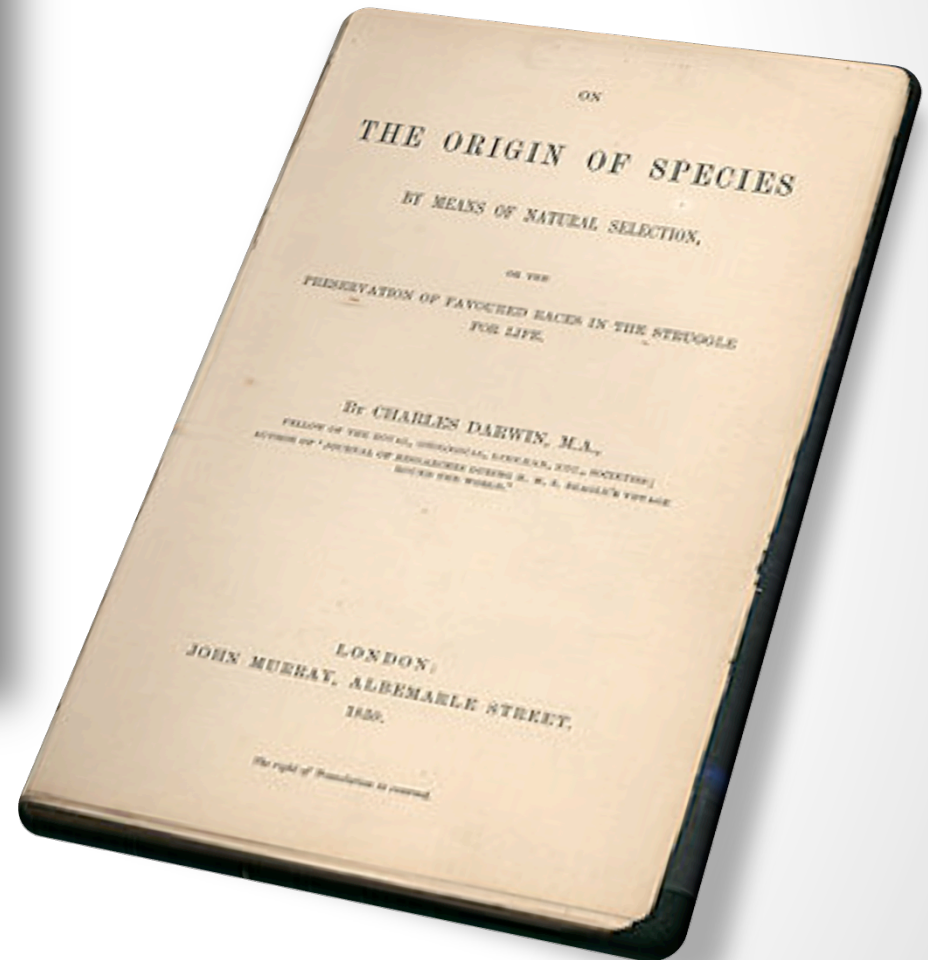
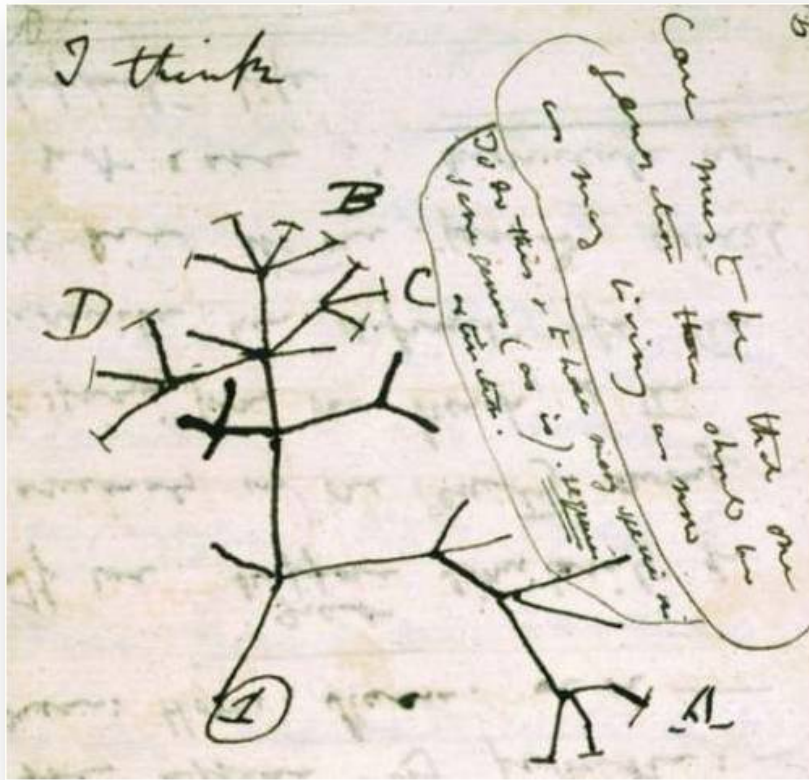
# Alfred Rusell Wallace

- While Darwin was busy “in his garden”, **Wallace** (1823-1913) reached to the same conclusions
  - He was in the Malay Islands.
  - Derived the theory of evolution from Malthus self-regulation of human population studies.
- In 1858 Darwin and Wallace published together their theory.



*Alfred R. Wallace*

# The Origin of Species [...]



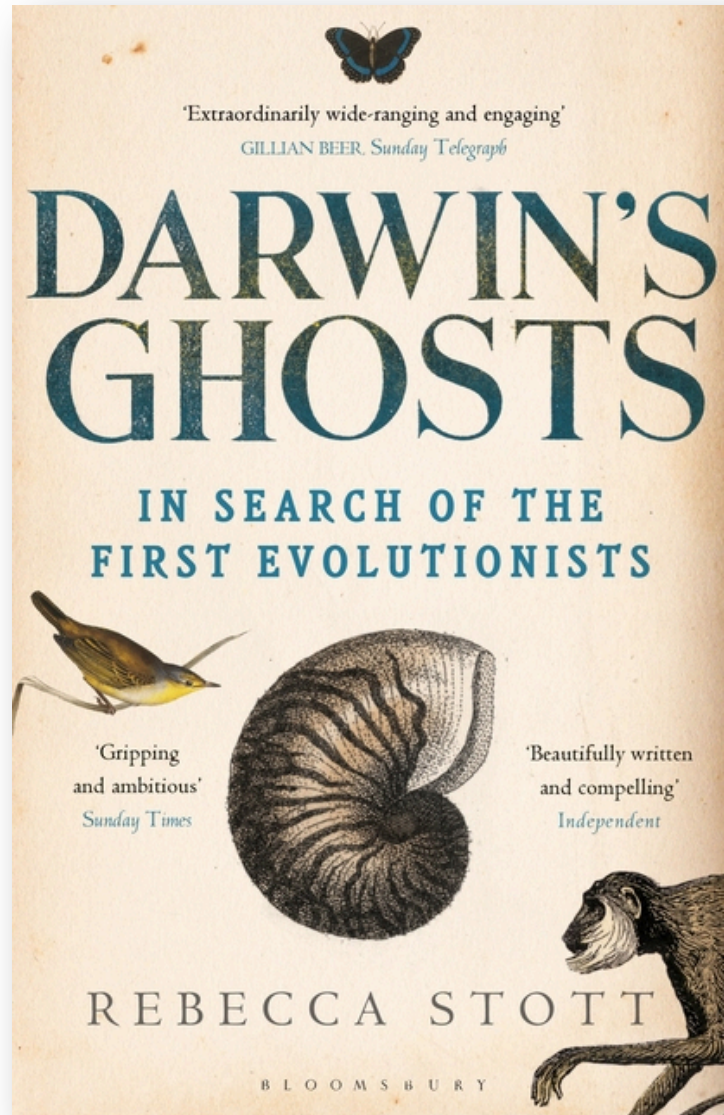
# Theories derived

1. Evolution.
2. Common ancestors.
3. Multiplication of species.
4. Gradualism.
5. Natural selection.

**Darwin was a trending topic!**



# Darwin was not the only one



# Optimization

# Optimization

A key subject:

- Computer Science
- Artificial Intelligence
- Operations Research
- Engineering
- ...

**To optimize is an imprecise term that essentially means “make better”**

# Optimization

- “The process of finding the best solution for a given **optimization problem** with a given resource and temporal **budget**.”
- Optimization problem:
  - Has a number of feasible solutions.
  - There is a clear notion of quality of solutions.
  - The best solution: global optimum



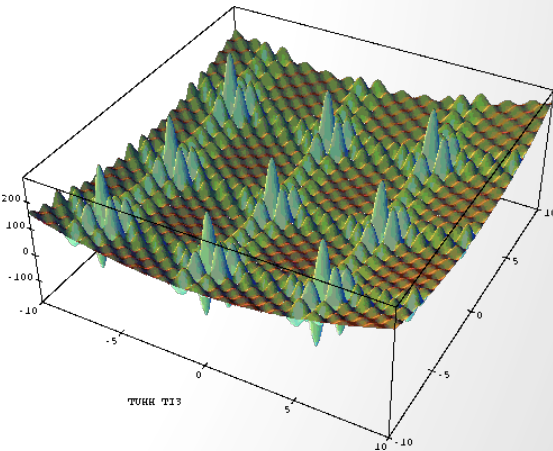
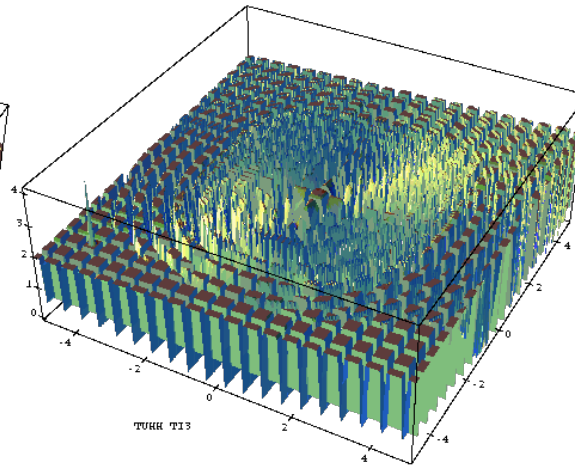
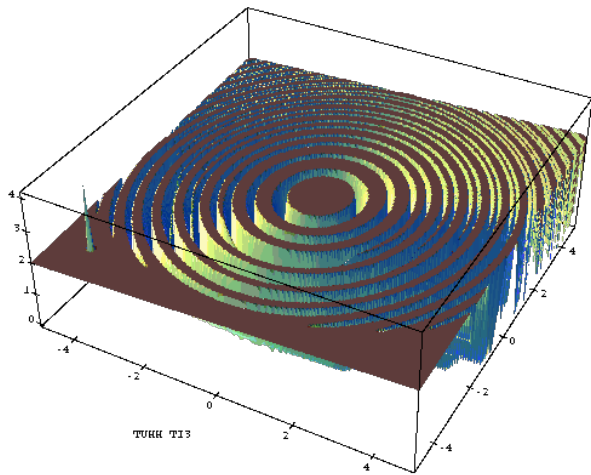
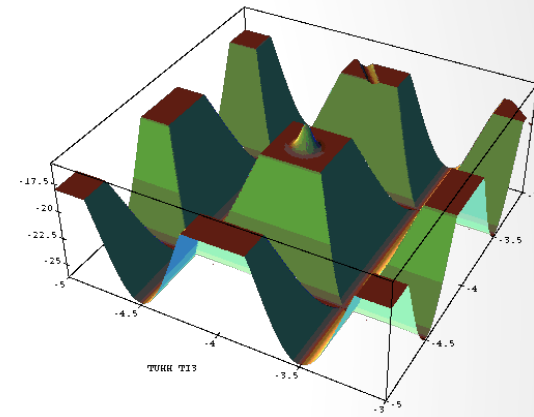
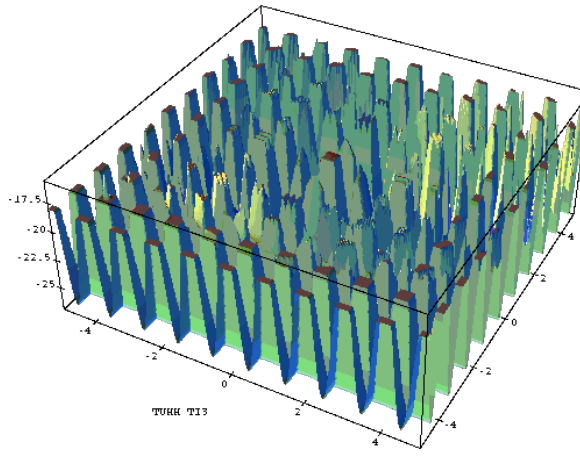
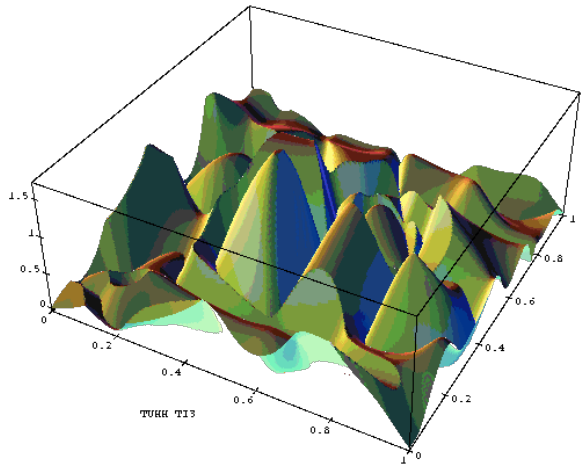
# “Hard” and “easy” problems

- **Tractable:** if there is an algorithm that solves it in polynomial time.
- **Intractable (hard):** if there is no algorithm that solves the problem in polynomial time, NP problems.

# Problemas de Optimización

- We are interested in “hard problems”
- Not warranted that the solution can be found.
- Properties of the problem is unknown.
- We need **metaheuristics**:
  - “Reduced” computational complexity.
  - Do not ensure convergence to the global optimum.

# Optimization problems



# Classes of problems and solvers

- Optimization problems:
  - Combinatorial vs numerical
  - Single vs multi-objective
- Algorithms:
  - Exact:
    - Linear programming
    - Dynamic programming
    - “Branch-and-bound”
  - Approximated or (meta)heuristic

# Iterative Stochastic Methods

## 1. Start:

- **Generate and evaluate** an initial collection of candidate solutions,  $S$ .

## 2. Production:

- Select elements of  $S$ . **Produce and evaluate** a new set of candidate solutions  $S'$  by means of modifications of the selected elements.

## 3. Replacement:

- **Replace** some elements of  $S$  with some elements of  $S'$  and return to 2.

# Why are these methods used?

- Easy to explain and implement.
  - A few lines of pseudocode describe the essential elements of most of these algorithms.
- They are multi-purpose.
  - Do not have strong a priori requirements.
- Proven success.
- Easy to adapt to particular problems with problem dependent (local) methods.

# Montecarlo search

```
t = 0;
result = createNewSolution();
evaluate(result);
while notFinished() do
    a = createNewSolution();
    evaluate(a);
    if a isBetterThan result then
        result = a;
    t = t+1;
end_while
```

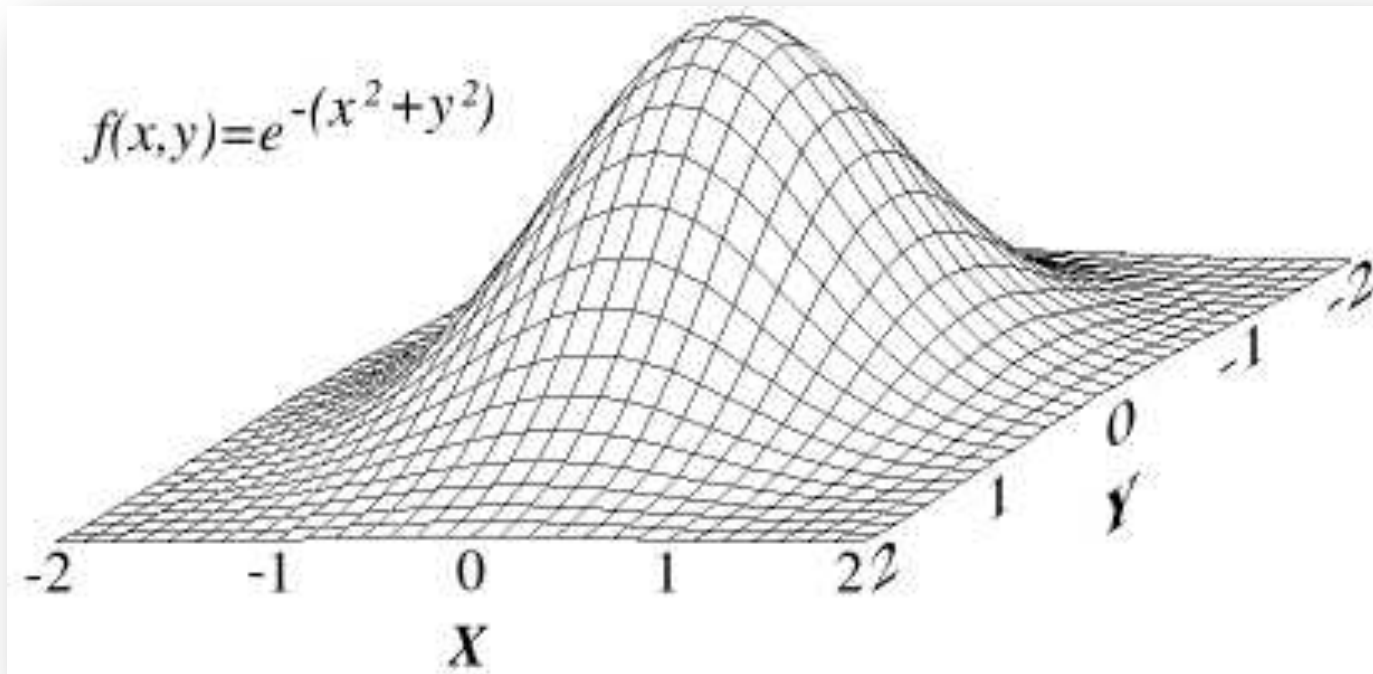
# Hill-Climber

```
t = 0;
result = createNewSolution();
evaluate(result);
while notFinished() do
    a = clone(result);
    mutate(a);
    evaluate(a);
    if a isBetterThan result then
        result = a;
    t = t+1;
end_while
```

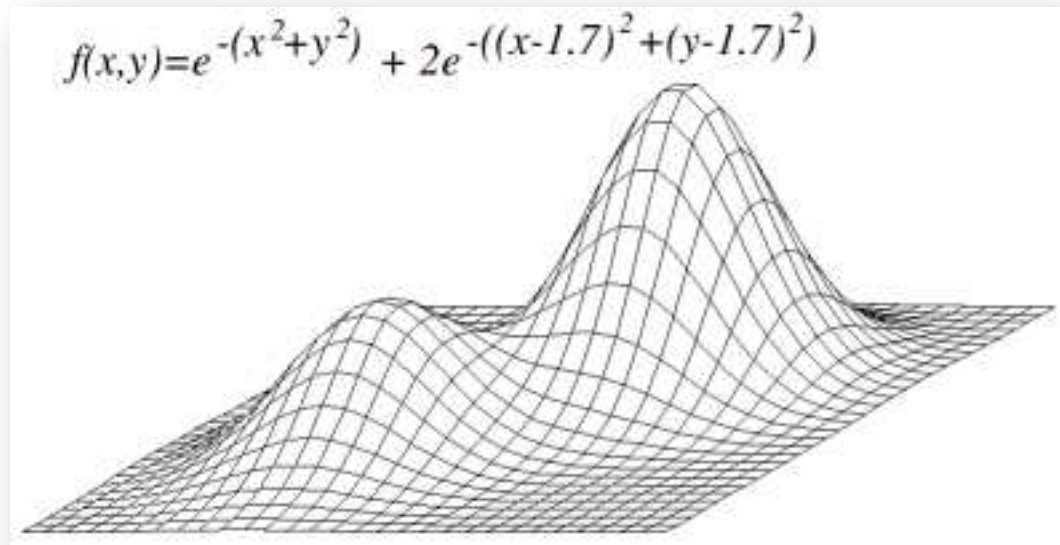


# Sample problem Montecarlo/H.C.

- Only one (global) optimum.



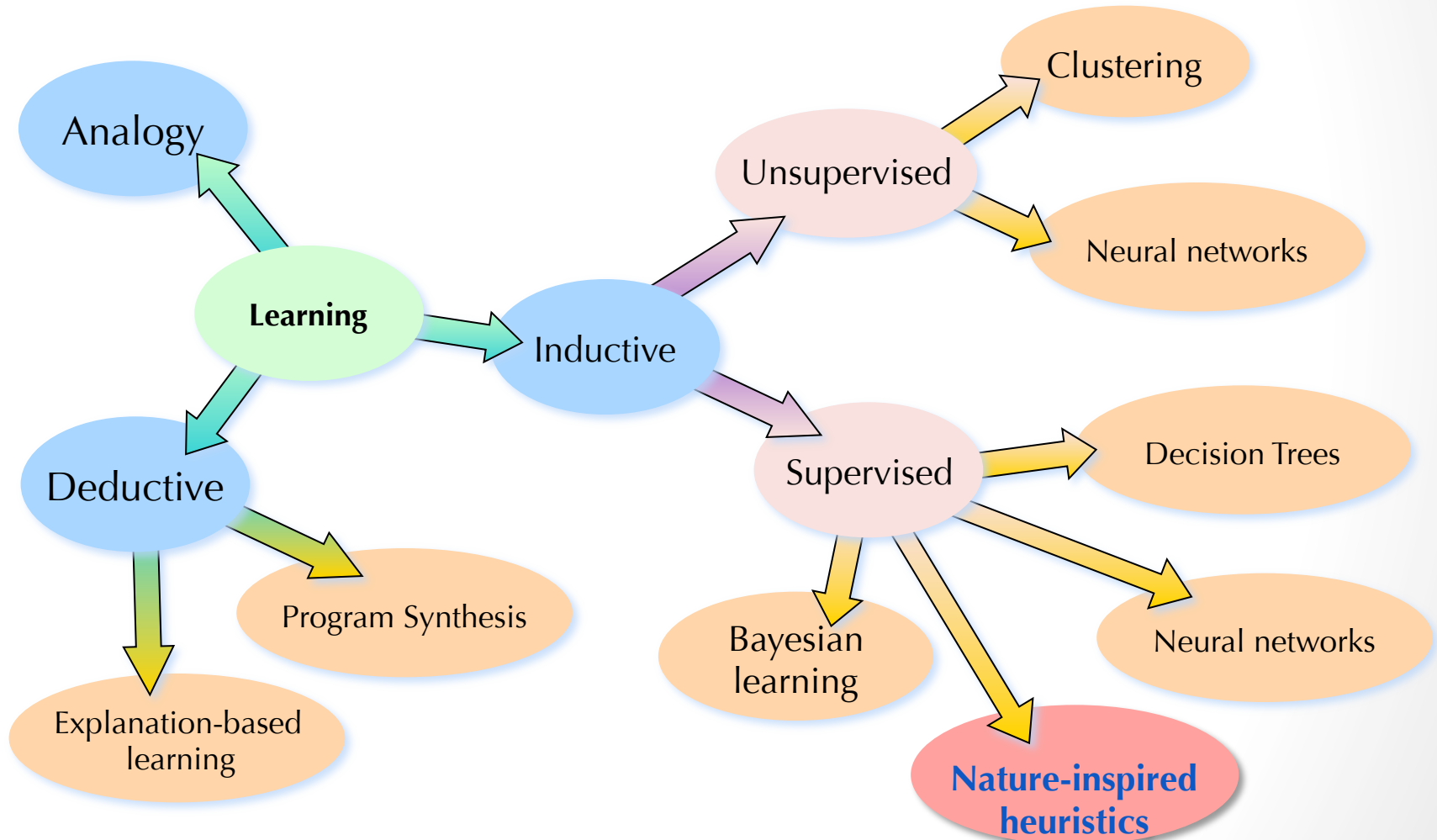
# Local optima



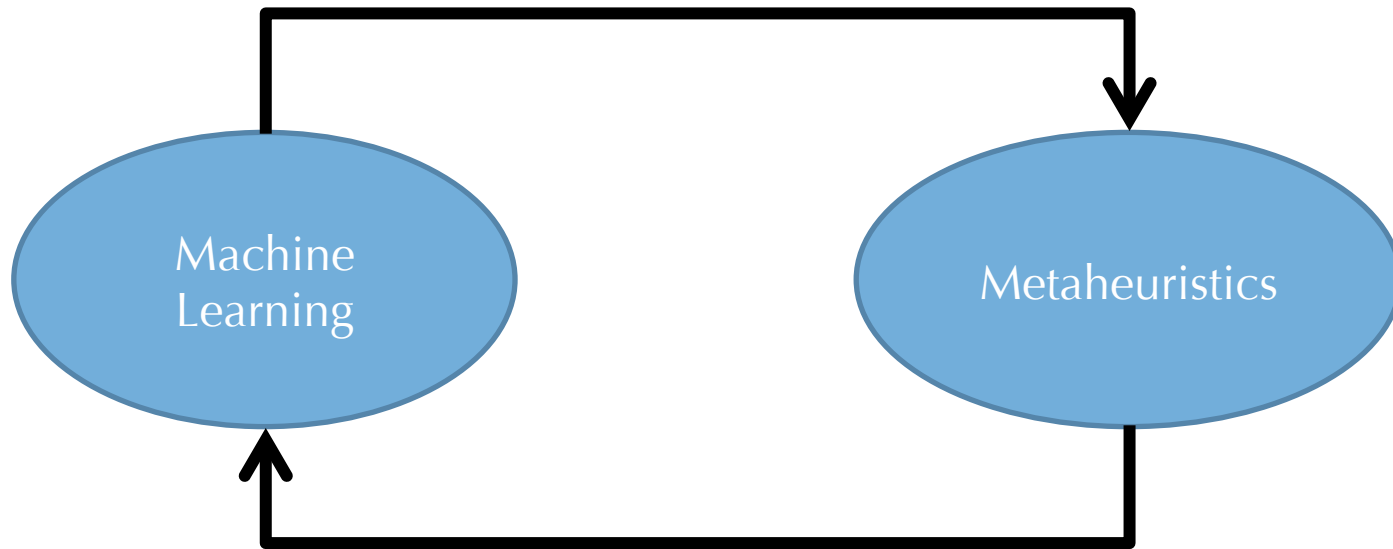
Repeat the algorithm with different initializations.

# Nature-inspired methods

# AI/Machine Learning

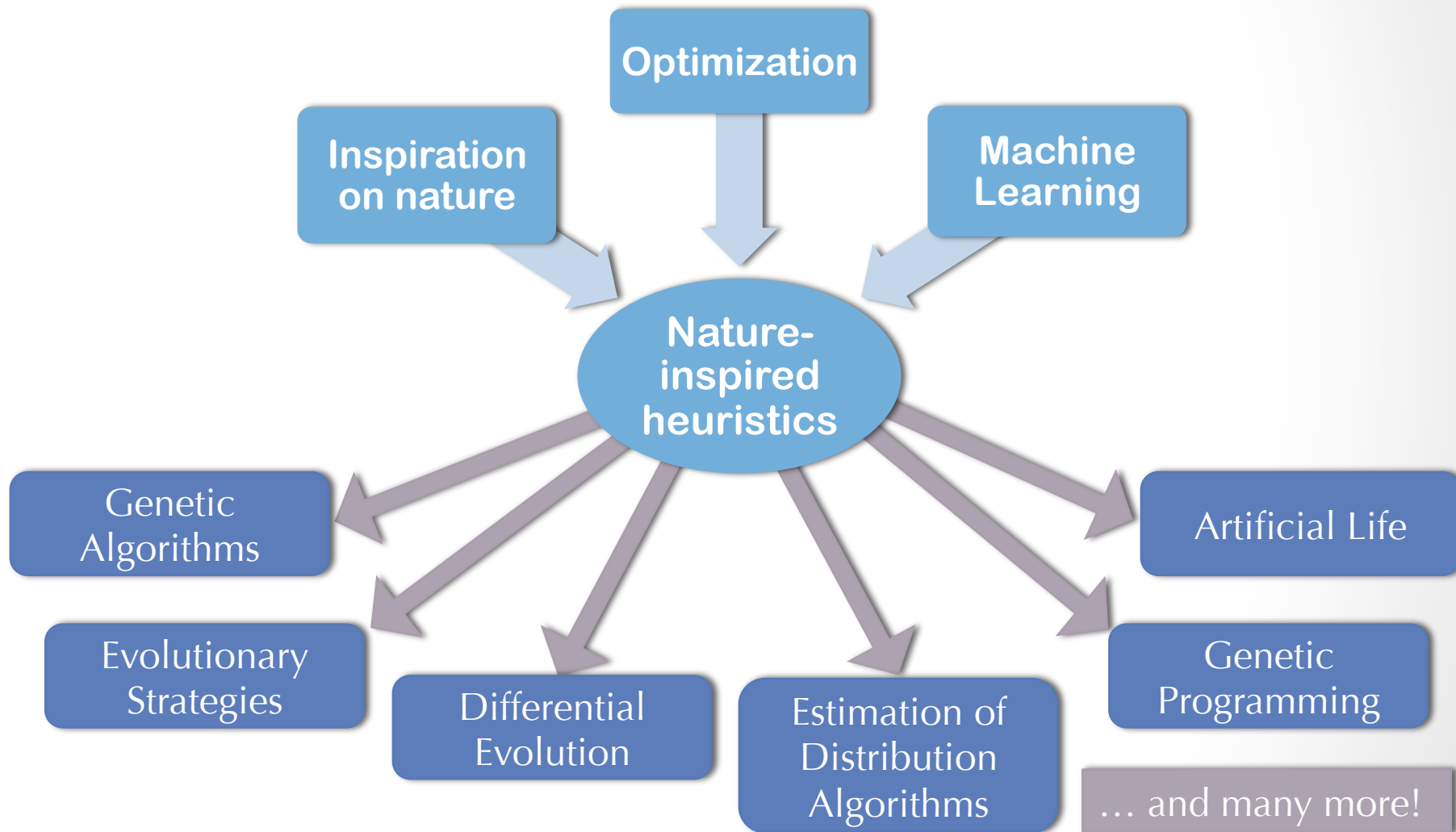


# Metaheuristics and ML



Machine learning approaches exploit metaheuristics and viceversa.

# Nature-inspired metaheuristics



# Nature's optimization algorithm?

- Feasible solutions are represented as a string (ADN).
- Populations of solutions.
- Evaluates every solution (individual) and eliminates the worst.
- **Natural selection** thanks to the survival of the fittest.
- New population combines surviving individuals:
  - Crossover.
  - Mutation.
- Repetition and lots of time.

# Evolutionary computing

- Computational simulation of the processes of evolution and natural selection.
- Mainly inspired by the theory of evolution.
  - Require little information of the problem
  - General purpose
  - Can contain/cooperate with other methodologies
  - Can be used in an “interactive mode”.
  - Inherent parallelism.
  - Robust with respect to data.



# Applications of EC

- As an engineering tool, for finding solutions in optimization problems.
  - Combinatorial and numerical optimization.
  - Planning and control
  - Engineering design
  - Data mining and machine learning applications.
- As a science tool
  - Simulation of real-world phenomena: artificial life, cellular automata, directed evolution, etc.

# Search

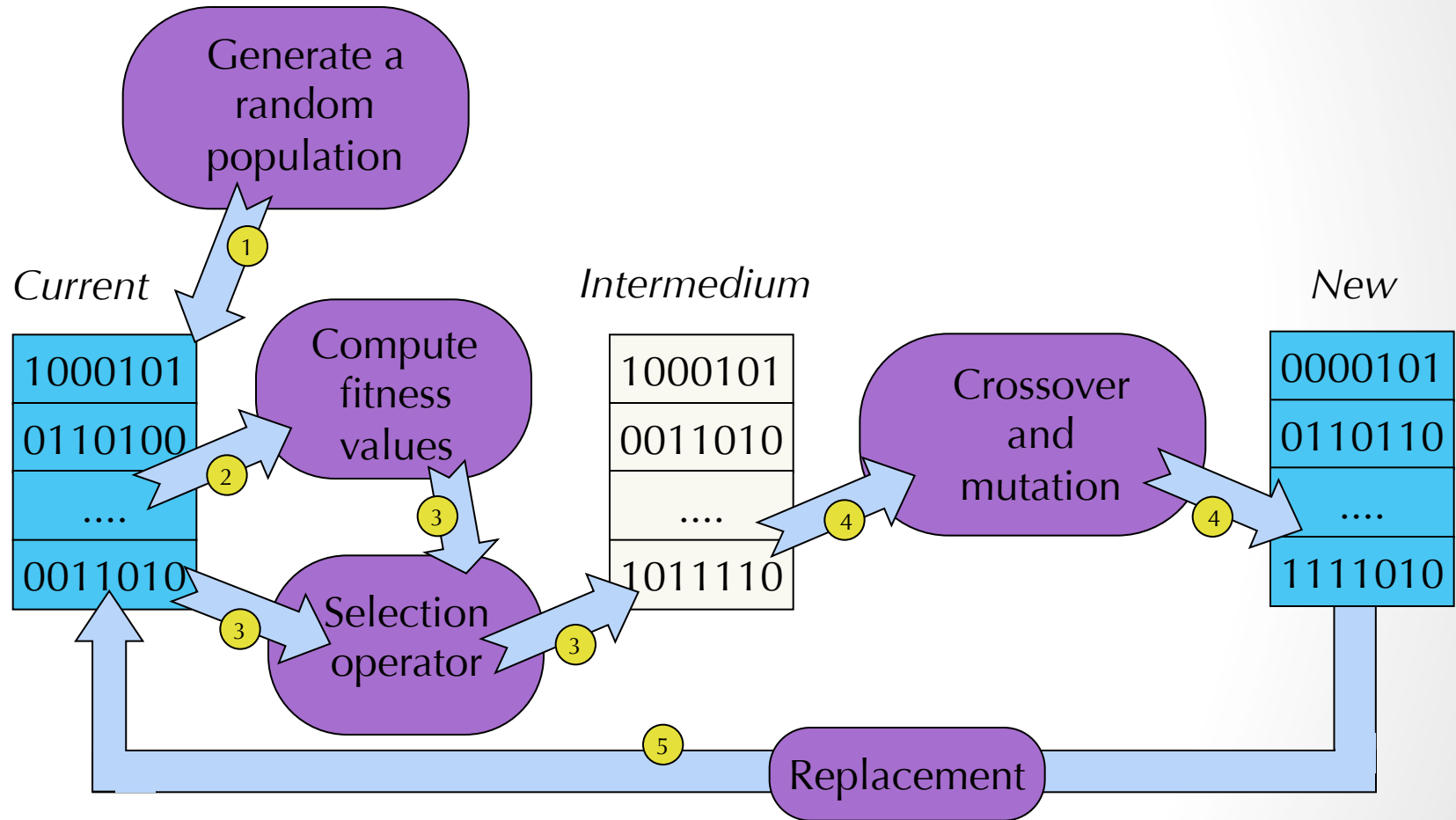
- **Search space:** set of all possible solutions.
  - Its size is an indicator of problem complexity.
- **Crossover operator:** combines characteristics of two or more individuals – **local search**.
- **Mutation:** generates new individuals with different characteristics – **global search**.
- Together they implement a pseudo-random walk:
  - Random, as operators are not deterministic.
  - Directed; as selection is controlled by the fitness function that tends to improve the quality of solutions.

# Genetic Algorithms

- **John Holland**, 1960s
- “Adaptation in natural and artificial systems”, 1975.



# Genetic Algorithm process



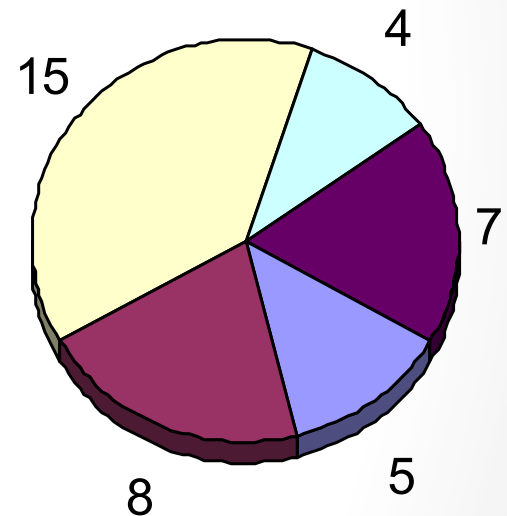
# Using GAs

- Modeling the problem
  1. Decide how to encode information.
  2. Create fitness function.  
**- This is a key part! -**
- Configure GAs
  1. Mating selection and replacement selection.
  2. Type of crossover and mutation.
  3. Parameters

# Selection Schemes

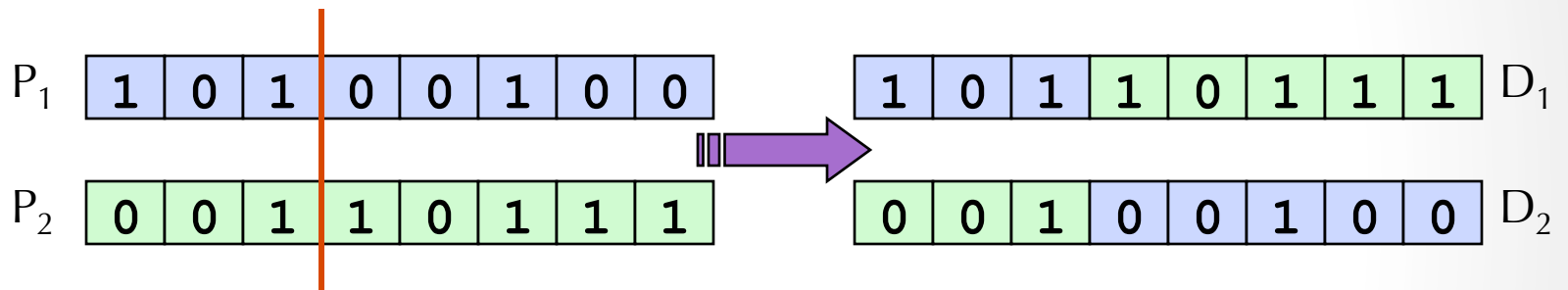
## Roulette wheel selection

- Proportional to fitness or ranking.
- stochastic sampling
  - roulette wheel selection
  - spin wheel N times
- stochastic universal sampling
  - roulette wheel selection
  - single spin, wheel has N equally spaced markers
- tournament selection
  - choose  $k$  candidates at random with uniform probability
  - pick best one for reproduction

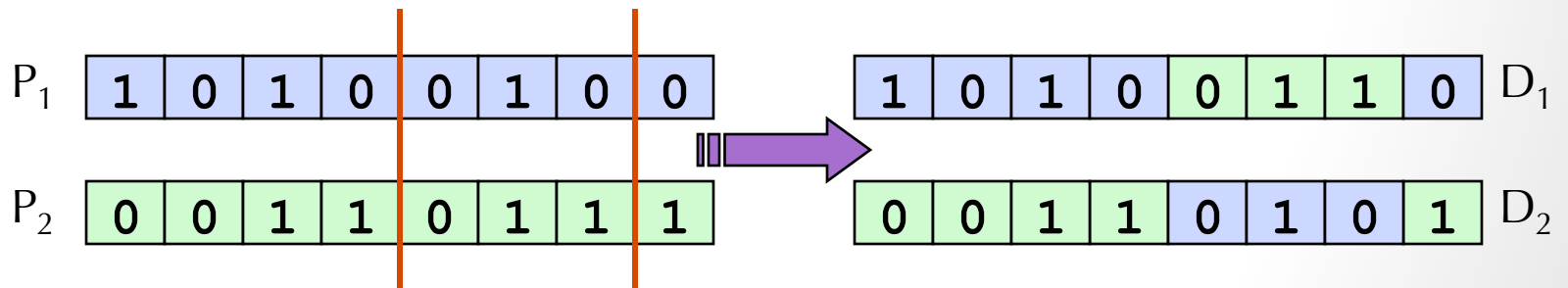


# Crossover

- 1 point cross-over



- 2-points cross-over



# Mutation

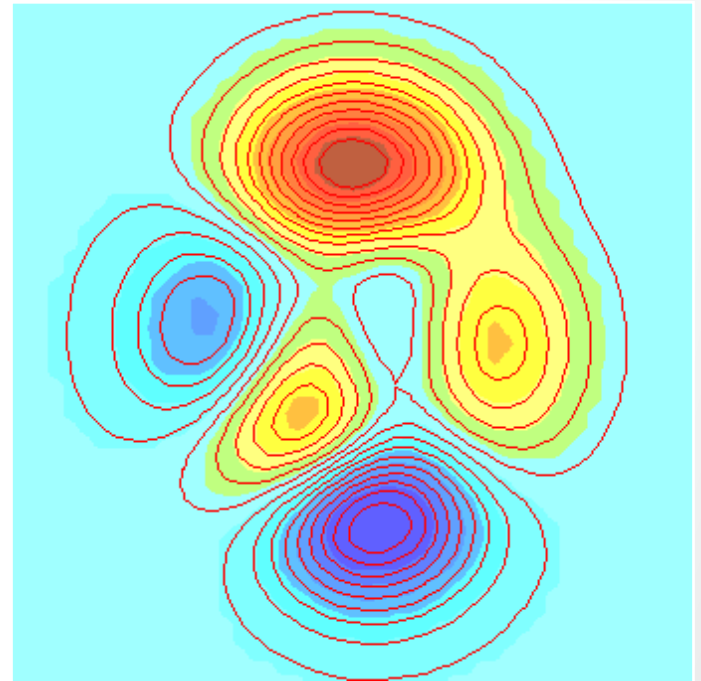
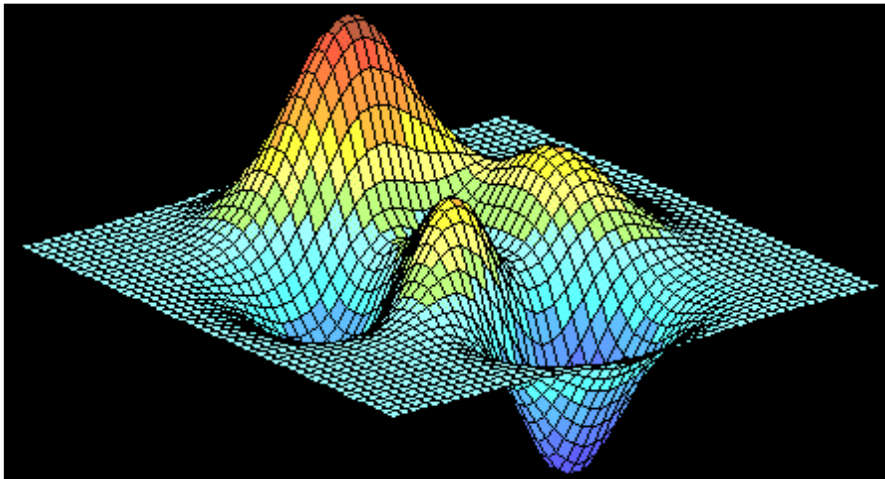
- Every gene is examined.
- An allele is mutated with a low probability,  $p_m$  (0.001-0.1)%





# GAs at work

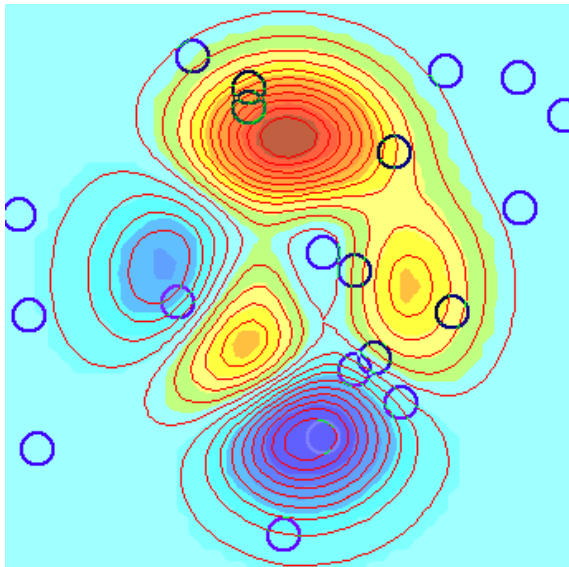
$$z = f(x, y) = 3(1 - x)^2 e^{-(x^2 + (y+1)^2)} - 10\left(\frac{x}{5} - x^3 - y^5\right) e^{-(x^2 + y^2)} - \frac{1}{3} e^{-((x+1)^2 + y^2)}$$



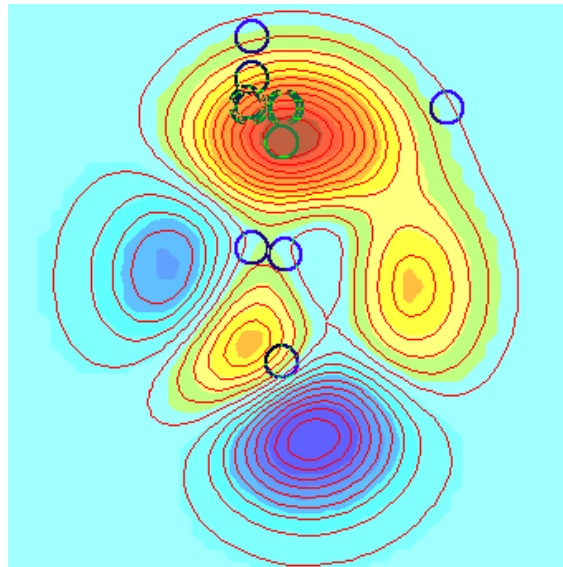
# GAs at work

- Population as iterations advance

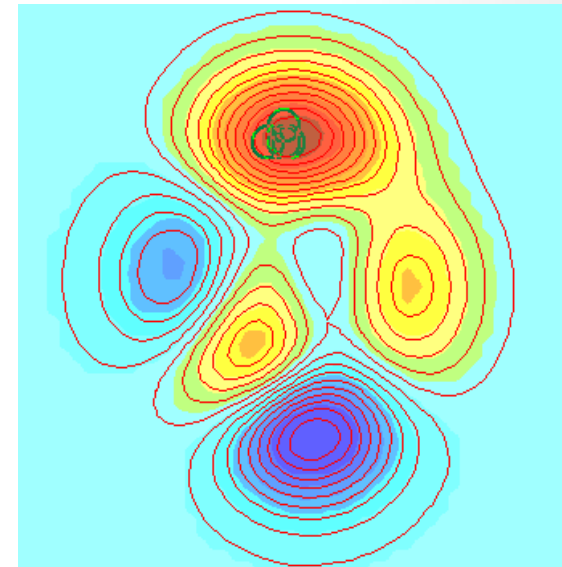
■  $t = 0$



■  $t = 5$



■  $t=10$



# Advanced GAs

- Diploid crossover.
- Multi-objective approaches.
- Knowledge-based methods.
- Multiple populations
- Coevolution
- Parallelization

<http://boxcar2d.com/>

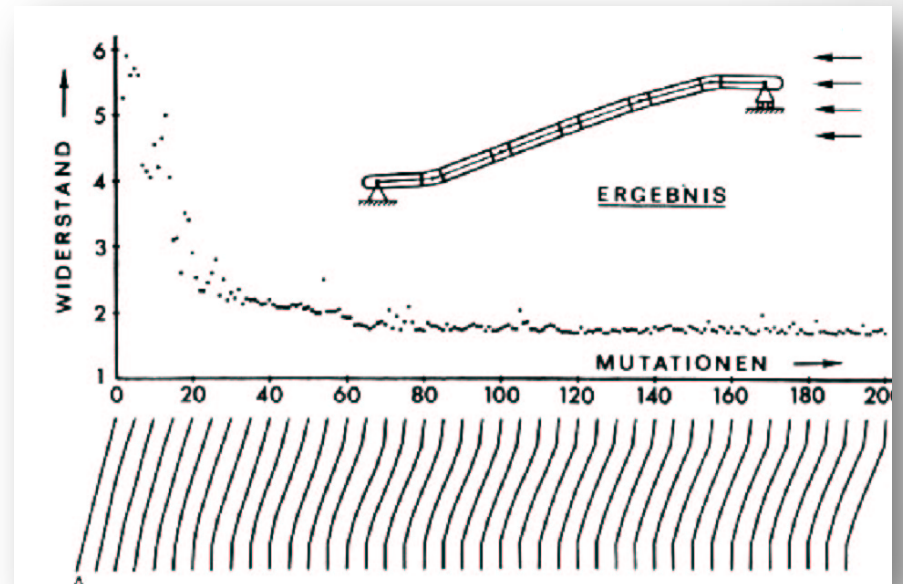
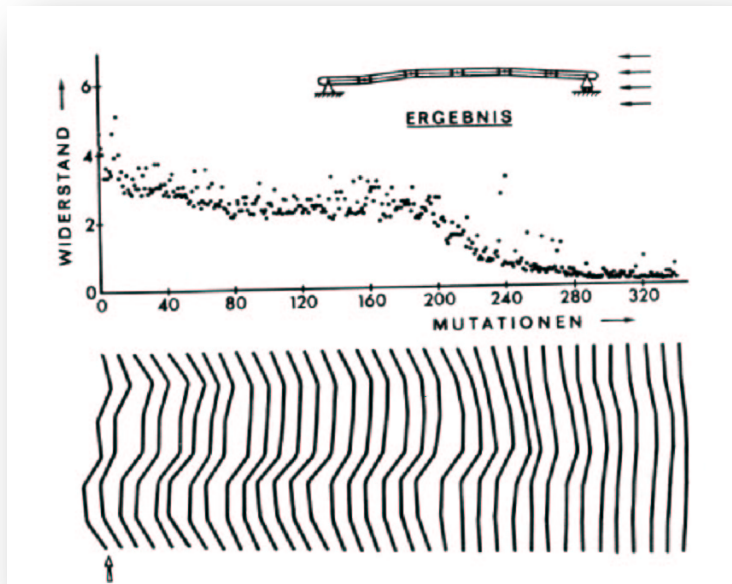
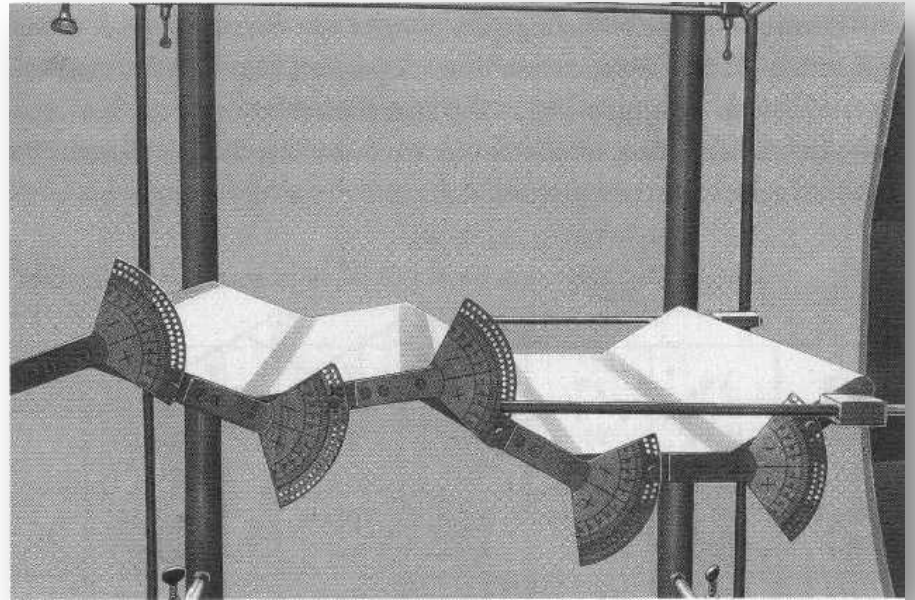
# Evolutionary Strategies

# Evolutionary Strategies

- Proposed in the 60s by Rechenberg and Schwefel.
- Method of parametric (numeric) optimization.
- Only mutation, with self-adaptation.
- Classes:
  - Simple EE (population of two)
  - Multiple EE (more elements)
- Characteristics
  - Fast
  - Solid theoretical foundation
  - Good results in numerical optimization.

# Initial steps

- Airfoil profile



# ES technical summary tableau

Representation	Real-valued vectors
Recombination	Discrete or intermediary
Mutation	Gaussian perturbation
Parent selection	Uniform random
Survivor selection	$(\mu, \lambda)$ or $(\mu + \lambda)$
Specialty	Self-adaptation of mutation step sizes

# Simple (1,1) Pseudocode

Set  $t = 0$

Create initial point  $\mathbf{x}^t = \langle \mathbf{x}_1^t, \dots, \mathbf{x}_n^t \rangle$

**repeat**

Draw  $z_i$  from a normal distr. for all  $i = 1, \dots, n$

$$\mathbf{y}_i^t = \mathbf{x}_i^t + z_i$$

**IF**  $f(\mathbf{x}^t) < f(\mathbf{y}^t)$  **THEN**

$$\mathbf{x}^{t+1} = \mathbf{x}^t$$

**ELSE**

$$\mathbf{x}^{t+1} = \mathbf{y}^t$$

**END\_IF**

Set  $t = t+1$

**until** *endCondition()*



# Representation

- Chromosomes consist of three parts:
  - Object variables:  $x_1, \dots, x_n$
  - Strategy parameters:
    - Mutation step sizes:  $\sigma_1, \dots, \sigma_{n_\sigma}$
    - Rotation angles:  $\alpha_1, \dots, \alpha_{n_\alpha}$
- Not every component is always present
- Full size:  $\langle x_1, \dots, x_n, \sigma_1, \dots, \sigma_n, \alpha_1, \dots, \alpha_k \rangle$
- where  $k = n(n-1)/2$  (no. of  $i, j$  pairs)

# Mutation

- Main mechanism: changing value by adding random noise drawn from normal distribution
- $x'_i = x_i + N(0, \sigma)$ 
  - $\sigma$  is part of the chromosome  $\langle x_1, \dots, x_n, \sigma \rangle$
  - $\sigma$  is also mutated into  $\sigma'$
- Thus: mutation step size  $\sigma$  is coevolving with the solution  $x$

# Mutate $\sigma$ first

- Net mutation effect:  $\langle x, \sigma \rangle \rightarrow \langle x', \sigma' \rangle$
- Order is important:
  - first  $\sigma \rightarrow \sigma'$  (see later how)
  - then  $x \rightarrow x' = x + N(0, \sigma')$
- Rationale: new  $\langle x', \sigma' \rangle$  is evaluated twice
  - Primary:  $x'$  is good if  $f(x')$  is good
  - Secondary:  $\sigma'$  is good if the  $x'$  it created is good
- Reversing order would not work

# Mutation case 1:

## Uncorrelated mutation with one $\sigma$

- Chromosomes:  $\langle x_1, \dots, x_n, \sigma \rangle$
- $\sigma' = \sigma \cdot \exp(\tau \cdot N(0,1))$
- $x'_i = x_i + \sigma' \cdot N(0,1)$
- Typically the “learning rate”  $\tau \propto 1/n^{1/2}$
- And we have a boundary rule  $\sigma' < \varepsilon_0 \Rightarrow \sigma' = \varepsilon_0$

# Mutation case 2:

## Uncorrelated mutation with $n$ $\sigma$ 's

- Chromosomes:  $\langle x_1, \dots, x_n, \sigma_1, \dots, \sigma_n \rangle$
- $\sigma'_i = \sigma_i \cdot \exp(\tau' \cdot N(0,1) + \tau \cdot N_i(0,1))$
- $x'_i = x_i + \sigma'_i \cdot N_i(0,1)$
- Two learning rate parameters:
  - $\tau'$  overall learning rate
  - $\tau$  coordinate wise learning rate
- $\tau \propto 1/(2n)^{1/2}$  and  $\tau \propto 1/(2n^{1/2})^{1/2}$
- And  $\sigma'_i < \varepsilon_0 \Rightarrow \sigma'_i = \varepsilon_0$

# Mutation case 3:

## Correlated mutations

- Chromosomes:  $\langle x_1, \dots, x_n, \sigma_1, \dots, \sigma_n, \alpha_1, \dots, \alpha_k \rangle$
- where  $k = n \cdot (n-1)/2$
- and the covariance matrix  $C$  is defined as:
  - $c_{ii} = \sigma_i^2$
  - $c_{ij} = 0$  if  $i$  and  $j$  are not correlated
  - $c_{ij} = 1/2 \cdot (\sigma_i^2 - \sigma_j^2) \cdot \tan(2 \alpha_{ij})$  if  $i$  and  $j$  are correlated
- Note the numbering / indices of the  $\alpha$ 's

# Correlated mutations cont'd

The mutation mechanism is then:

- $\sigma'_i = \sigma_i \cdot \exp(\tau' \cdot N(0,1) + \tau \cdot N_i(0,1))$
- $\alpha'_j = \alpha_j + \beta \cdot N(0,1)$
- $\mathbf{x}' = \mathbf{x} + \mathbf{N}(\mathbf{0}, \mathbf{C}')$ 
  - $\mathbf{x}$  stands for the vector  $\langle x_1, \dots, x_n \rangle$
  - $\mathbf{C}'$  is the covariance matrix  $\mathbf{C}$  after mutation of the  $\alpha$  values
- $\tau \propto 1/(2n)^{1/2}$  and  $\tau' \propto 1/(2n^{1/2})^{1/2}$  and  $\beta \approx 5^\circ$
- $\sigma'_i < \varepsilon_0 \Rightarrow \sigma'_i = \varepsilon_0$  and
- $|\alpha'_j| > \pi \Rightarrow \alpha'_j = \alpha'_j - 2\pi \text{sign}(\alpha'_j)$

# Recombination

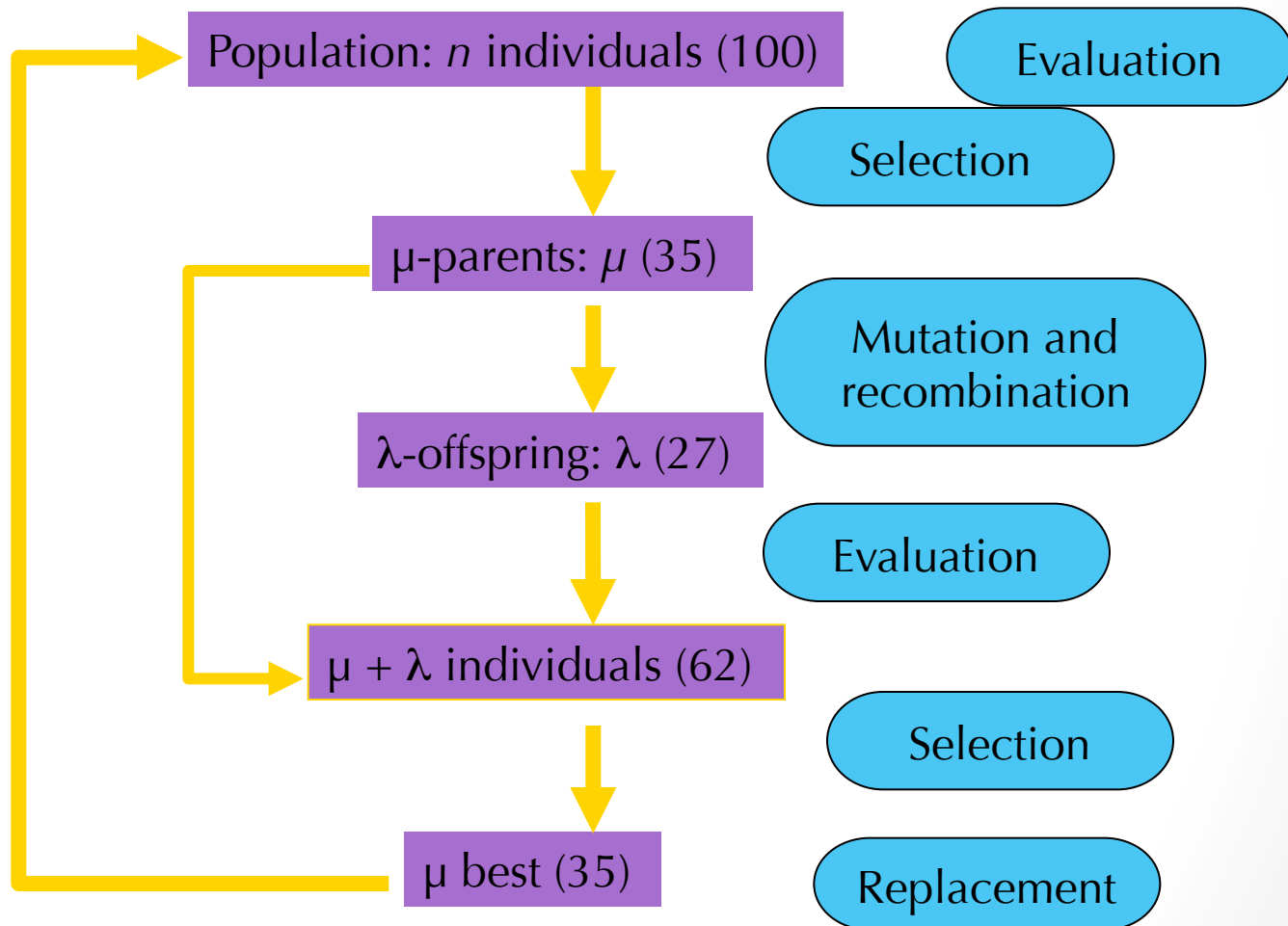
- Creates one child
- Acts per variable / position by either
  - Averaging parental values, or
  - Selecting one of the parental values
- From two or more parents by either:
  - Using two selected parents to make a child.
  - Selecting two parents for each position.



# Names of recombinations

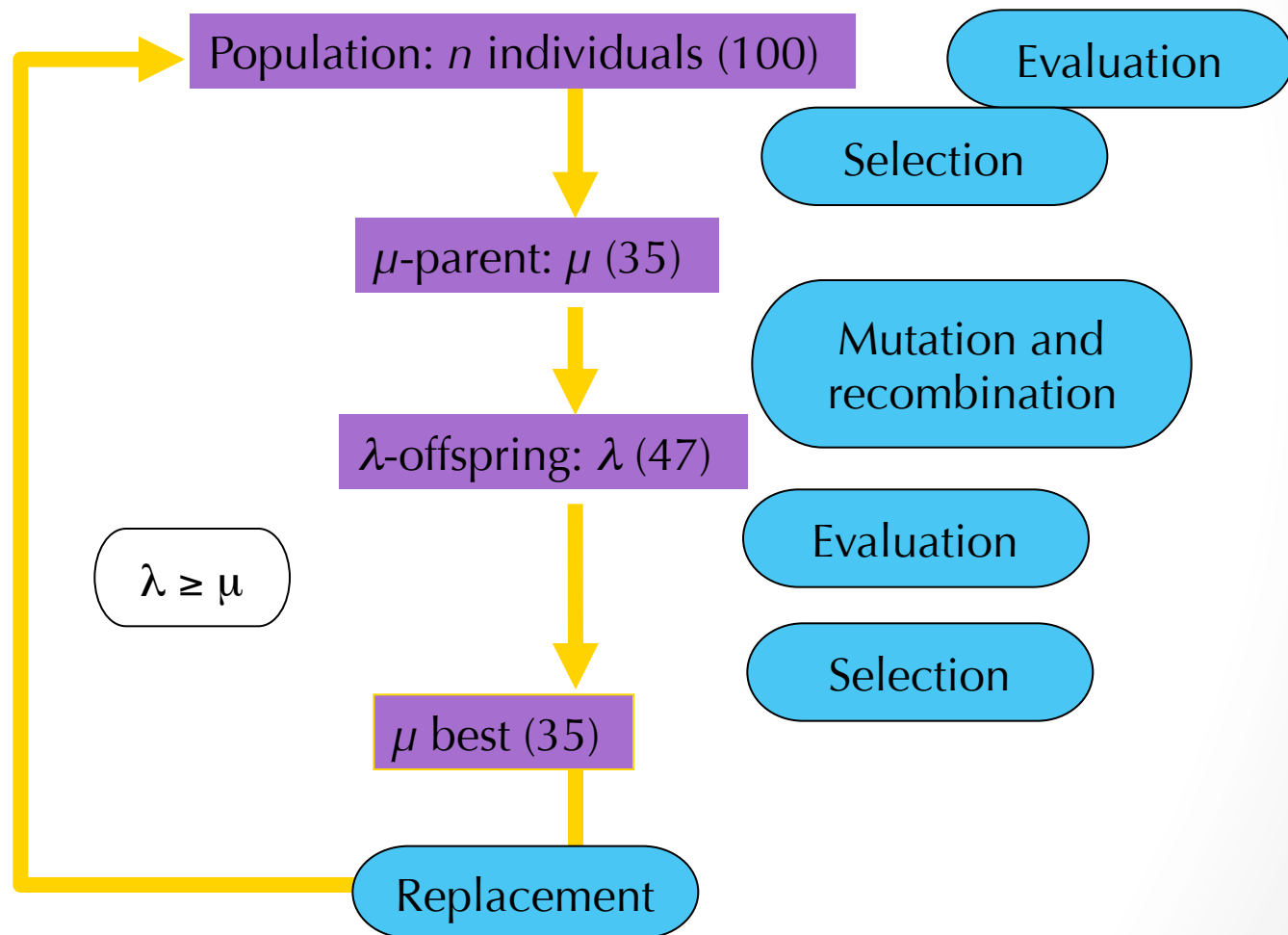
	Two fixed parents	Two parents selected for each $i$
$z_i = (x_i + y_i)/2$	Local intermediary	Global intermediary
$z_i$ is $x_i$ or $y_i$ chosen randomly	Local discrete	Global discrete

# $(\mu+\lambda)$ Evolutionary Strategies

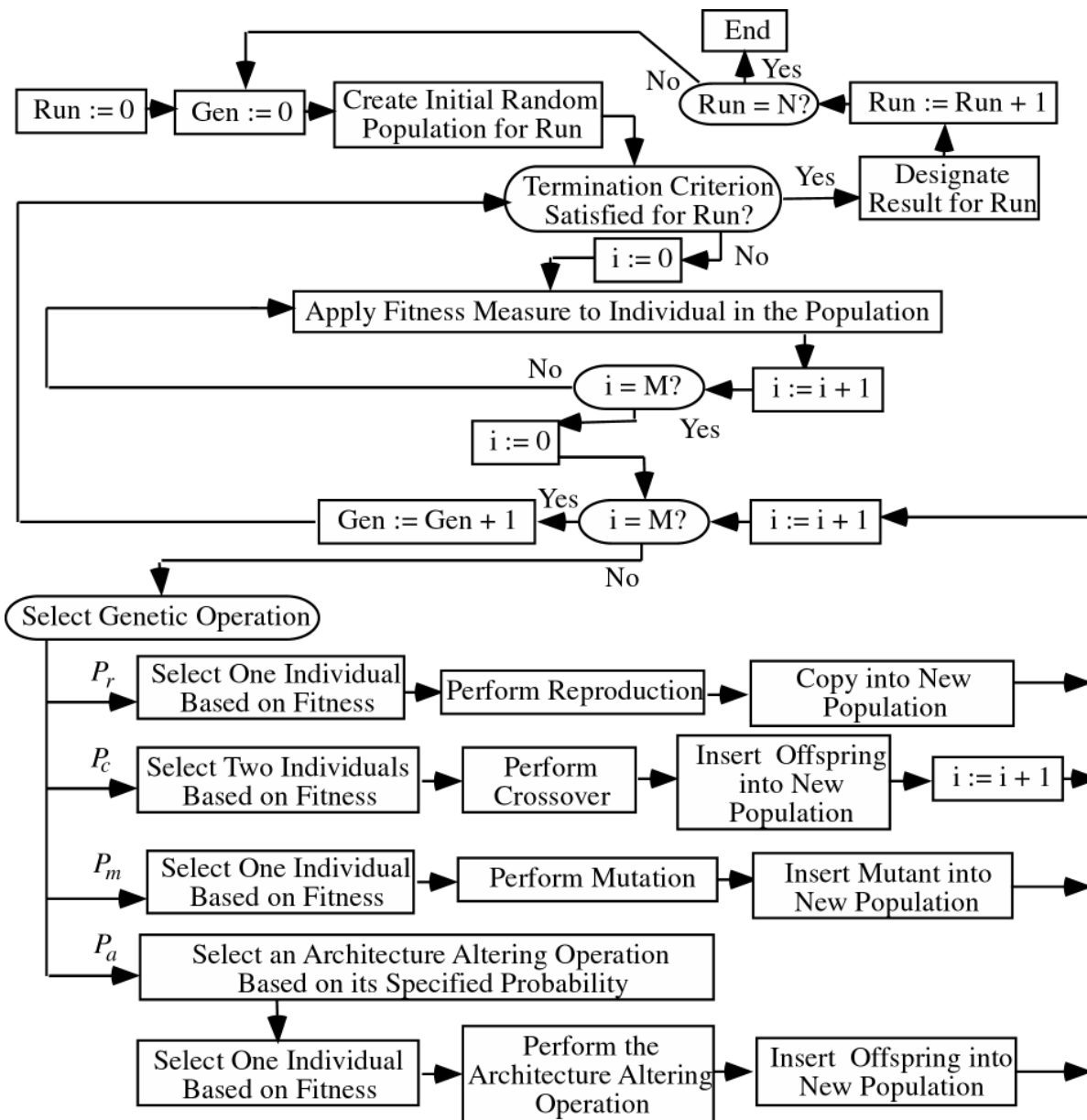


# Estrategias Evolutivas

## Tipo ( $\mu$ , $\lambda$ )



# Genetic Programming



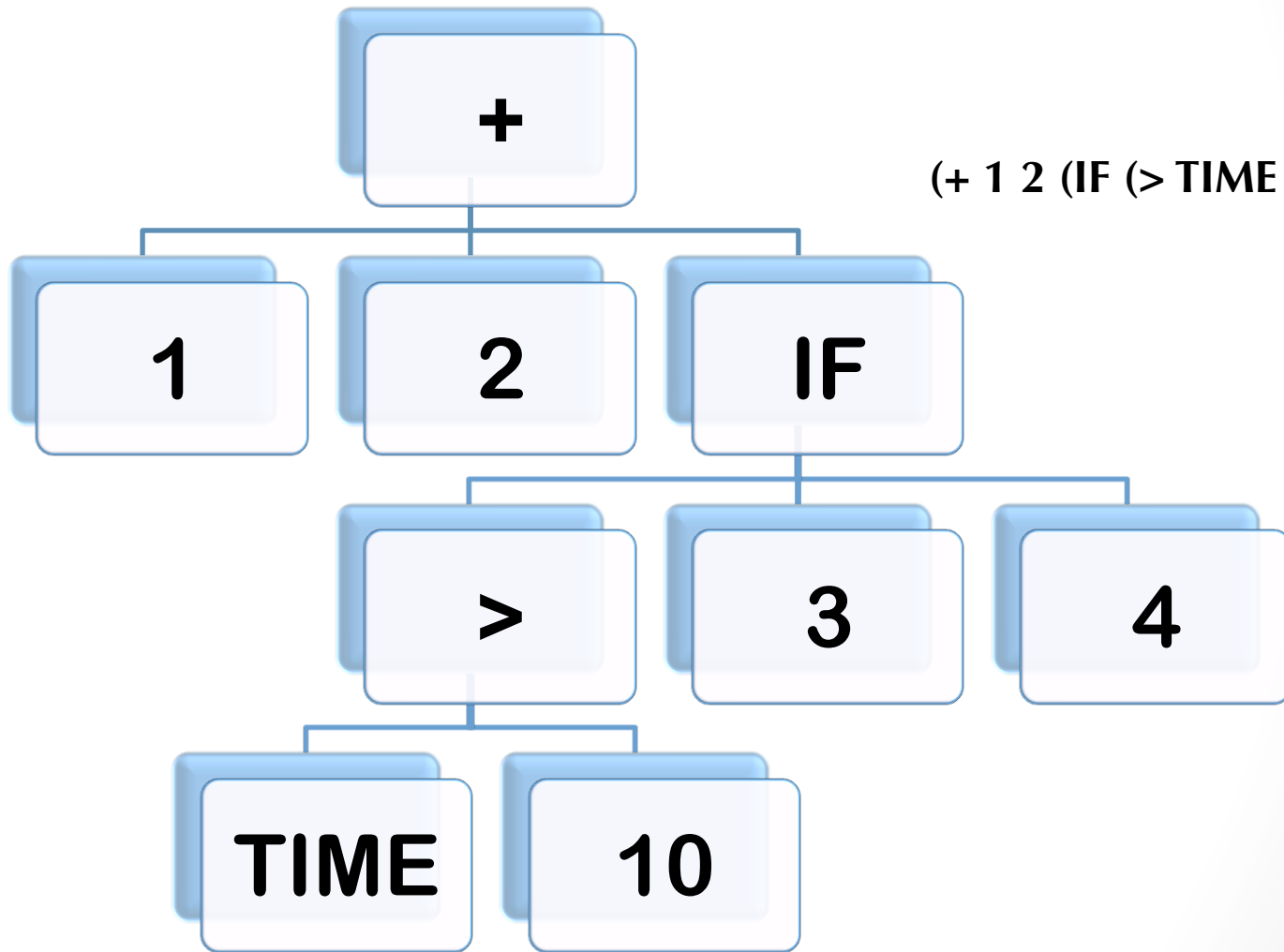
# GP'S Flow Diagram

# C program

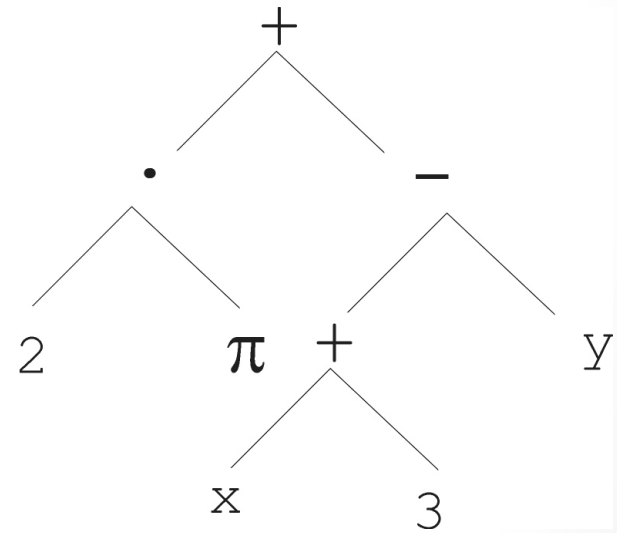
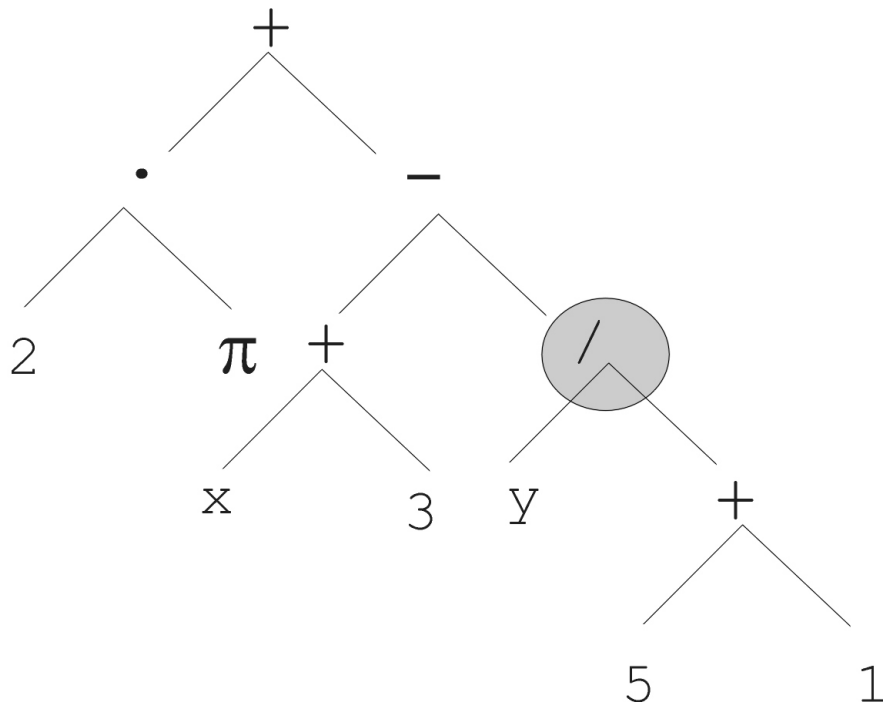
```
int foo (int time)
{
    int temp1, temp2;
    if (time > 10)
        temp1 = 3;
    else
        temp1 = 4;
    temp2 = temp1 + 1 + 2;
    return (temp2);
}
```

time	result
0	6
1	6
2	6
3	6
4	6
5	6
6	6
7	6
8	6
9	6
10	6
11	7
12	7

# Tree representation

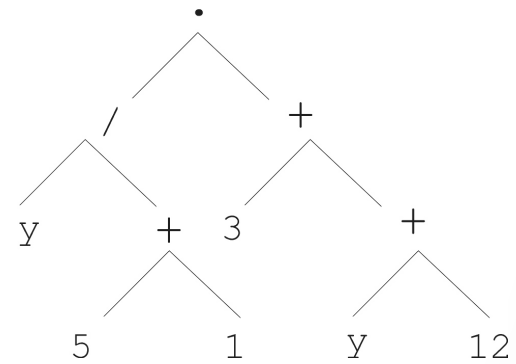
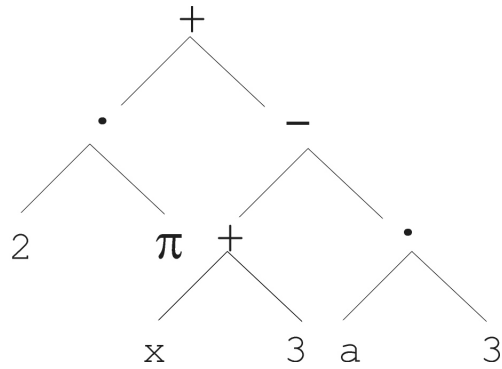
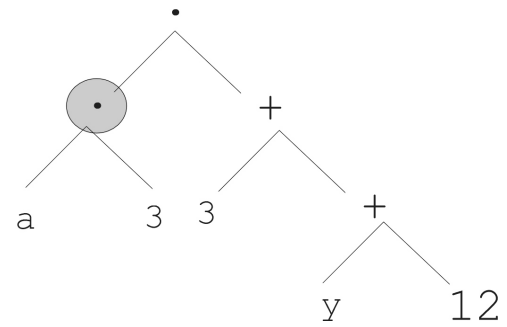
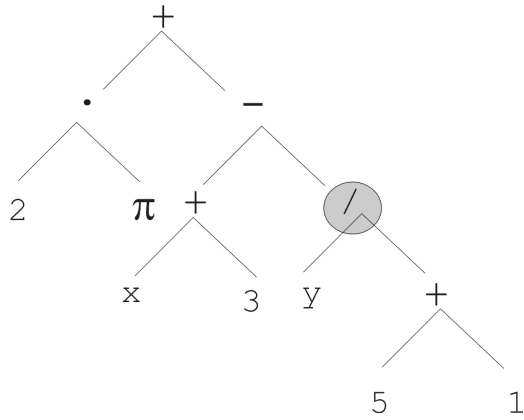


# Mutation example





# Crossover

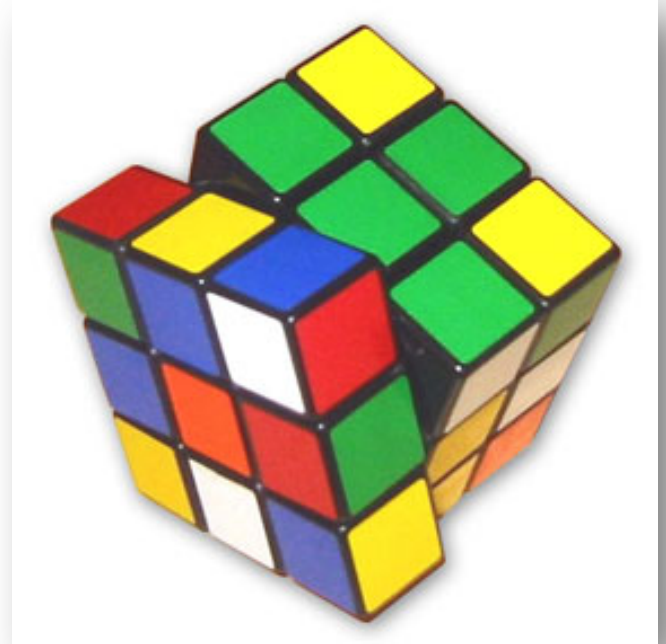


# Fitness

- How to measure the quality of a problem?
  - Number of errors, impact of the errors, computing time, computational complexity, etc.
- Bloating.

# Our experience with GP

- I directed an undergraduate thesis on GP.
- UC3M GECCO 2009 GP Rubik's cube team.
- We were the only participants!



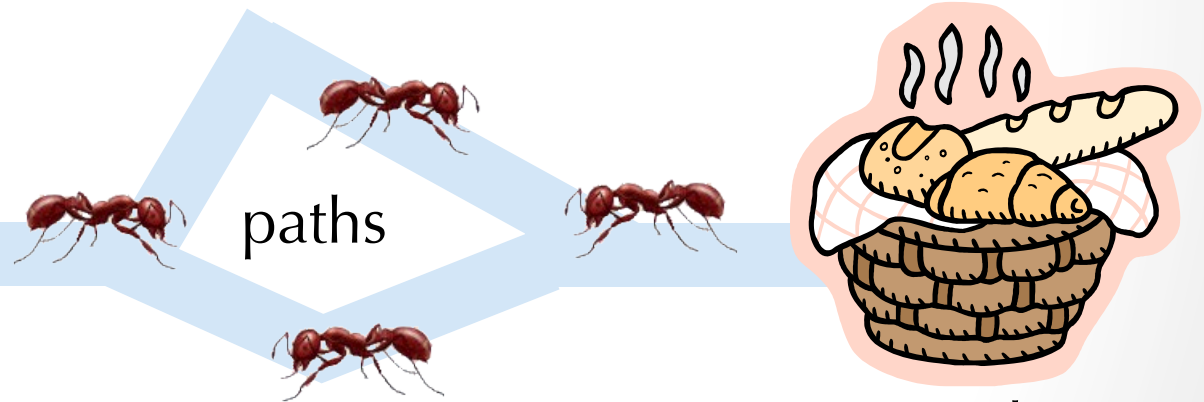
# Colonia de Hormigas

# More inspiration on nature

- Ant colony optimization.

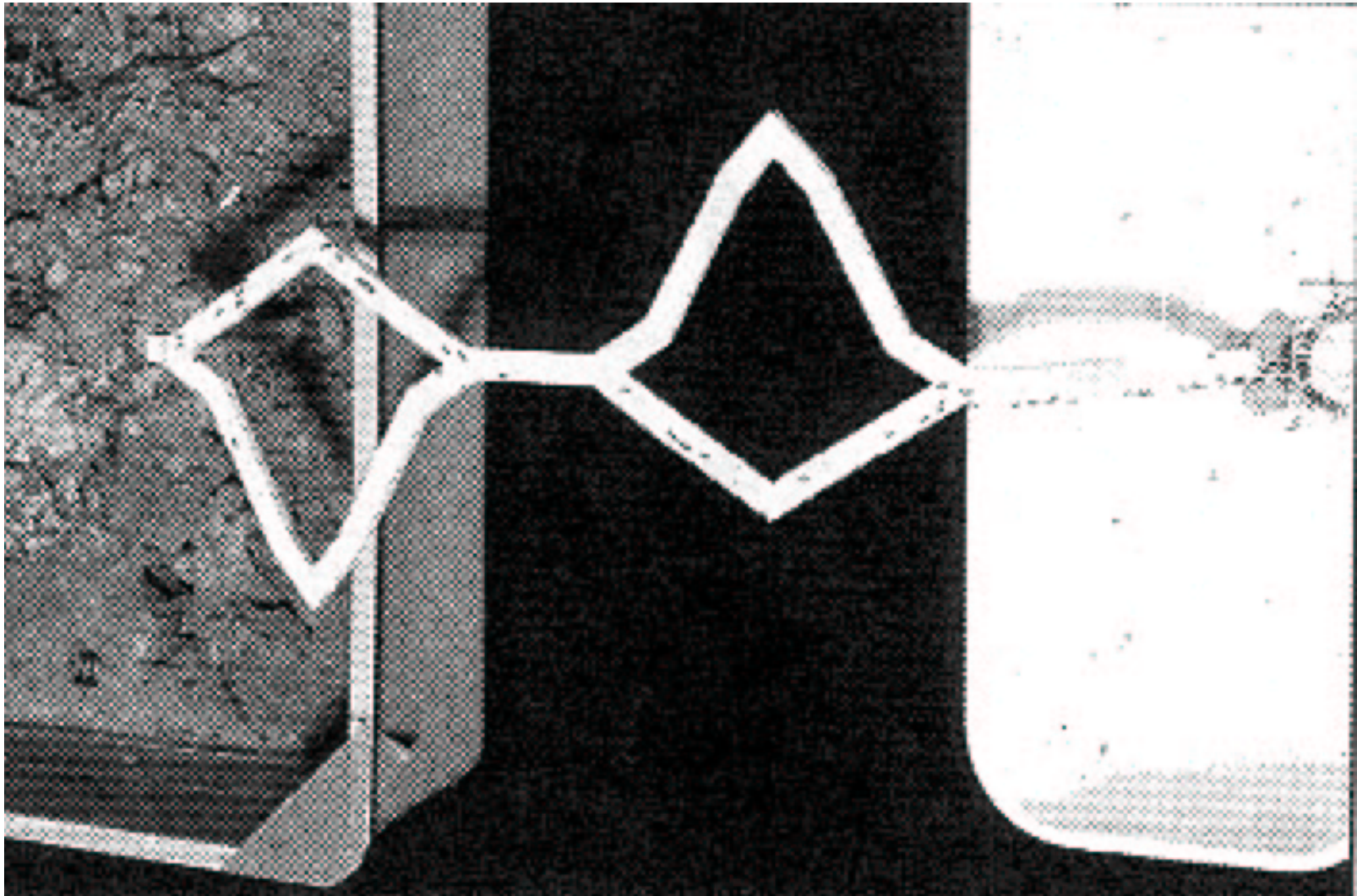


Nest



Food

# Ant colony in action



# More formally

- For a connected graph  $G=(N,A)$  the ant colony find the shortest path between two nodes.
- There is an “artificial pheromone footprint” associated with every arc in  $A$ .
- Ants can “read” and “write” that footprint.
- Highly transited arcs have a higher footprint.

# Final Remarks



# Final remarks

- Differential evolution.
- Estimation of distribution algorithms.
- Particle swarms.
- These approaches have seen many important practical results.
- Inspiration from nature does not stop here!

# Homework!

- Read:
  - Von Zuben, Fernando J. "Computação evolutiva: uma abordagem pragmática."
- Start getting familiarized with IPython, numpy, scipy, scikit.learn, inspyred and DEAP.