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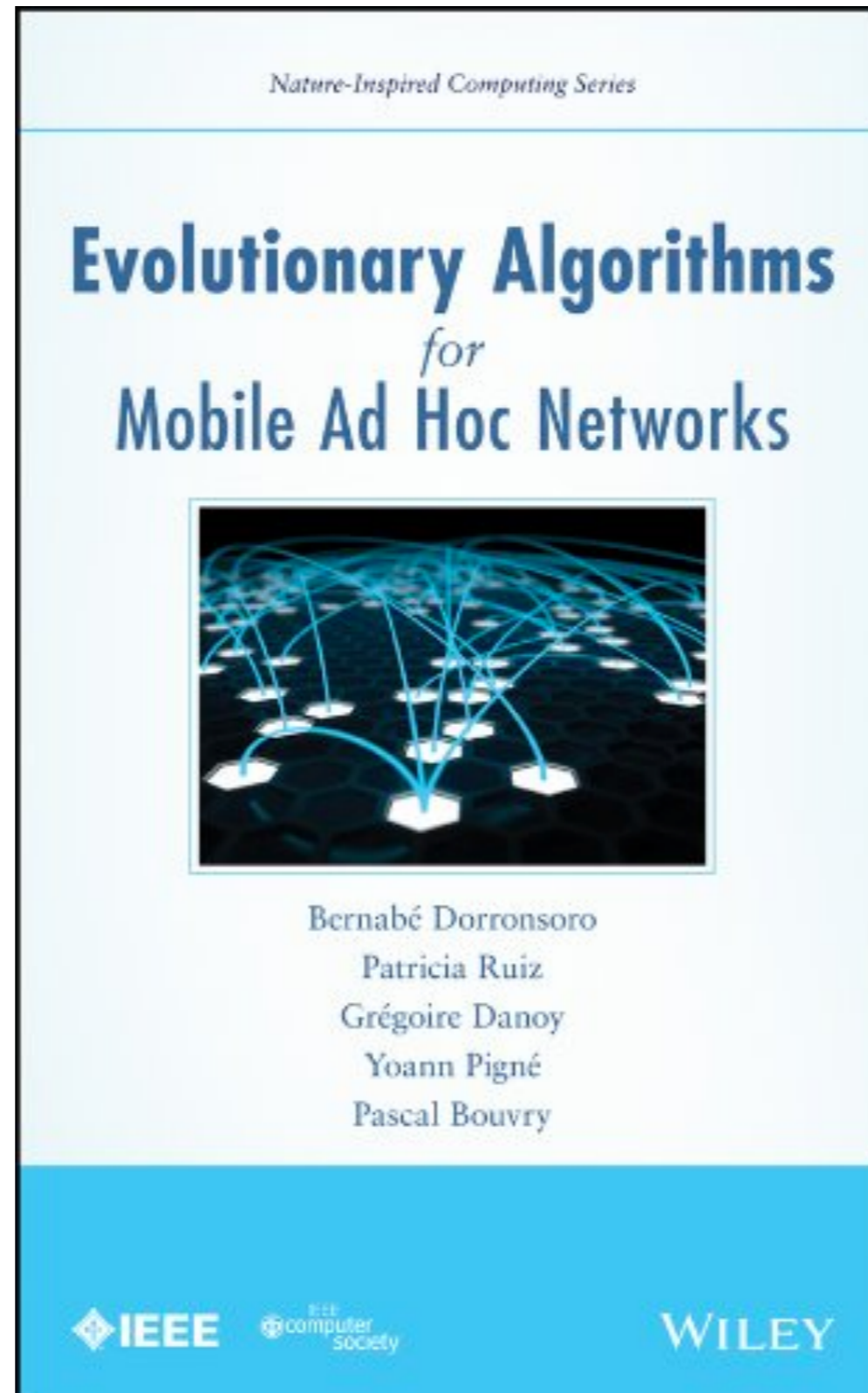
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# **An Optimization Framework for Mobile Ad Hoc Networks**

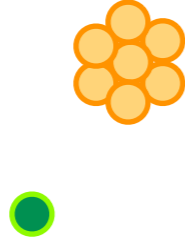
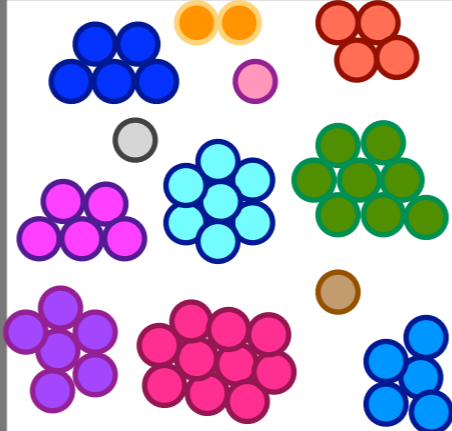
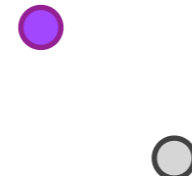
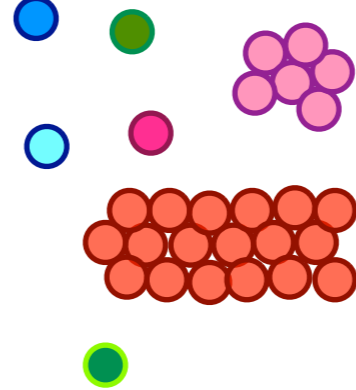
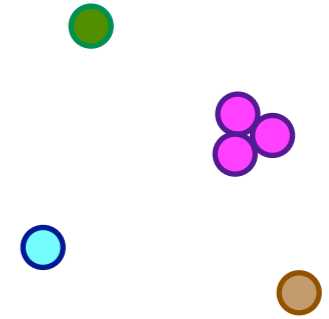













# **On the Need of Optimization for Mobile Ad Hoc Networks**



- Energy efficiency
- Broadcast
- Routing
- Network topology
  - Connectivity
  - Clustering
  - Node deployment
- Selfishness
- Security
- Quality of Service

- We can characterise them in terms of
  - Operation mode
    - ▶ Offline
    - ▶ Online
  - Knowledge
    - ▶ Global
    - ▶ Local
  - Approach
    - ▶ Centralized
    - ▶ Decentralized

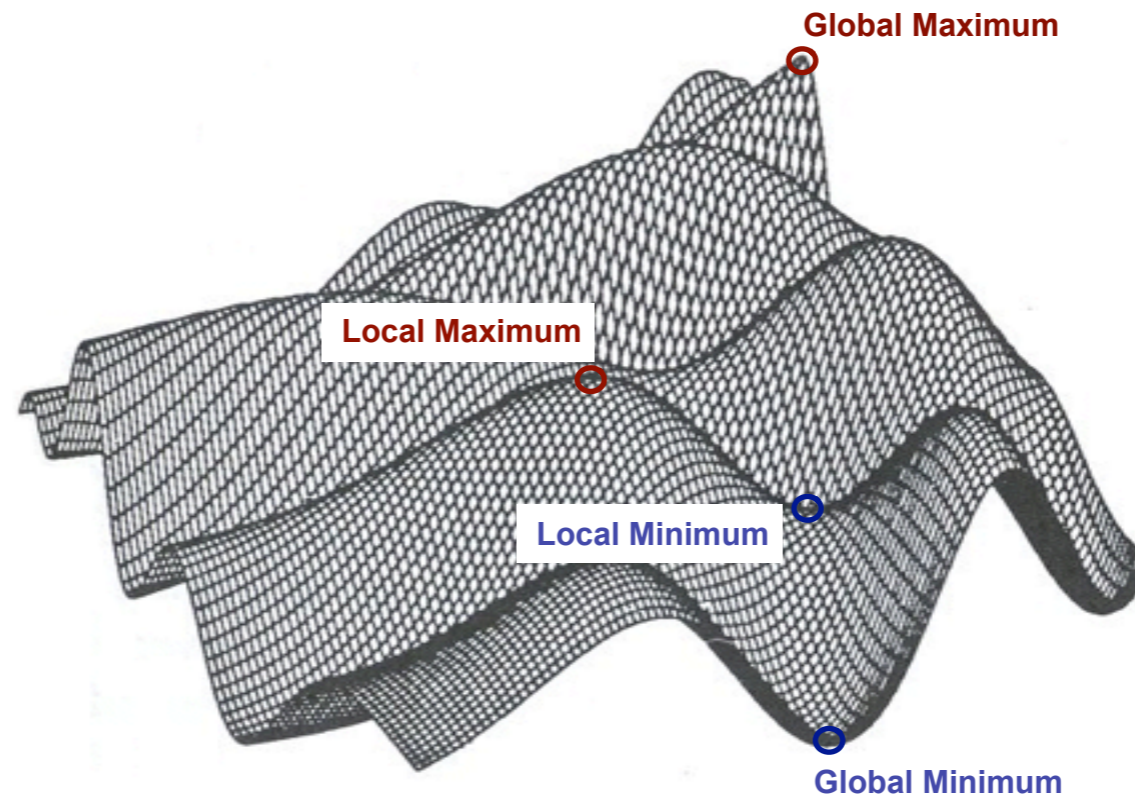
	Centralised local	Centralised global	Decentralised local	Decentralised global
Offline				
Online				

-  Protocol optimisation
-  Broadcasting
-  Mobility
-  Topology Ctrl: Sleep mode
-  Clustering
-  Selfishness
-  Topology Ctrl: Power allocation
-  Routing
-  Security
-  Topology Ctrl: Node deployment
-  Multipath Routing
-  Others
-  Topology Ctrl: Connectivity
-  Multicast Routing

# Single-objective Optimization

**Solutions Space**

$$Opt(f(\vec{x})) = \{\vec{x}^* | \forall \vec{x} \in M : \begin{cases} f(\vec{x}) \leq f(\vec{x}^*) & \text{Maximization} \\ f(\vec{x}) \geq f(\vec{x}^*) & \text{Minimization} \end{cases}\}$$





- Complete methods

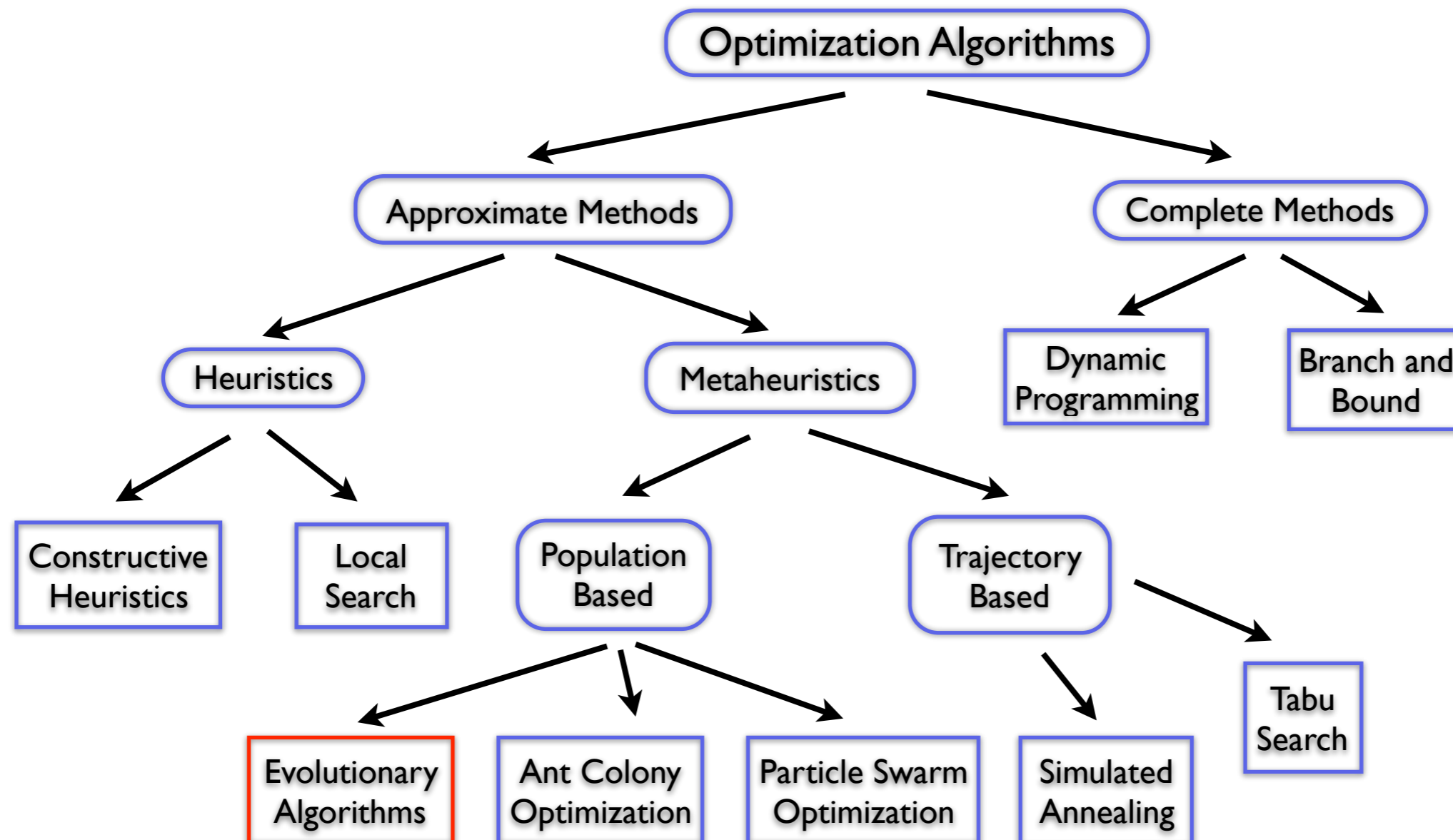
They guarantee to find for every finite size instance of a CO problem an optimal solution in bounded time

Only CO problems!

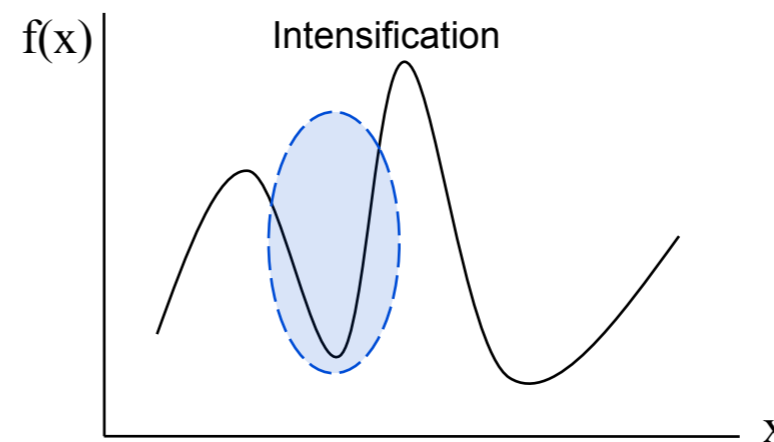
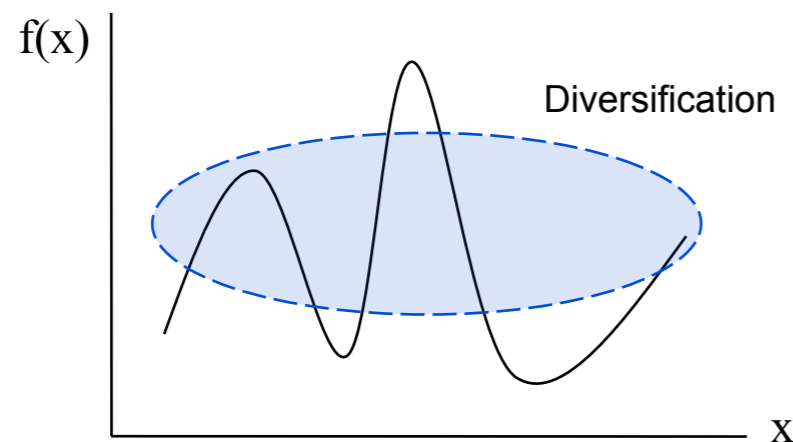
- Approximate methods

No guarantee of finding an optimal solution

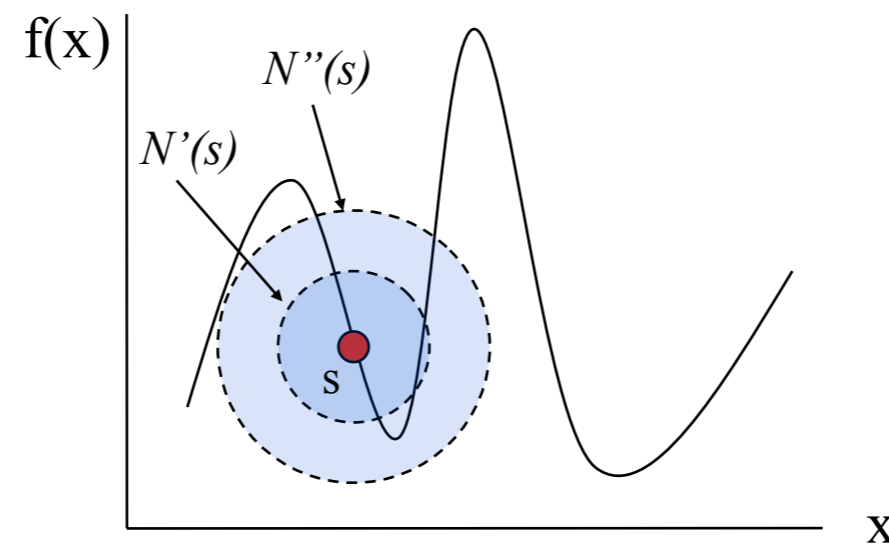
Combinatorial and Continuous



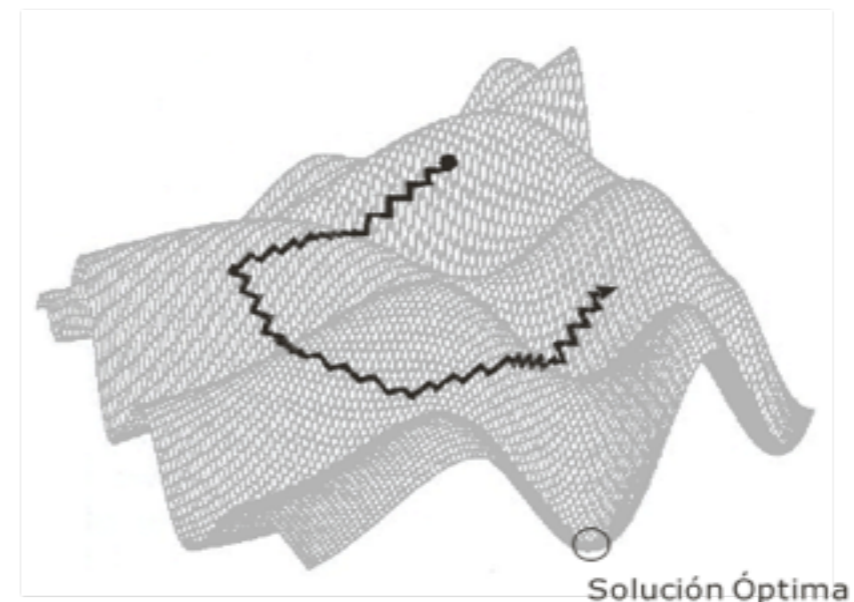
- Metaheuristics must achieve a balance between diversification and intensification
  - Diversification: exploration of the search space
  - Intensification: exploitation of promising regions of the search space



- Given a solution  $s$ , the neighborhood of  $s$ ,  $N(s)$ , is the set of solutions of the search space that can be reached using some kind of transformation on  $s$



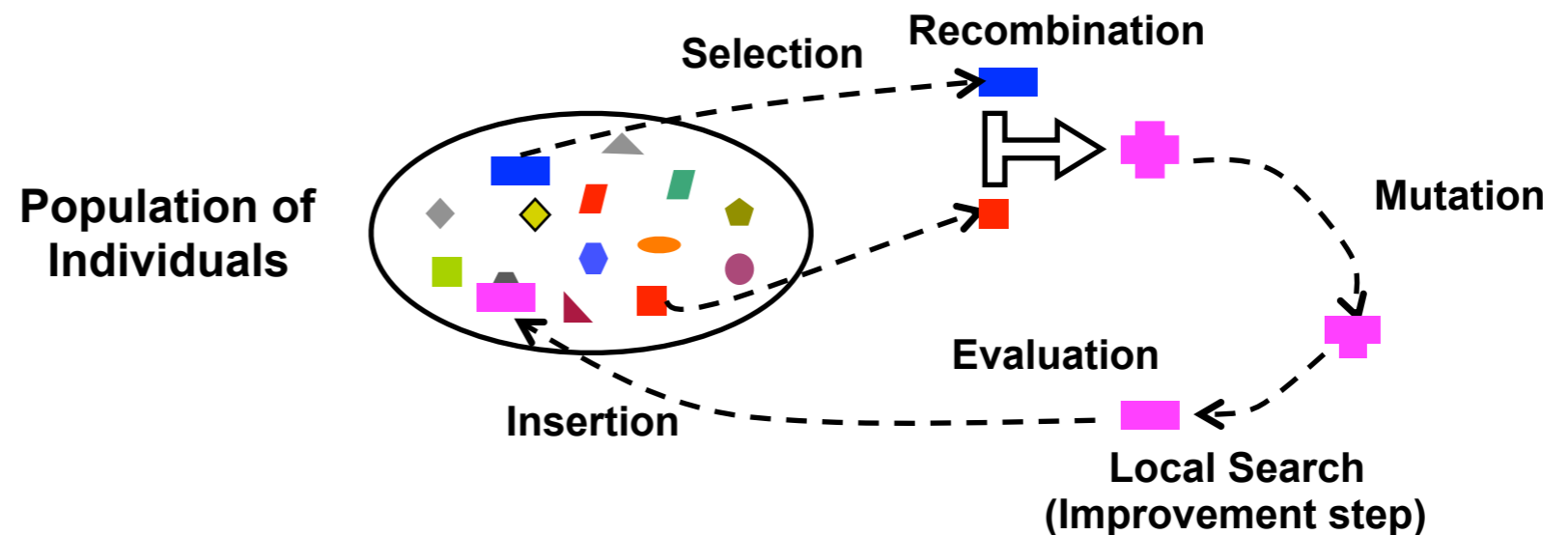
- Evolutionary Algorithms (EAs): Useful optimization techniques for complex problems
  - Show a good tradeoff between exploration and exploitation
- Based in population
  - Individuals → Potential solutions to the problem
    - Fitness value: ¿How good is the individual?
  - Variation operators → Allow the evolution of the population
    - Recombination: Interchange of genetic material
    - Mutation: Generation of new genetic material



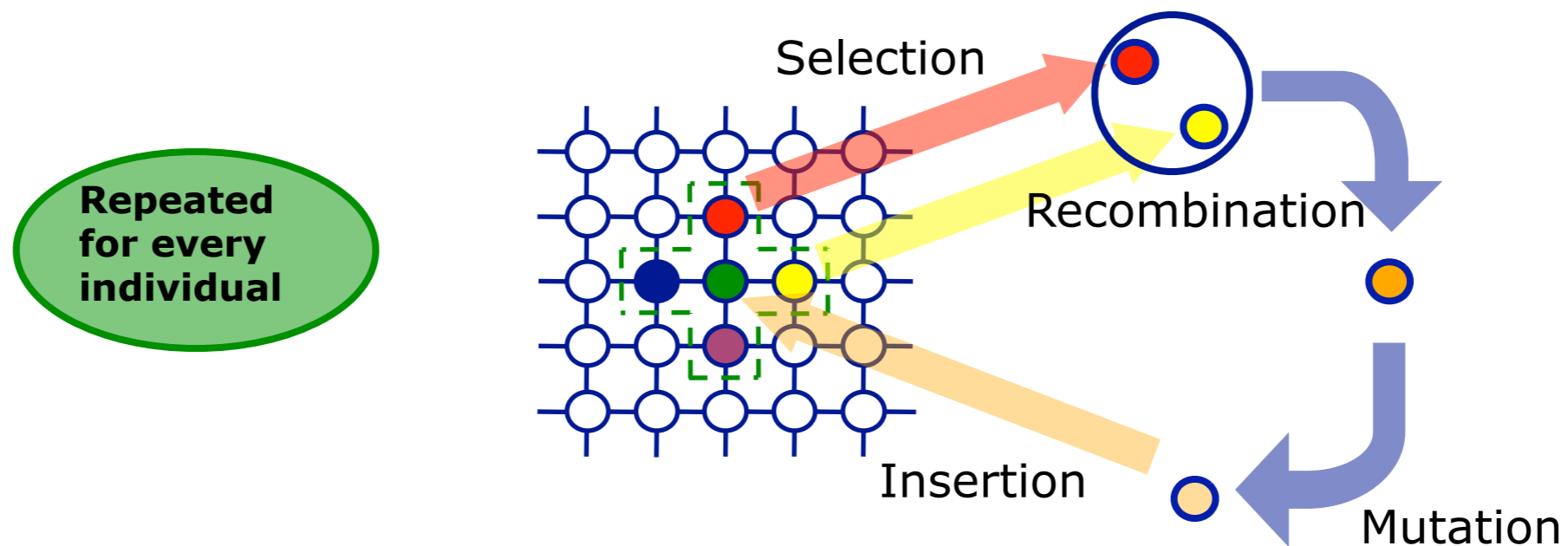
- Population evolution
  - Improvement of the quality of solutions
  - Guided by the fitness function

Panmictic

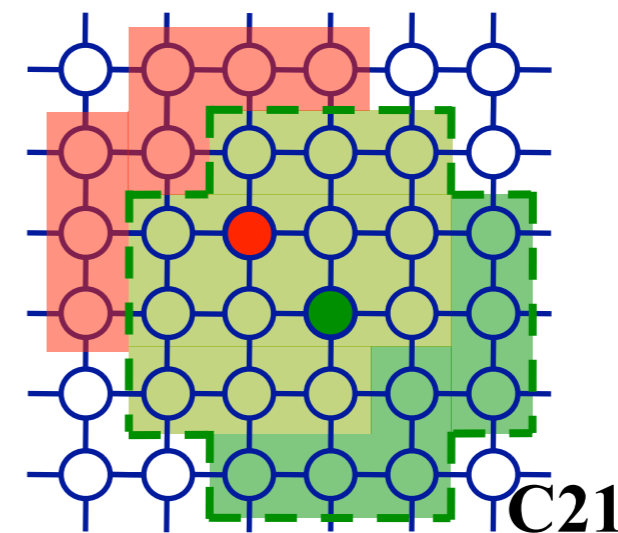
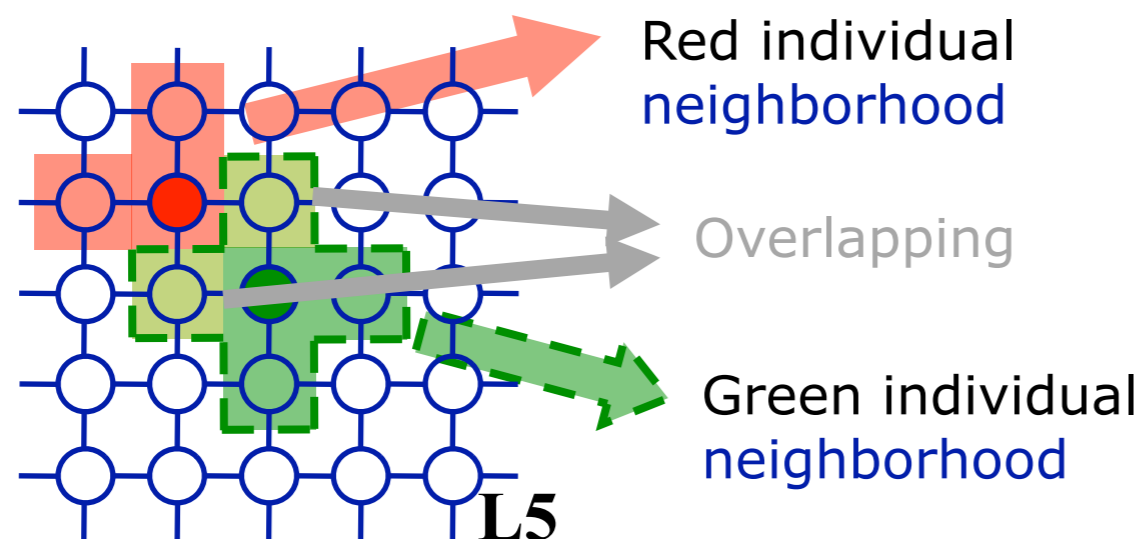
- Application operators
  - Stochastic
  - Generic



- Spatially structured population (2-D)
- Breeding loop applied inside small neighborhoods



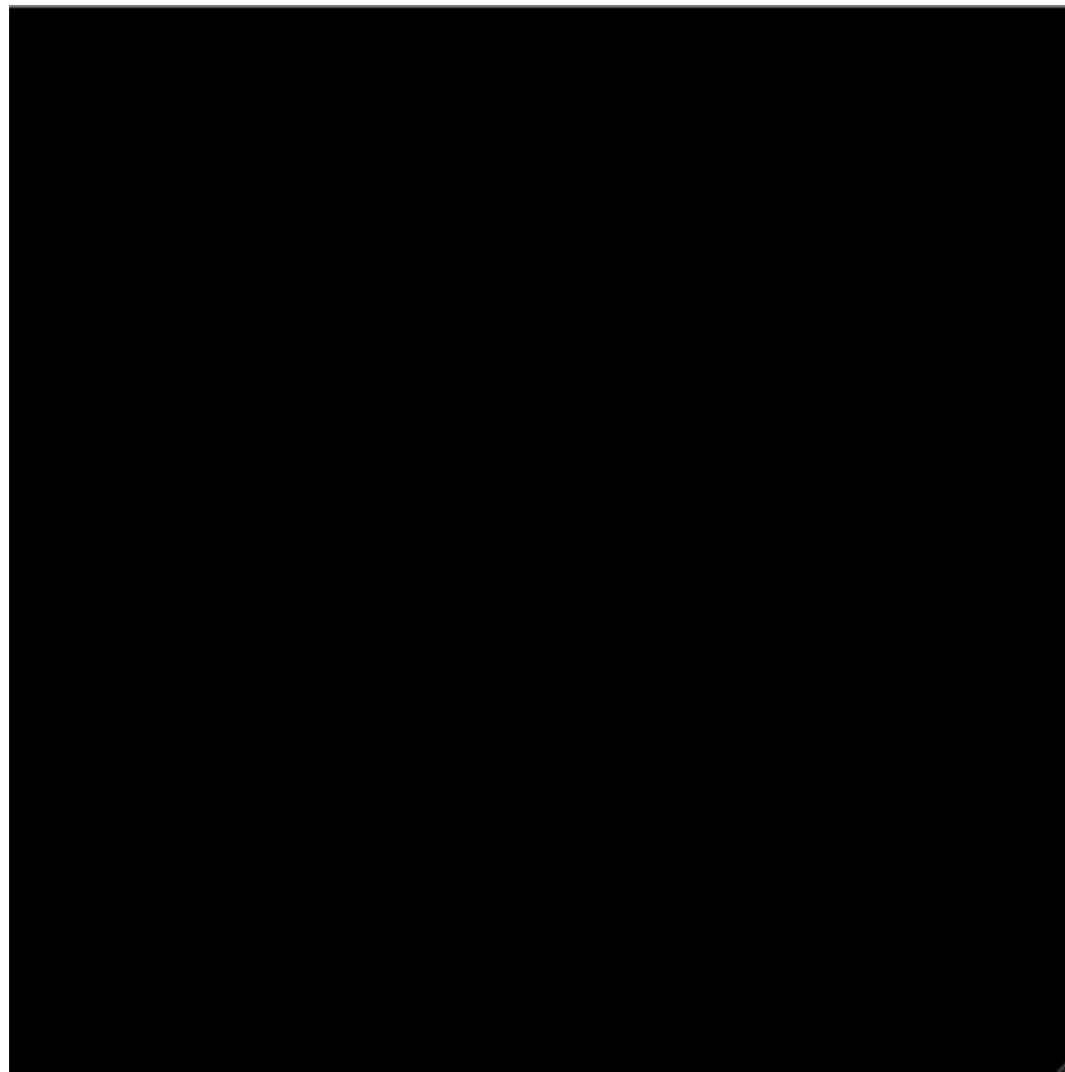
- Spatially structured population (2-D)
- Breeding loop applied inside small neighborhoods
- Overlapped neighborhoods → Smooth diffusion
- Isolation by distance among individuals in the population
- Appropriate exploration/exploitation tradeoff
  - Exploitation: Inside neighborhoods
  - Exploration: Neighborhood borders





## MAXUT100 Problem

cGA with L5

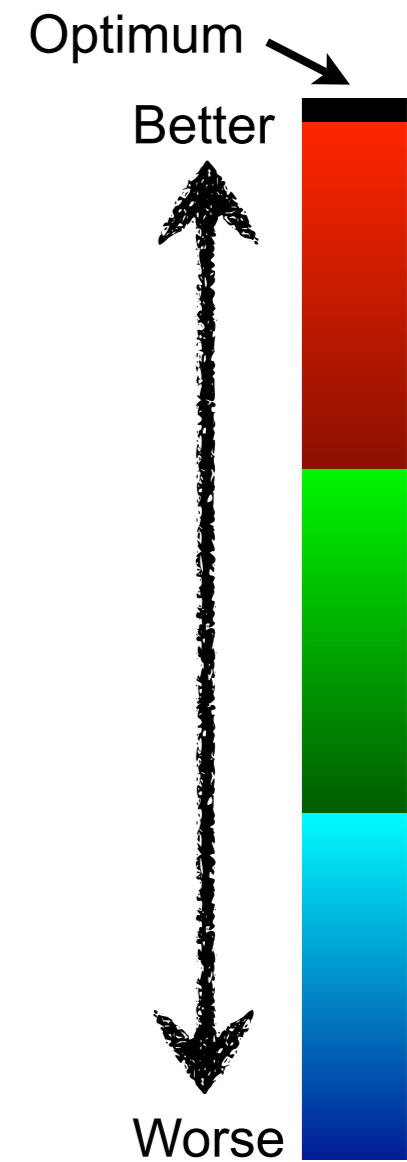


Optimum (1077.0) after 33 s

genGA



Converges to 967.0 after 24s



## MAXUT100 Problem

cGA with L5

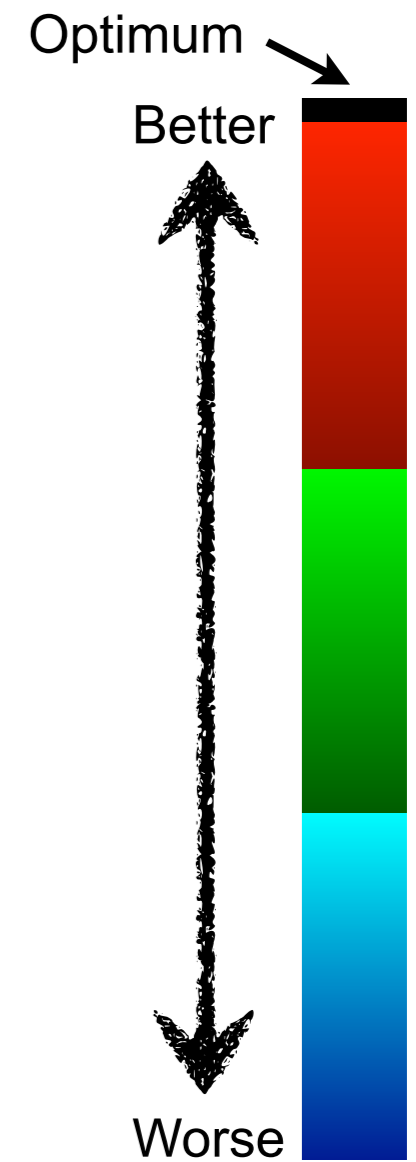


Optimum (1077.0) after 33 s

genGA



Converges to 967.0 after 24s



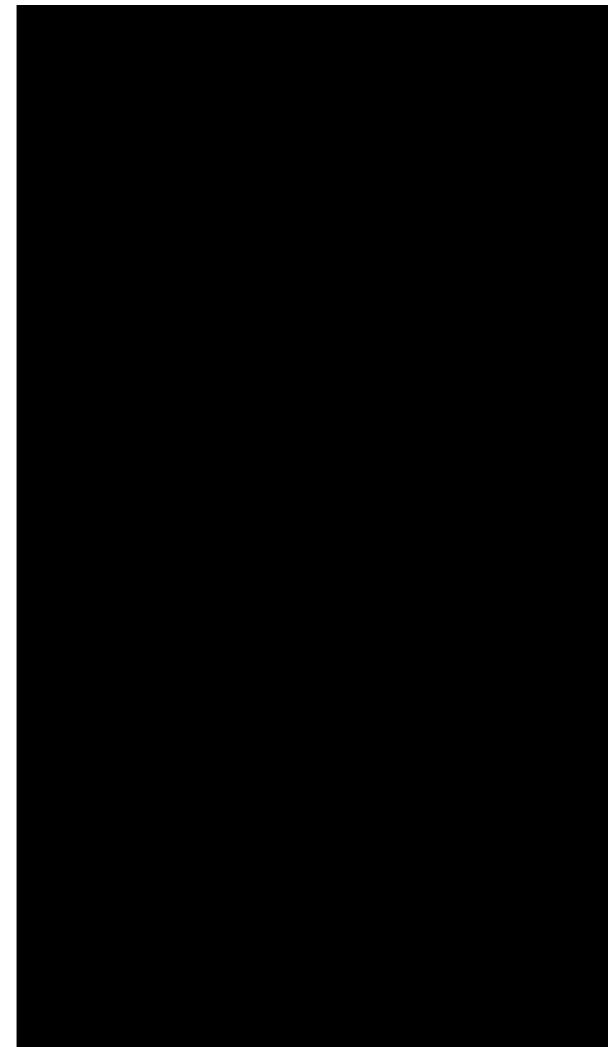
## MAXUT20\_01 Problem

cGA with L5

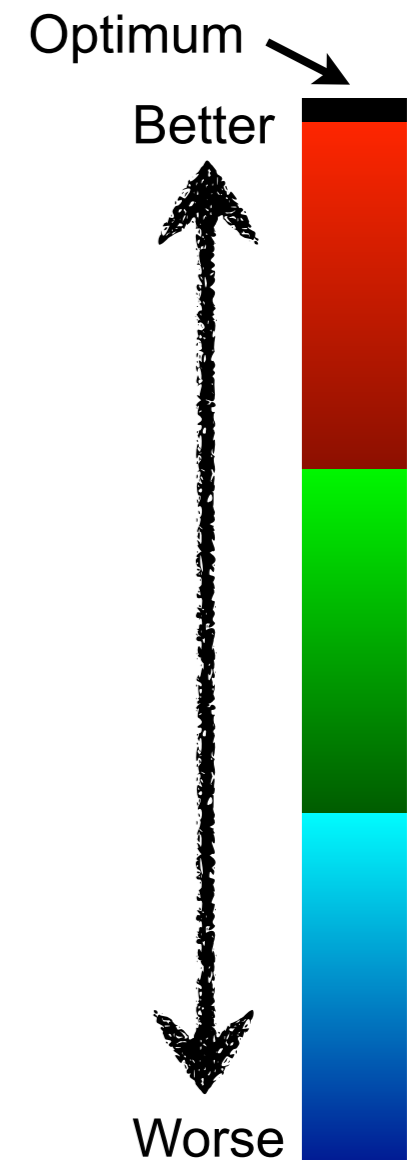


Optimum (10.1198) after 1.9 s

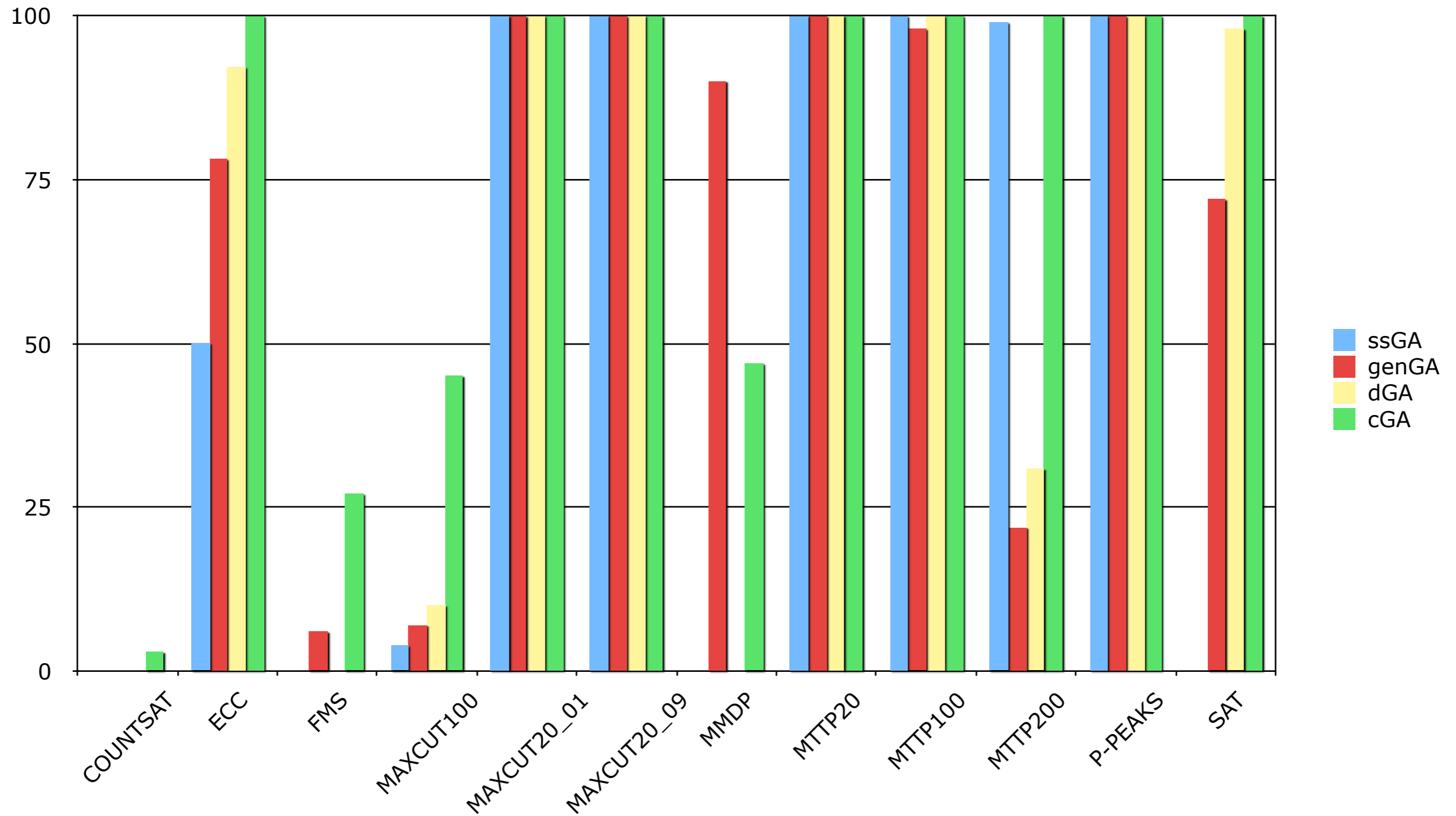
genGA

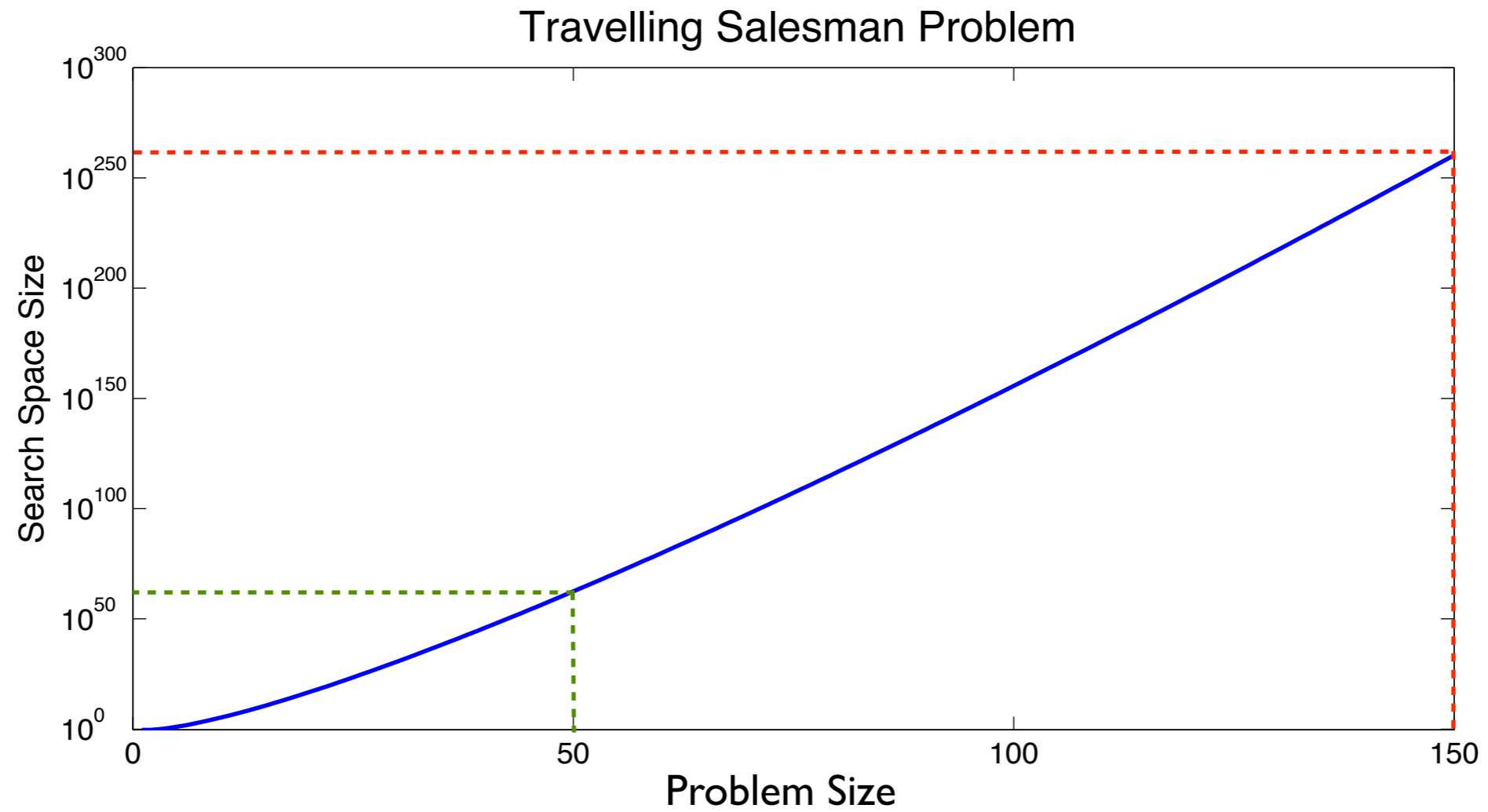


Optimum after 18.5s



### Percentage of Successful Runs



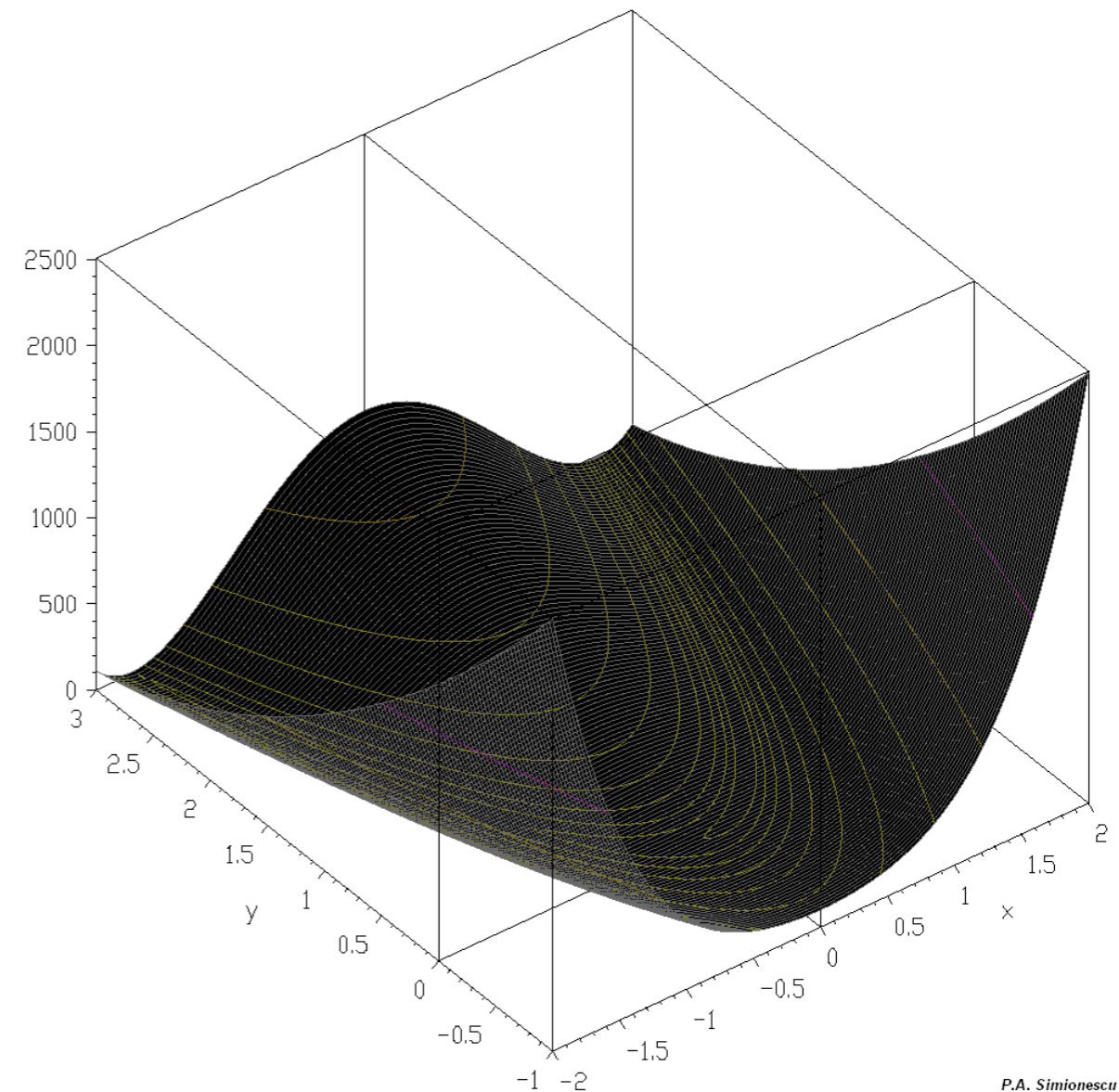


- Part of De Jong's five function test suite
- Continuous and unimodal

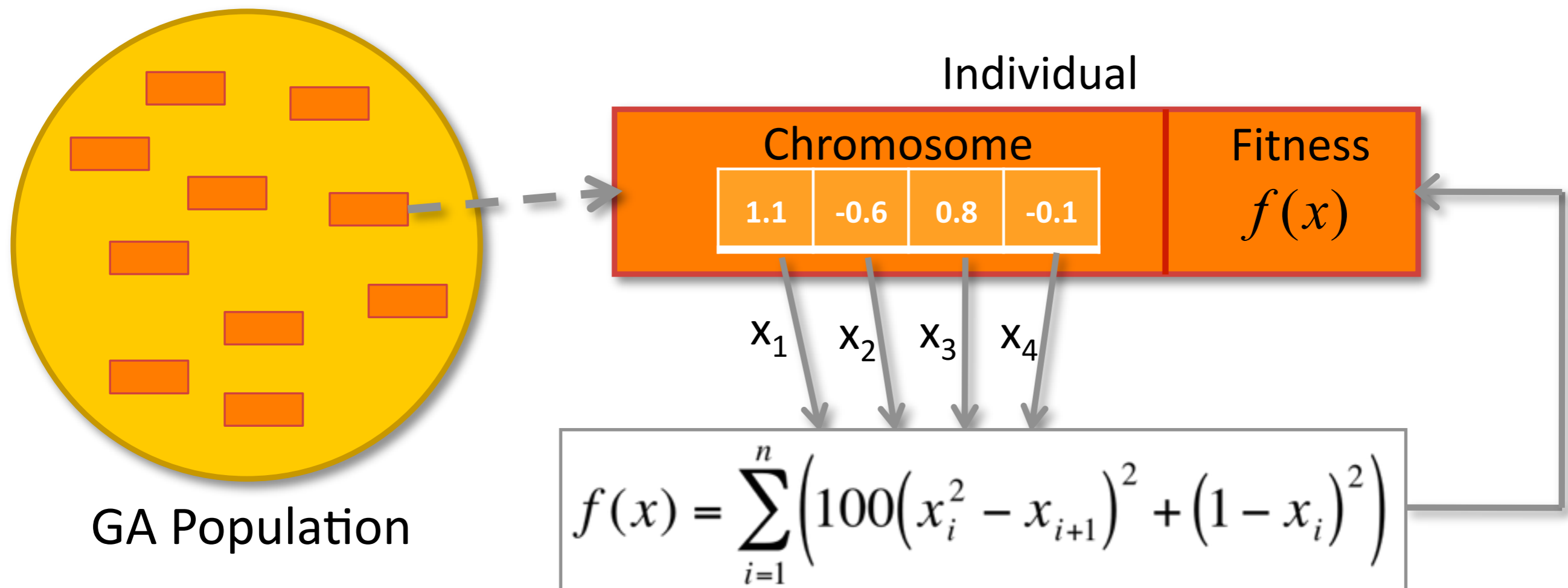
$$f(x) = \sum_{i=1}^n \left( 100(x_i^2 - x_{i+1})^2 + (1 - x_i)^2 \right)$$

with  $-2.12 \leq x_i \leq 2.12$

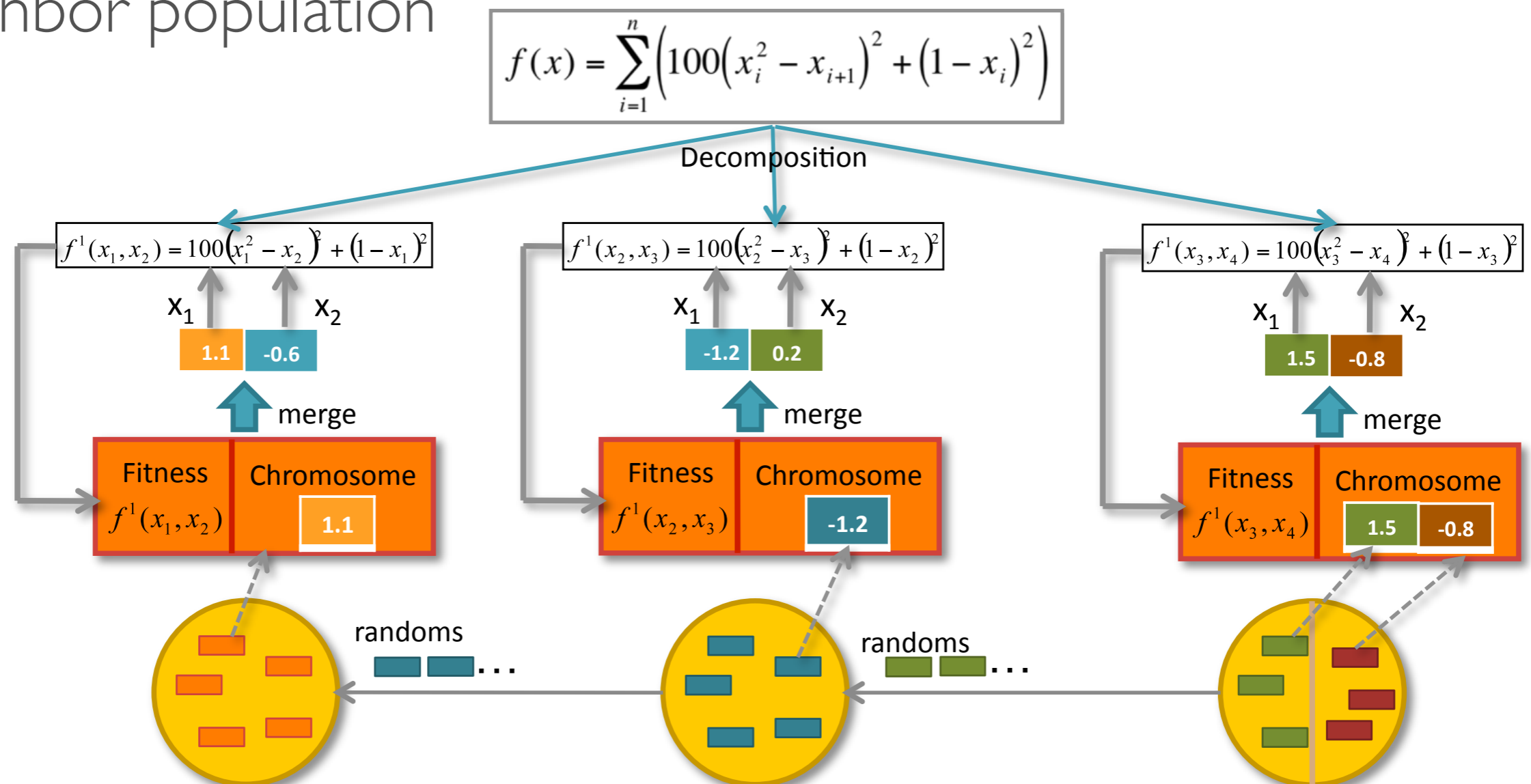
- Global minimum  $f(x^*) = 0$   
with  $x^* = (1, 1, \dots, 1)$



- A chromosome encodes a complete solution
- Solution evaluated on the global problem

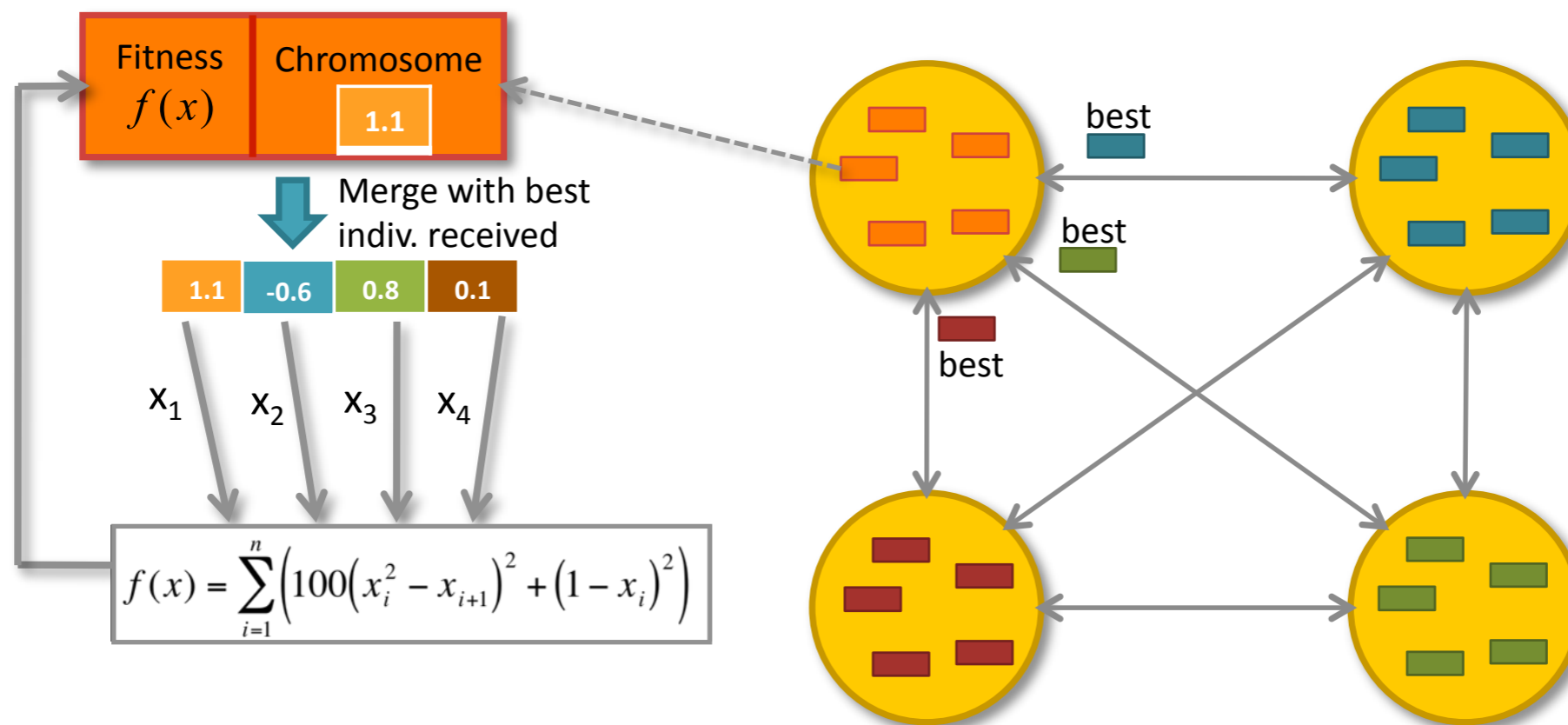


- Each node runs a subpopulation for a **subset of the N variables**
- Each population **evaluates** its individuals on a **local subproblem** using a random individual received from its neighbor population





- Each node runs a subpopulation for a **subset of the N variables**
- Each population **evaluates** each of its individuals **on the global fitness function** using the best individual received from each other subpopulation





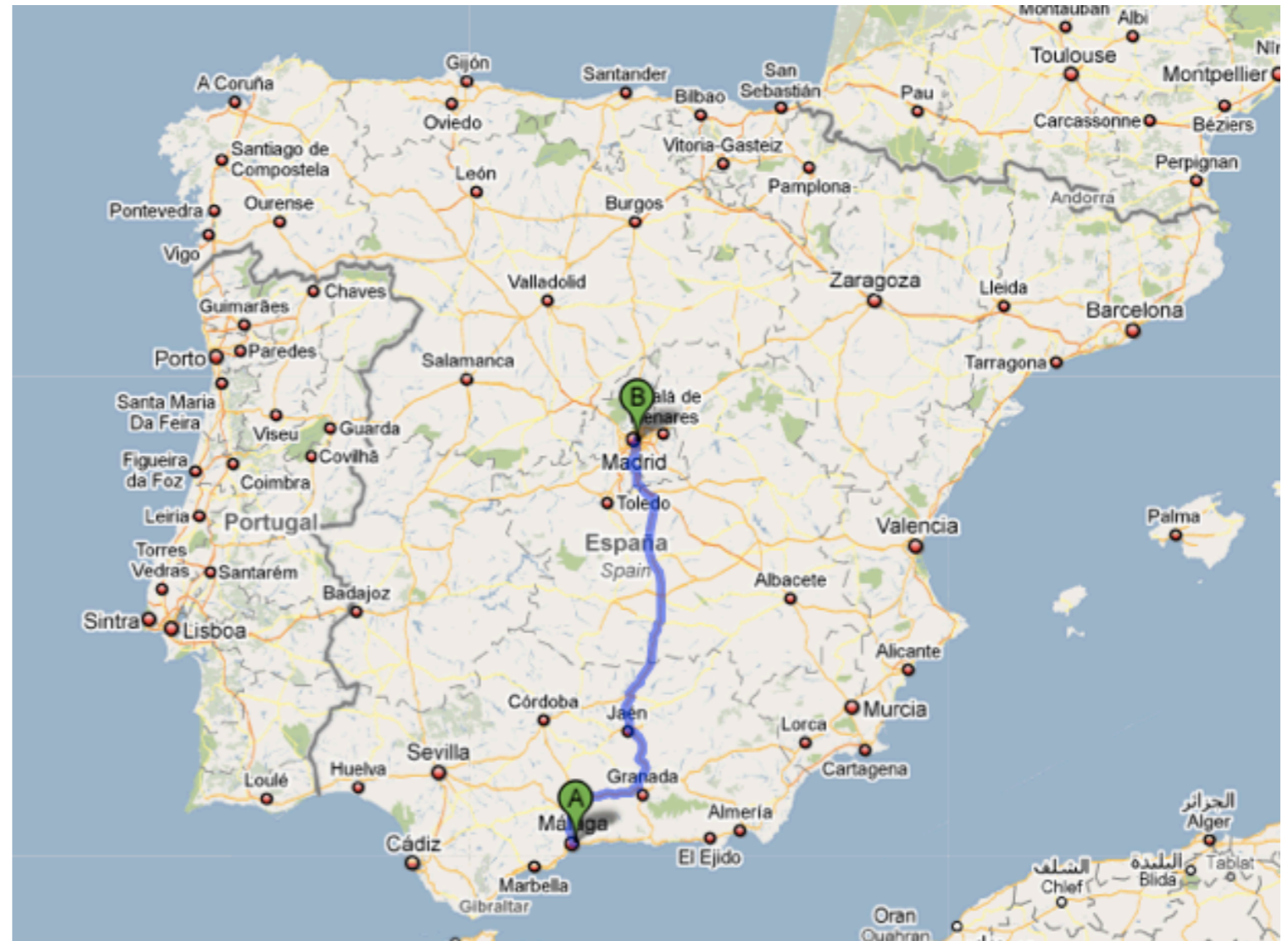
# **Multi-objective Optimization**

- Many real-world optimization problems require to optimize more than one objective at the same time
  - These objectives are usually **in conflict** among them
  - Improving one means worsening the others
- **Multi-objective** (or multi-criteria) optimization
  - Discipline focused on solving multiobjective optimization problems ( MOPs )

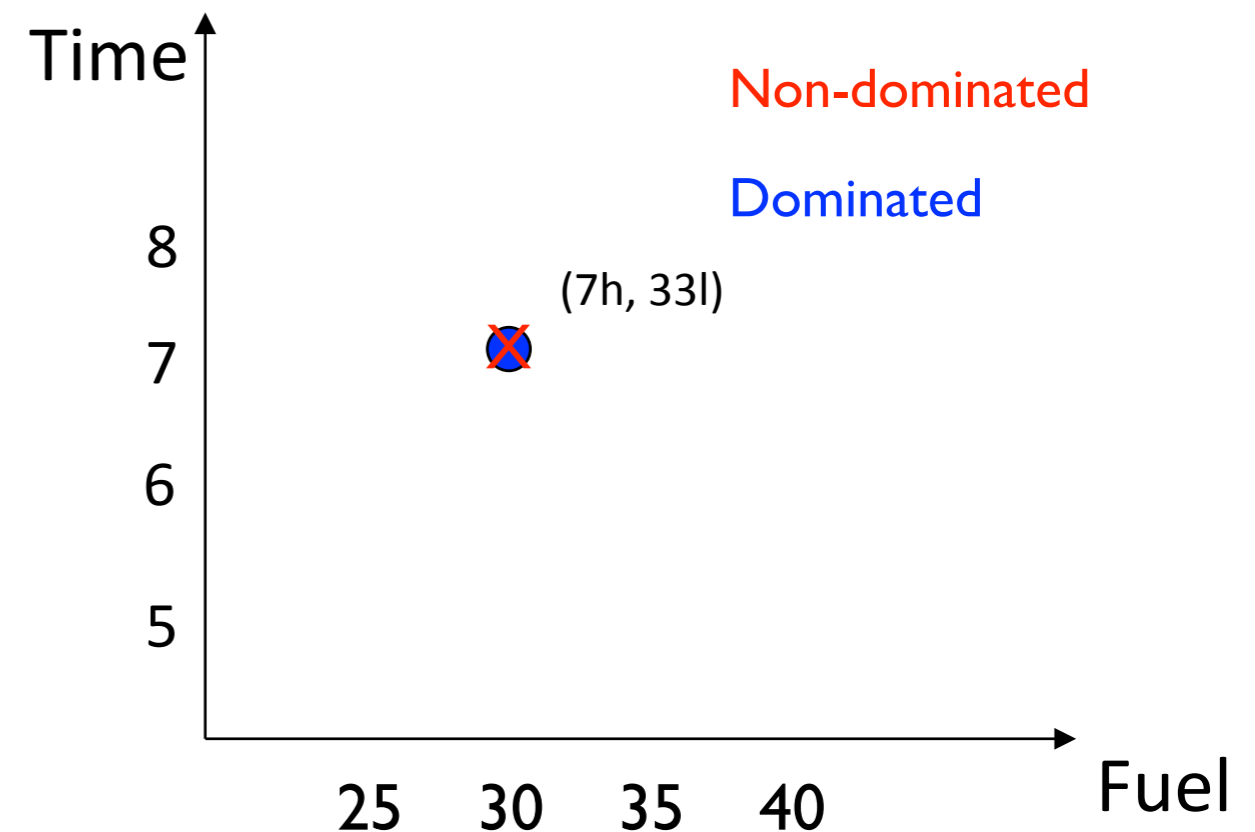
$$\begin{array}{lll} \text{Optimize} & f_m(\vec{x}) & m = 1, 2, \dots, m \\ \text{Subject to} & g_j(\vec{x}) \geq 0 & j = 1, 2, \dots, j \\ & h_k(\vec{x}) = 0 & k = 1, 2, \dots, k \end{array}$$

- Example: travelling by car from Málaga to Madrid (535 km)

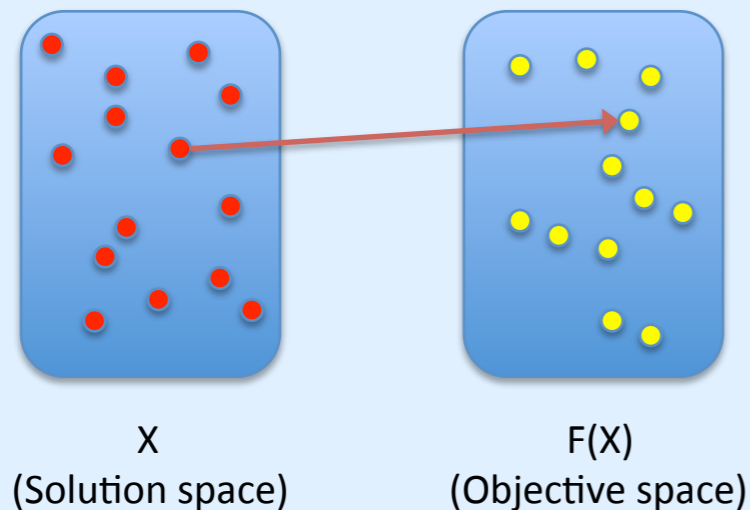
- Objective 1:
  - ▶ Minimizing time
- Objective 2:
  - ▶ Minimizing fuel
- Constraints:
  - ▶ Max. speed: 120 km/h
  - ▶ Min. speed: 60 km/h
- Decision variable:
  - ▶ mean car speed



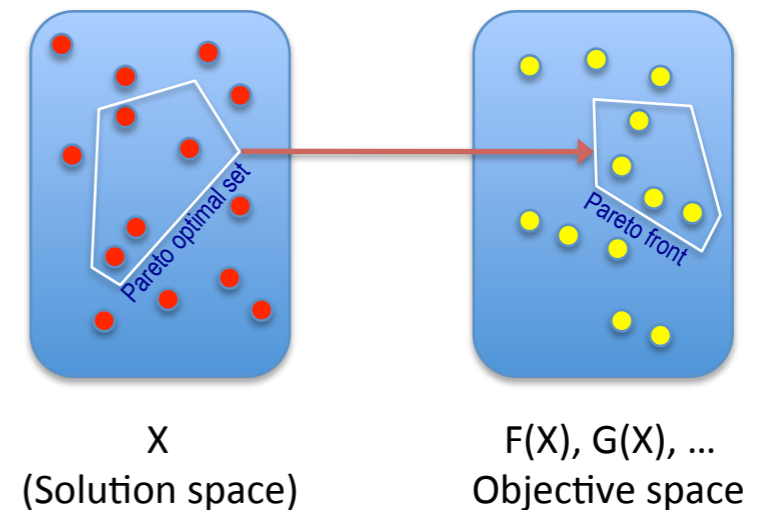
- Travelling by car from Málaga to Madrid (535 km)
  - Extreme solutions
    - ▶ Time: 5 hours, fuel: 9.0 litres
    - ▶ Time: 8 hours, fuel: 6.0 litres
  - Other solutions
    - ▶ Time: 5.5 hours, fuel: 7.5 litres
    - ▶ Time: 6 hours, fuel: 6.5 litres



- In single-objective optimization (SO)
  - The optimum is
    - ▶ One solution
    - ▶ Several ones with same quality



- In multi-objective optimization (MO)
  - The optimum (Pareto optimal set) is a set of (non-dominated) solutions



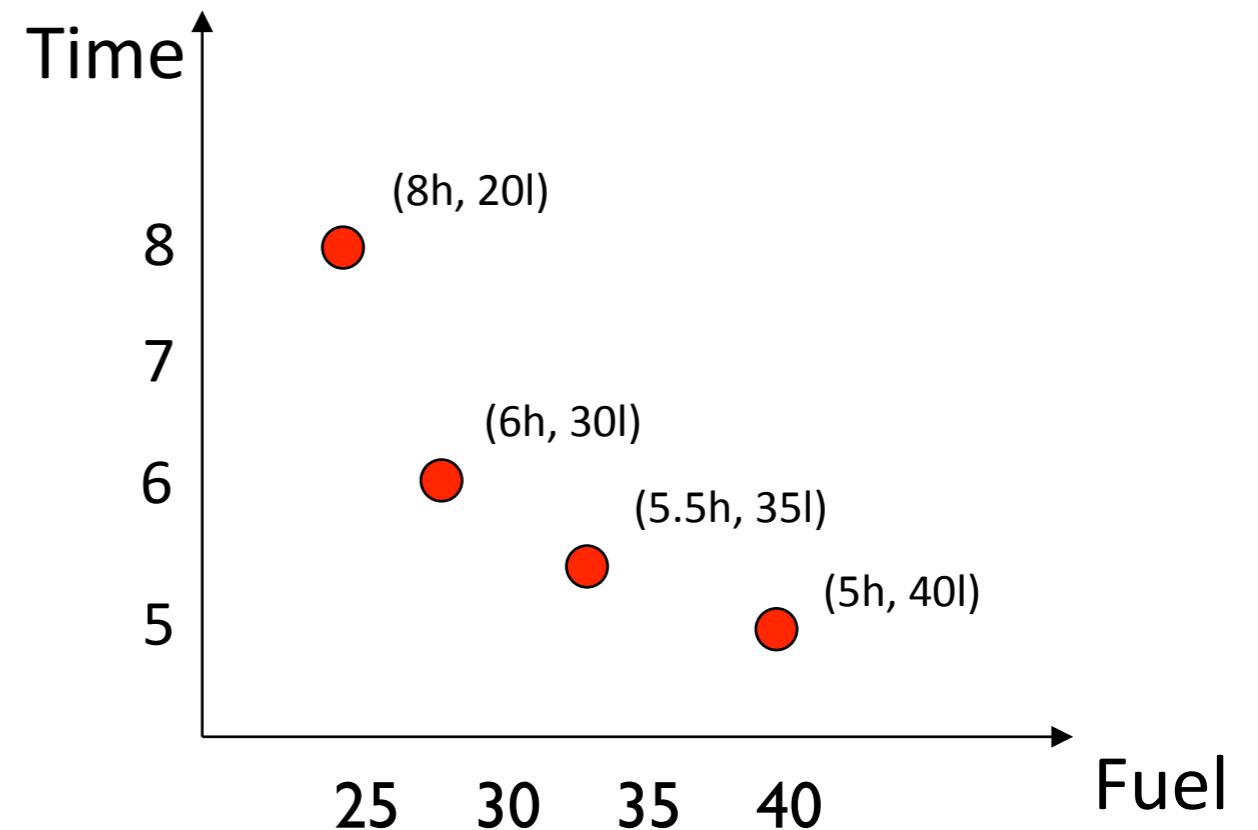
- Finding the Pareto front of a problem is not the last step in multi-objective optimization



- In practice, an expert in the domain (the decision maker) has to choose the best trade-off solution



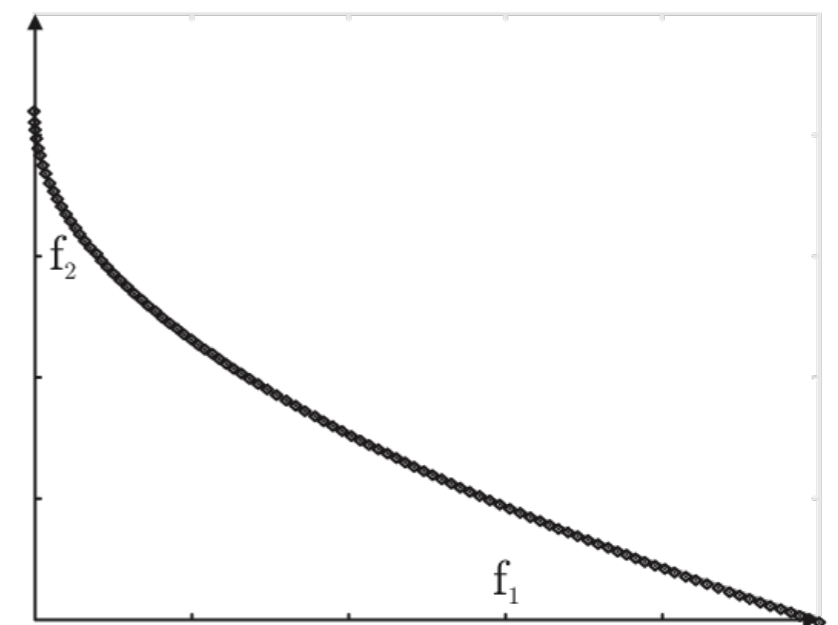
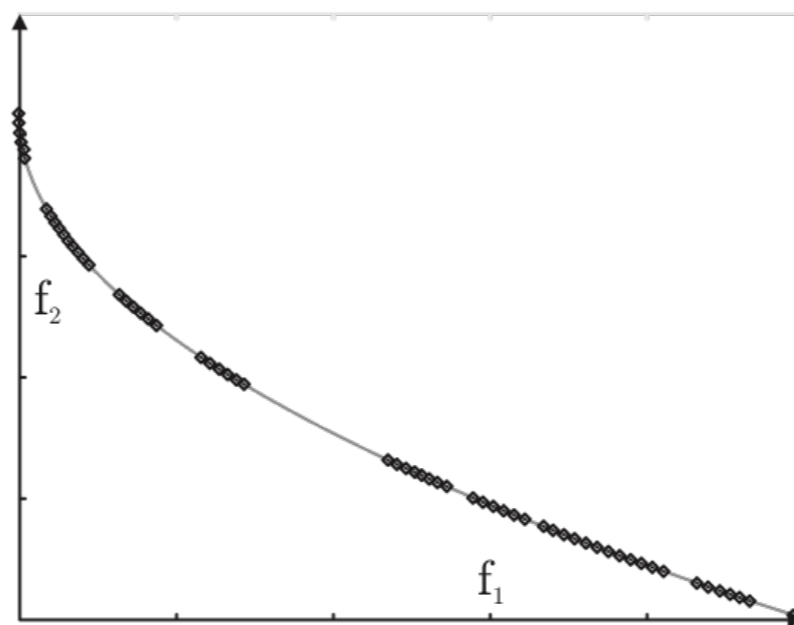
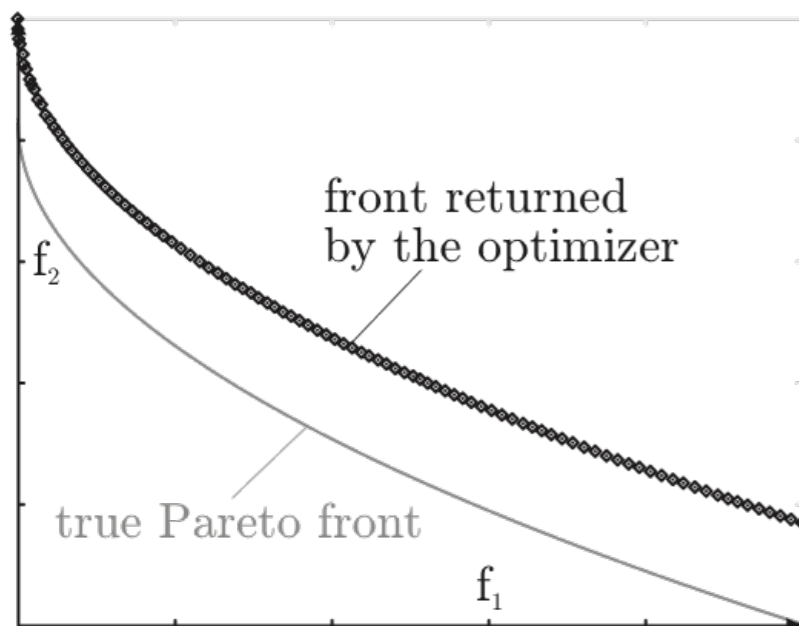
- In the example of traveling from Málaga to Madrid
- If time is important
  - Choose (5h, 40l)
- If consumption is important:
  - Choose (8h, 20l)
- Compromise solution:
  - (6h, 30l)
  - (5.5h, 35l)



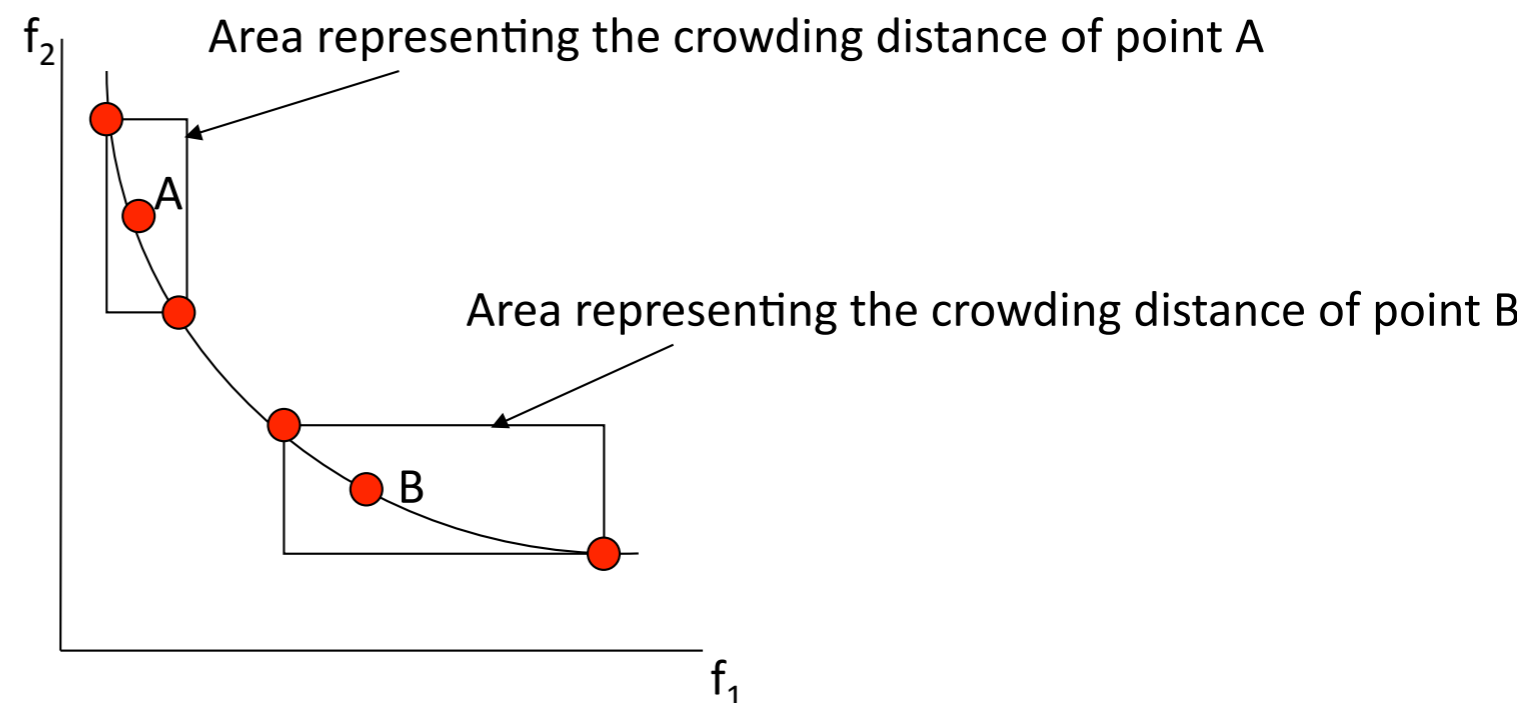
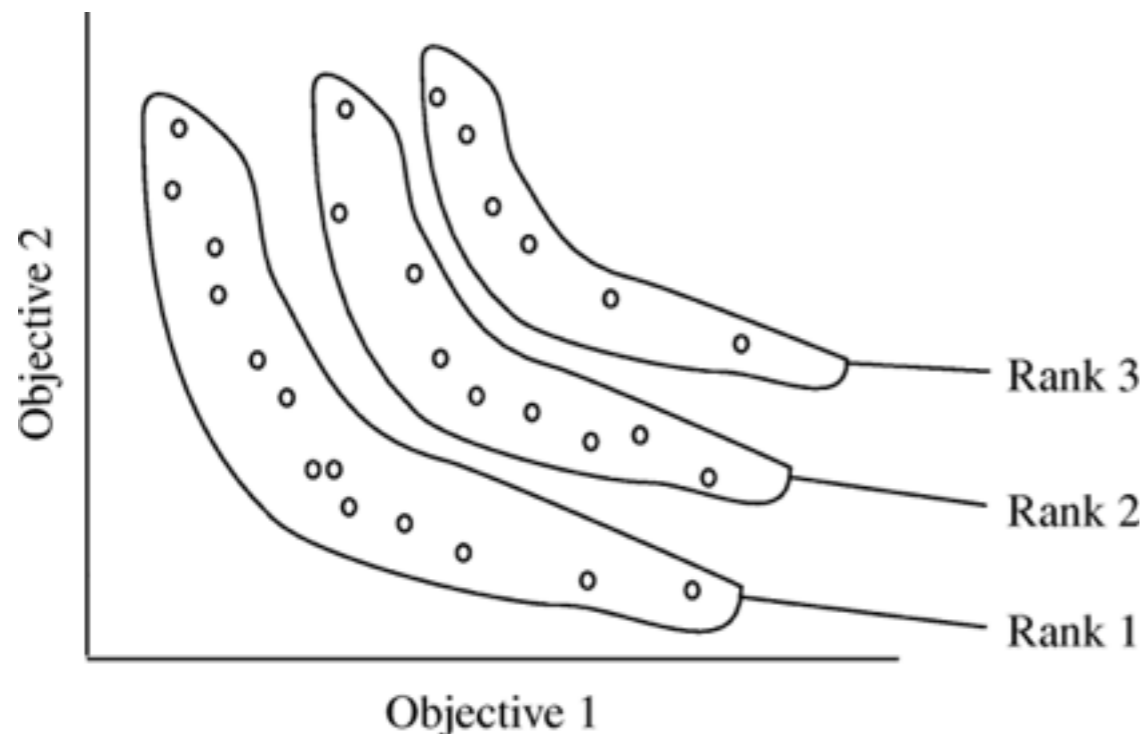


- The ideal goal is to obtain the Pareto front
- Unfortunately, this is unpractical in real-world problems
  - NP-hard complexity, non-linearity, epistasis, ...
  - Frequently, exact techniques are not useful
- Alternative: Use non-exact algorithms
  - E.g. Metaheuristics
  - These techniques provide an approximation to the Pareto front

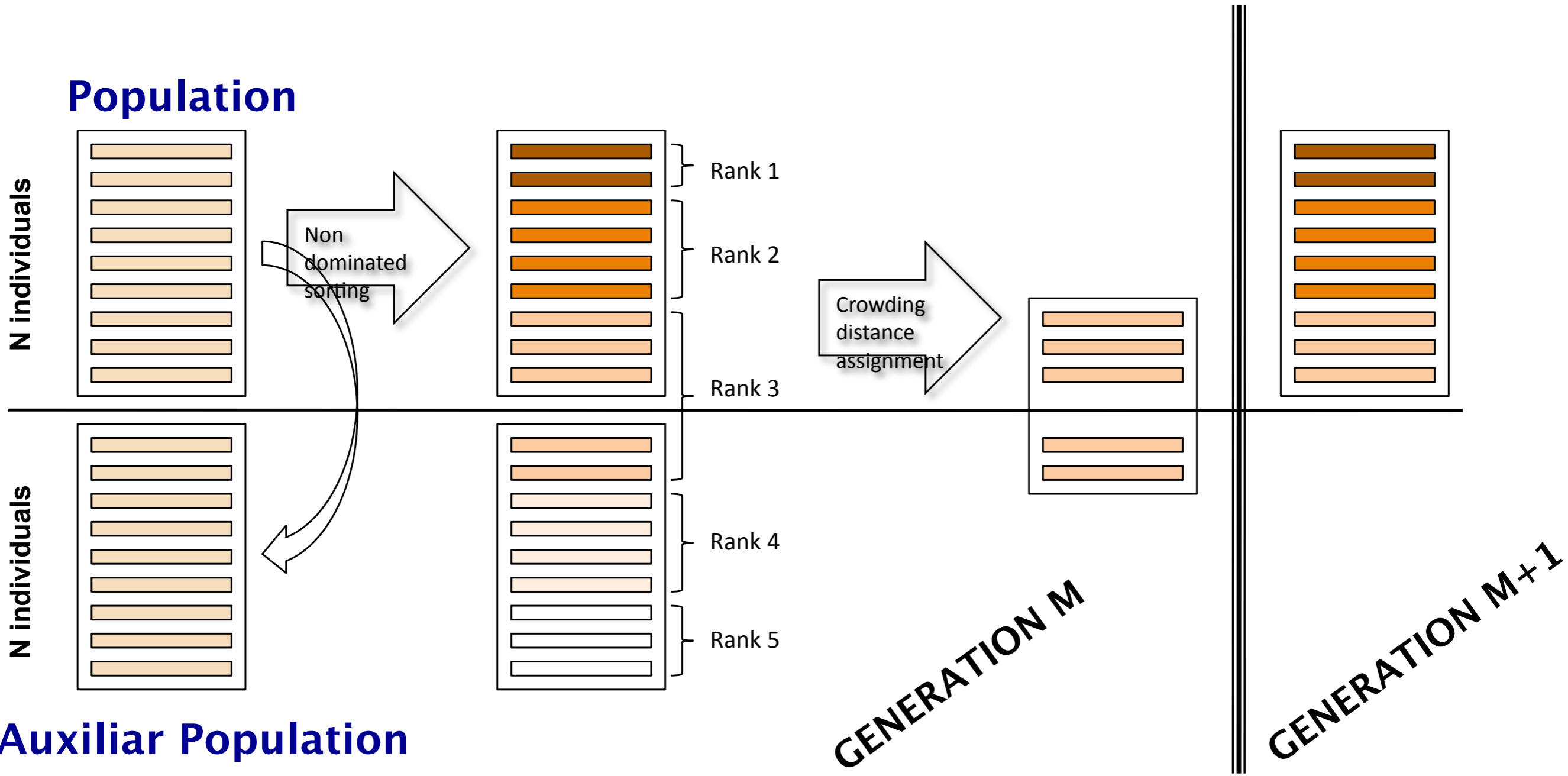
- The goal is to find the **Pareto front**
- Exact techniques are **not useful** in most cases
  - NP-hard complexity, non-linearity, epistasis , ...
- Rely on **approximation** techniques
  - Two key features to measure the **quality** of solutions
    - ▶ Convergence
    - ▶ Diversity

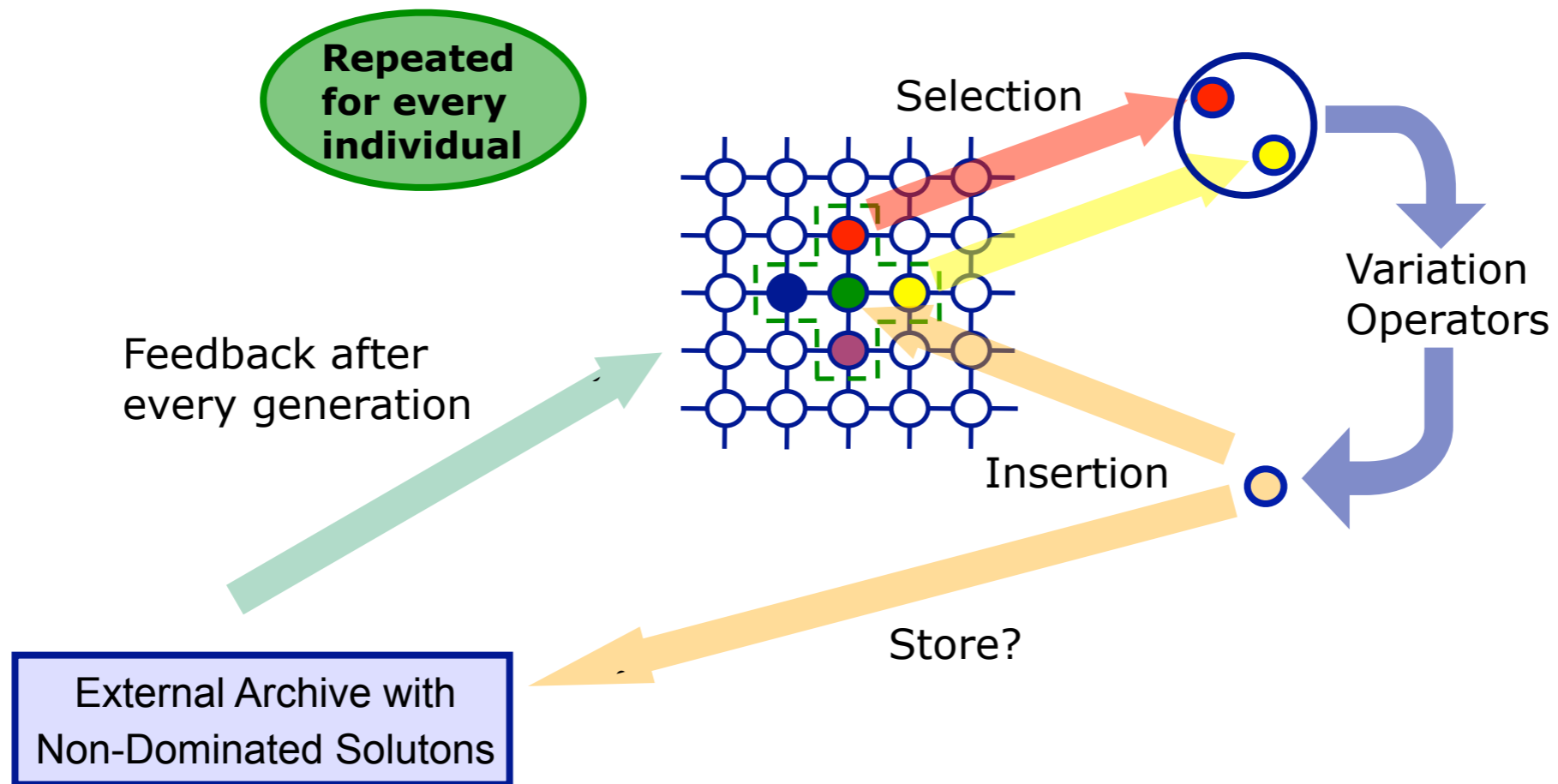


- Non-dominated Sorting Genetic Algorithm
- The most popular metaheuristic for multi-objective optimization
- Features
  - Ranking using non-dominated sorting
  - Crowding distance as density estimator

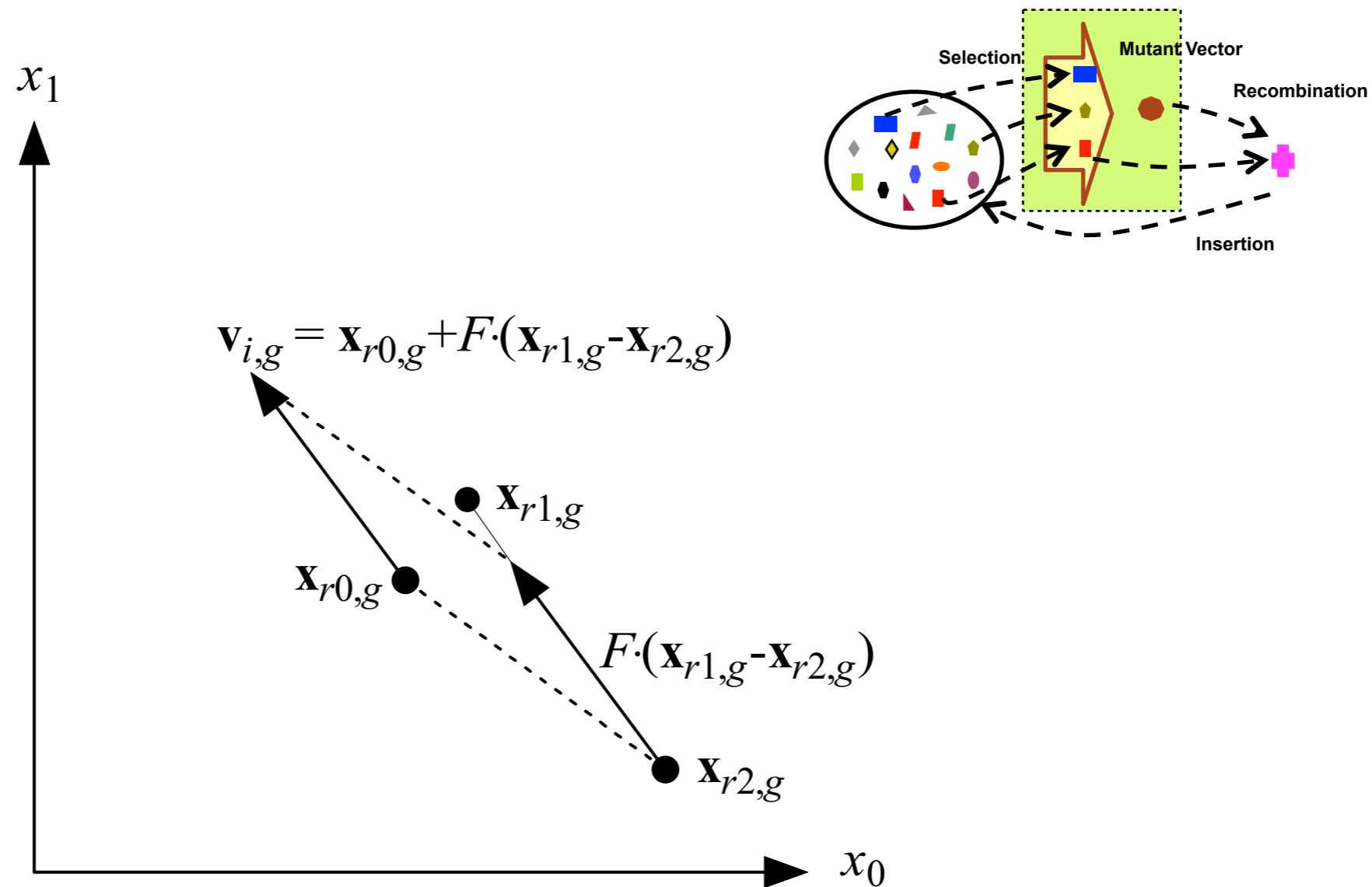


Point B is in a less crowded region than point A

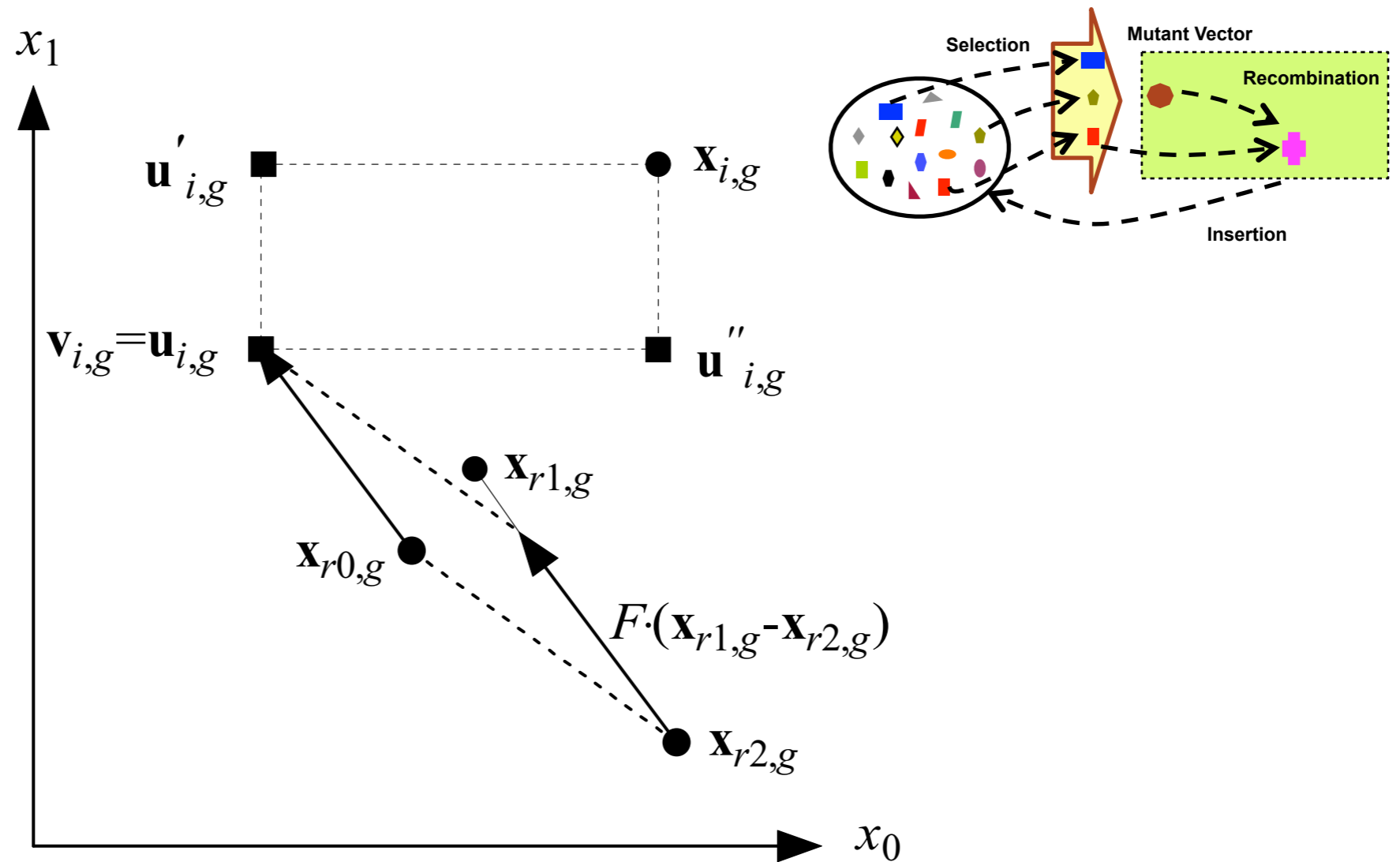




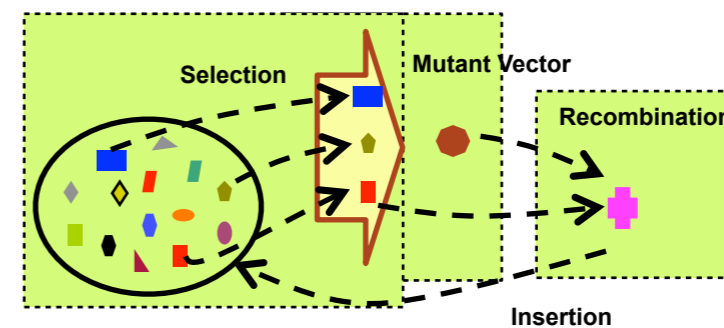
- Mutant vector generation



- Recombination



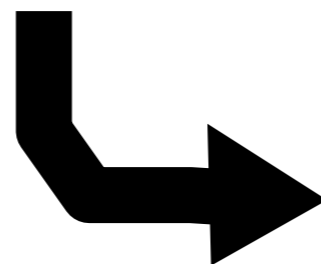
$X_{r0,g}$	1	5	-2,5	30,7	0,4
$X_{r1,g}$	5,7	8,9	0,3	-4,3	30,5
$X_{r2,g}$	4,2	7,2	3,3	8,2	6,7



$V_{i,g}$	1,75	5,85	-0,7	24,45	12,3
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$$U_{i,g} = \begin{cases} V_{i,g} & \text{With prob. CR} \\ X_{r0,g} & \text{otherwise} \end{cases}$$

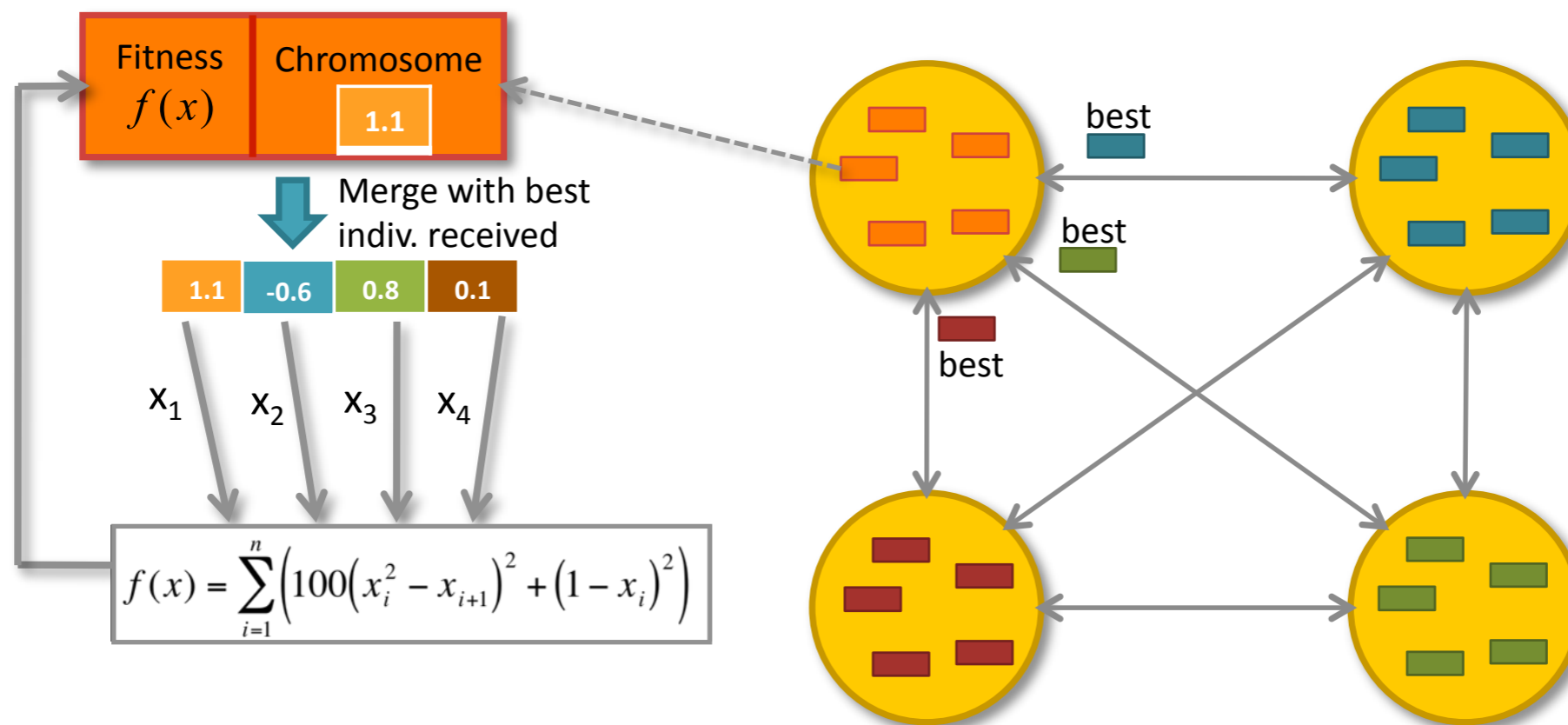
$X_{r0,g}$	1	5	-2,5	30,7	0,4
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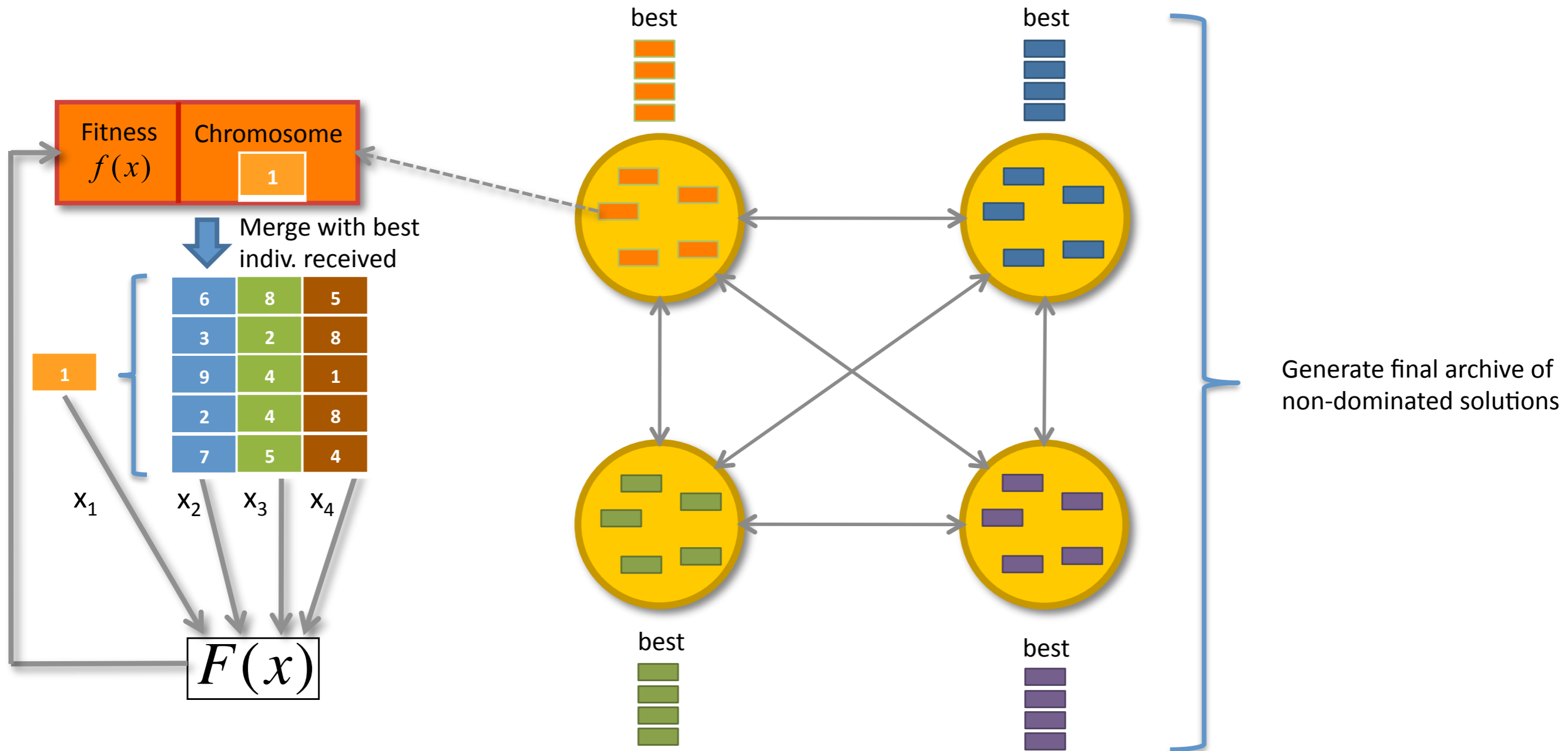


$U_{i,g}$	1,75	5	-0,7	30,7	0,4
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- Each node runs a subpopulation for a **subset of the N variables**
- Each population **evaluates** each of its individuals **on the global fitness function** using the best individual received from each other subpopulation





- Three CCMOEAAs designed
  - Based on NSGA-II: CCNSGAI
  - Based on SPEA2: CCSPEA2
  - Based on MOCcell: CCMOCcell

## NSGA-II

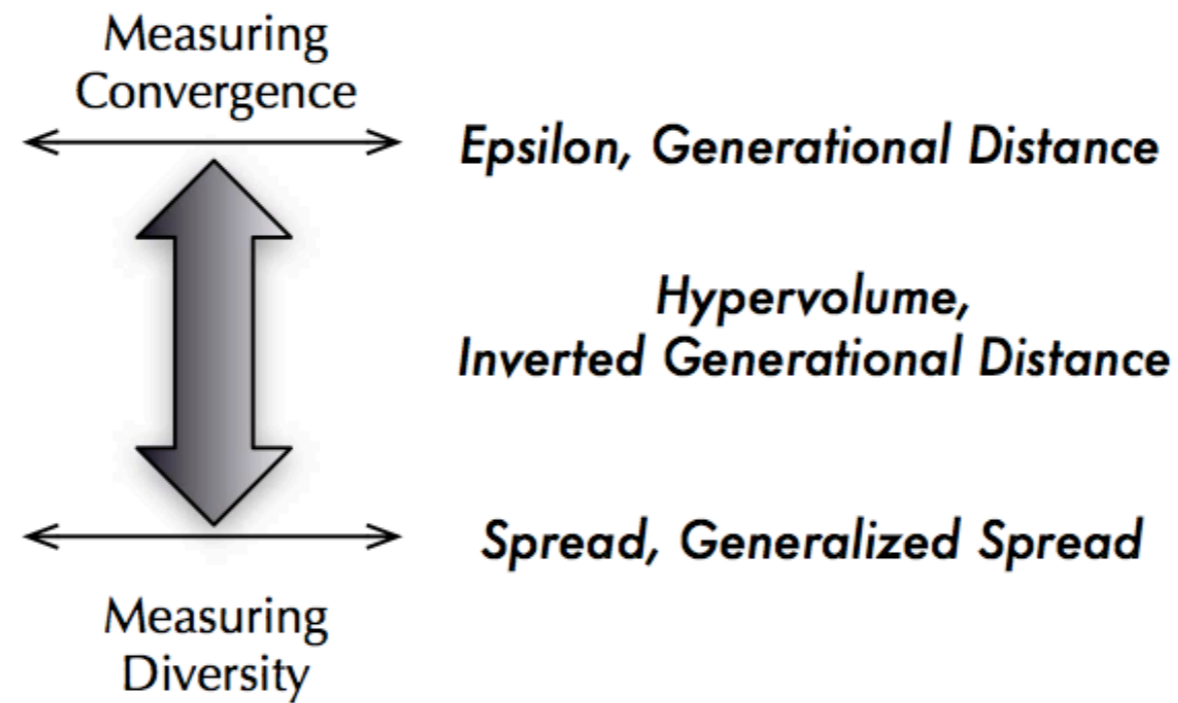
- Reference algorithm
- Panmictic population
- Selection of solutions
  - Ranking
  - Crowding

## SPEA2

- Panmictic population
- External archive
  - Strength raw fitness
  - k-nearest neighbors

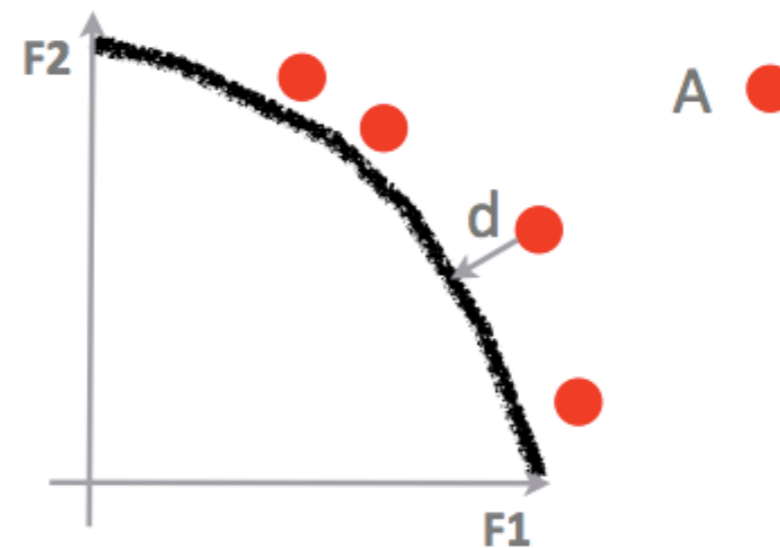
## MOCcell

- Cellular population
  - Only next individuals can interact
- External archive
  - Feedback to population



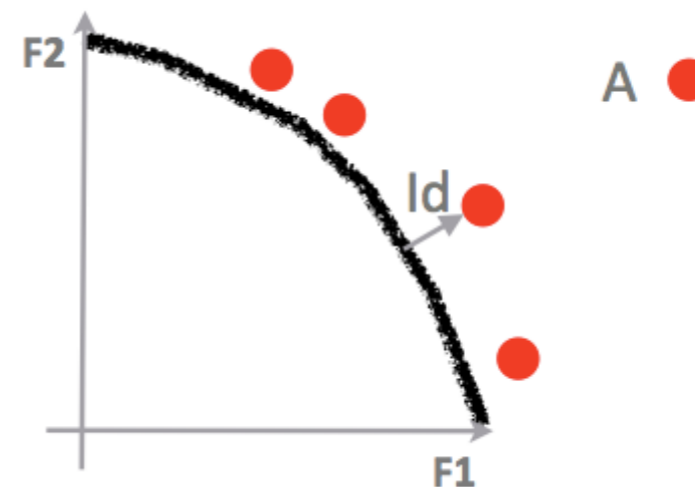
- Generational distance
  - Average distance of every solution of a front A to the Pareto front
  - Convergence to the true Pareto front

$$GD(A) = \frac{\sum_{1..n} d}{n}$$

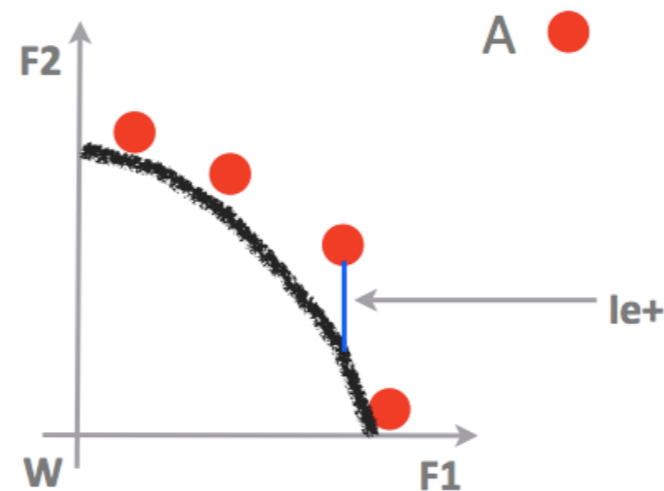


- Inverted generational distance
  - Average distance of every solution of the points of the Pareto front to of a front A
  - Convergence to the true Pareto front

$$\text{IGD}(A) = \frac{\sum_{1..n} Id}{n}$$

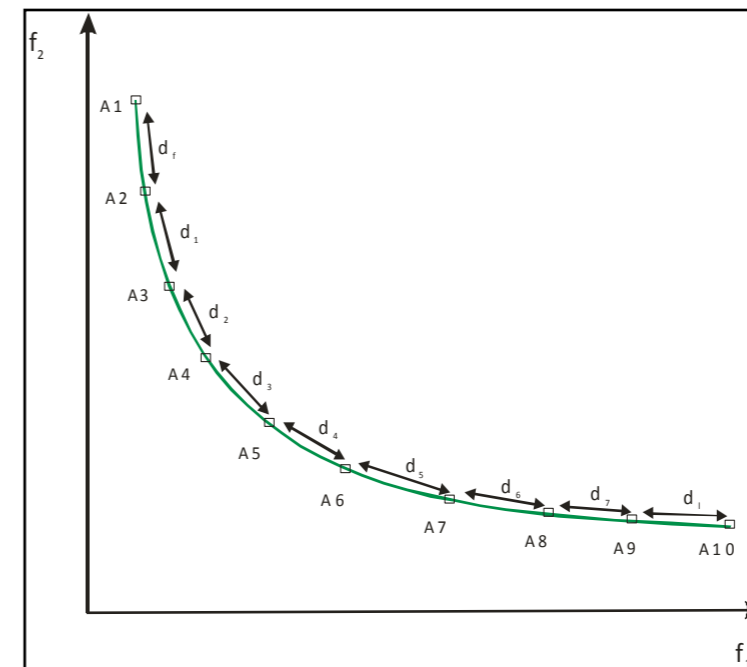


- Additive epsilon indicator
  - Convergence to the Pareto front
  - Given an approximation set  $A$ , this indicator is a measure of the smallest distance we would need to translate every solution in  $A$  so that it dominates the Pareto front



- Spread
  - Diversity of the solutions along the Pareto front

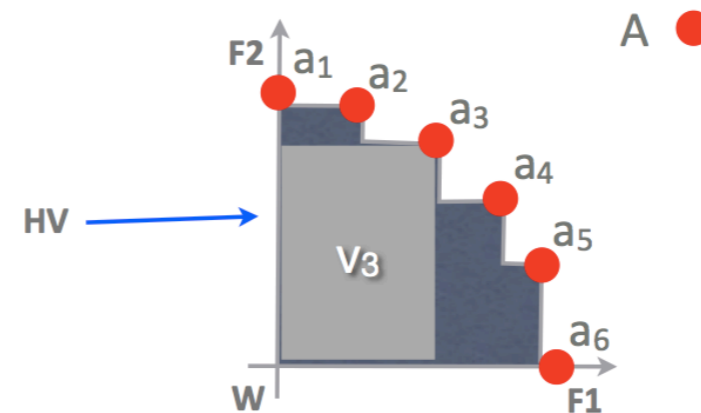
$$\Delta = \frac{d_f + d_l + \sum_{i=1}^{N-1} |d_i - \bar{d}|}{d_f + d_l + (N - 1)\bar{d}}$$





- Hypervolume
  - Takes into account both convergence and diversity
  - Measures the region dominated by the obtained front

$$HV = volume \left( \bigcup_{i=1}^{|Q|} v_i \right)$$



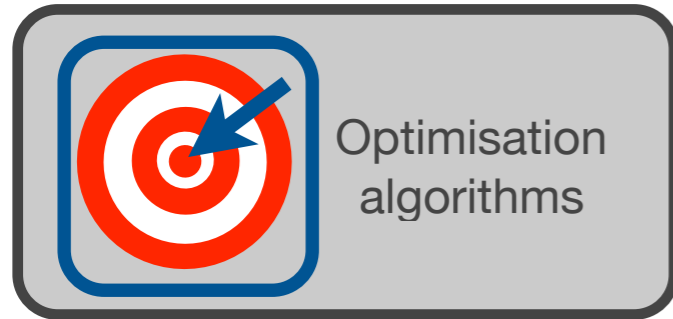
- All discussed metrics require the optimal Pareto front
  - Either for computations
  - Or to normalize the fronts
- What if we do not know it?
  - Build a reference Pareto front of (hopefully) quasi-optimal solutions
    - ▶ Run the problem with different algorithms
    - ▶ Run every algorithm a large number of times
    - ▶ Take the best non-dominated solutions found in all runs by all algorithms

- Metaheuristics are stochastic algorithms
  - Repeating the same experiment may lead to different results
  - It is necessary to apply a rigorous statistical methodology to assess the performance of a metaheuristic
- To draw firm conclusions, we need to look for statistical significance on the results

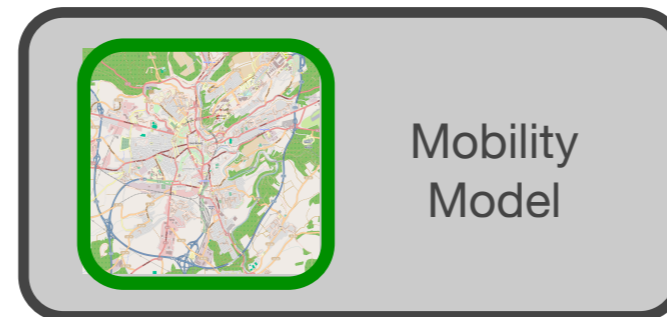
- Statistical significance
  - Large number of independent runs
  - Compute quality metrics
  - Statistical test on the results of the quality metrics
    - ▶ Non-parametric test: Wilcoxon unpaired signed-ranks test
  - Confidence level of 95%
    - ▶ Significance level of 5% or p-value under 0.05 in the statistical tests
    - ▶ This means that the differences are unlikely to have occurred by chance with a probability of 95%

# Optimization Framework for Mobile Ad Hoc Networks

Optimization algorithm



Mobility simulation



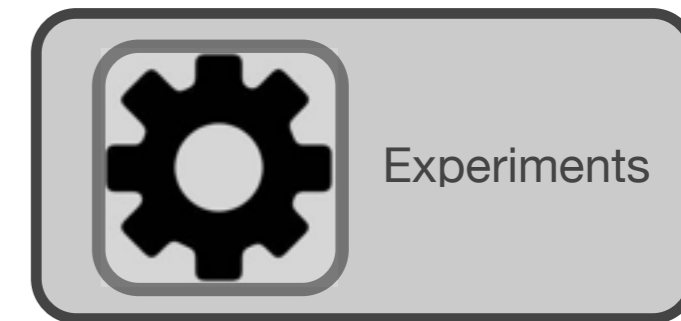
Network simulator



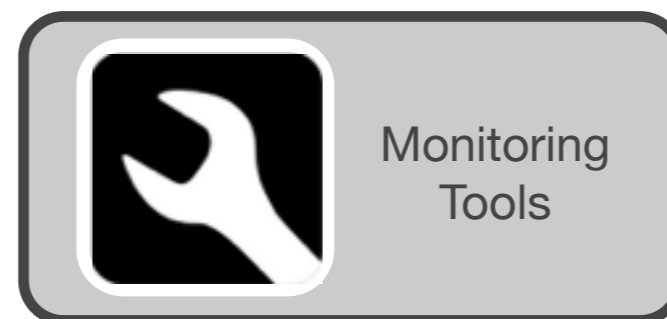
Protocol to optimize



Configuration of simulations

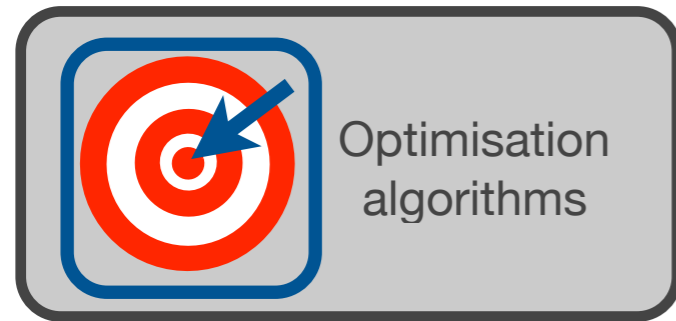


Performance measurements

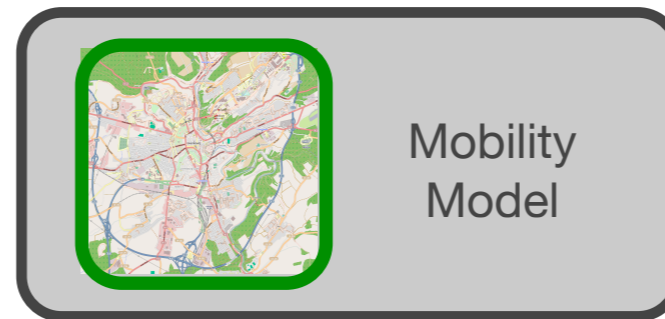


# Off-line optimization process

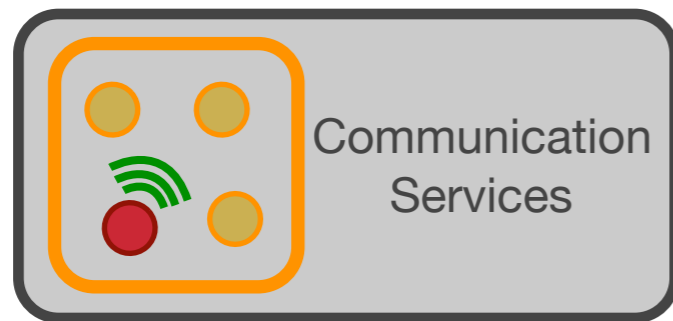
Optimization algorithm



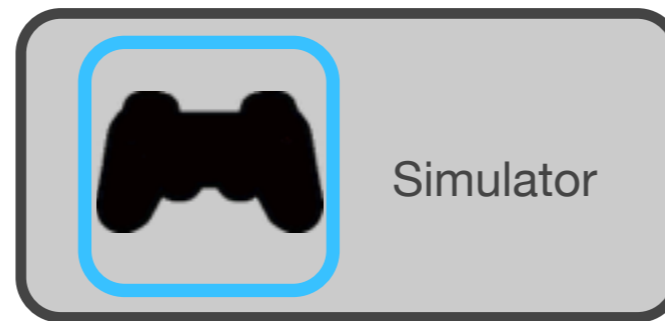
Mobility simulation



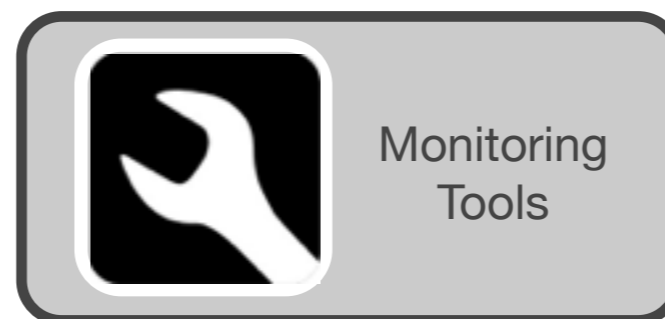
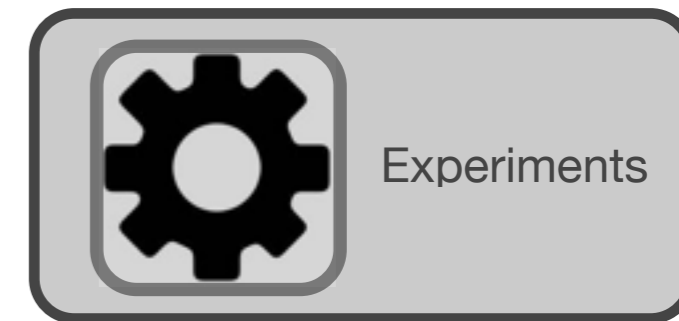
Network simulator



Protocol to optimize



Configuration of simulations



Performance measurements

