



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

Others

Metaheuristic Applications to Telecoms, Bioinf, Software, and other Domains



<http://neo.lcc.uma.es>

University of Málaga, Spain



Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

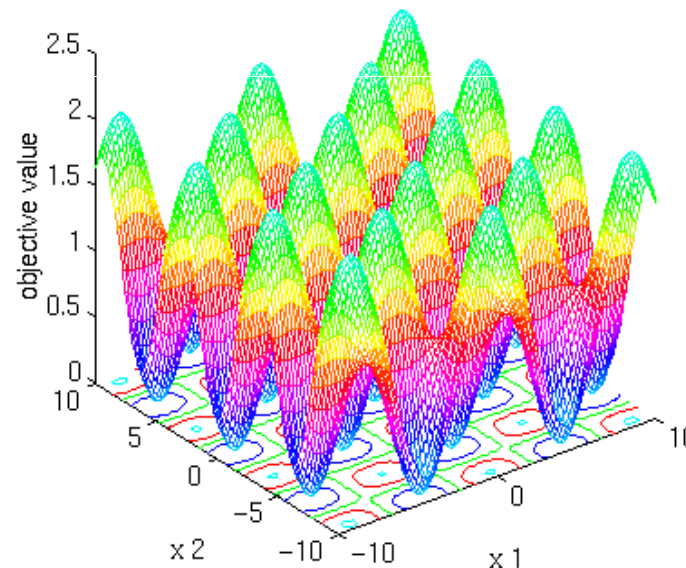
Software

Others

Introduction (I)

- **Objective of a global optimization problem:**

$$f(\vec{x}) \rightarrow \max : \text{find a vector } \vec{x}^* \\ \text{such that } \forall \vec{x} \in M : f(\vec{x}) \leq f(\vec{x}^*) := f^*$$



- **Minimizing is also possible**
- **Vectors can map to other data structures**



Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

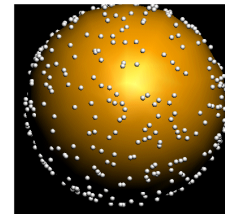
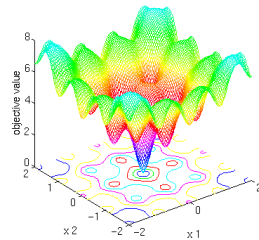
Bioinformatics

Software

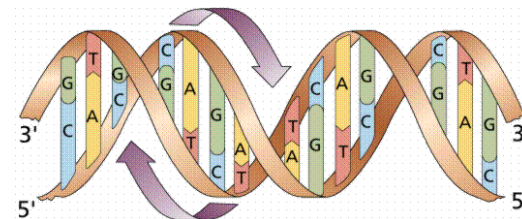
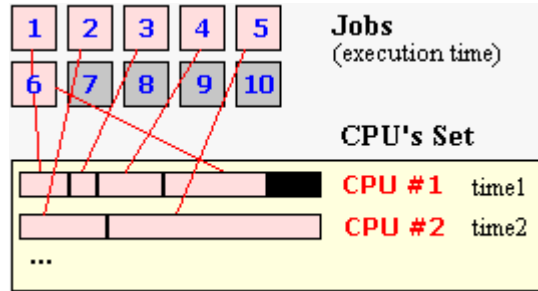
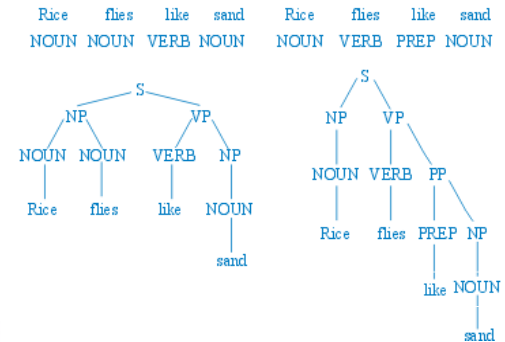
Others

Introduction (II)

Where can optimization problems be found?



Distribute "n" electrons on a sphere





Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

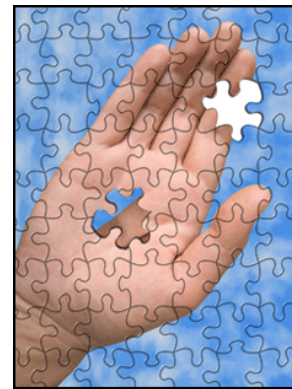
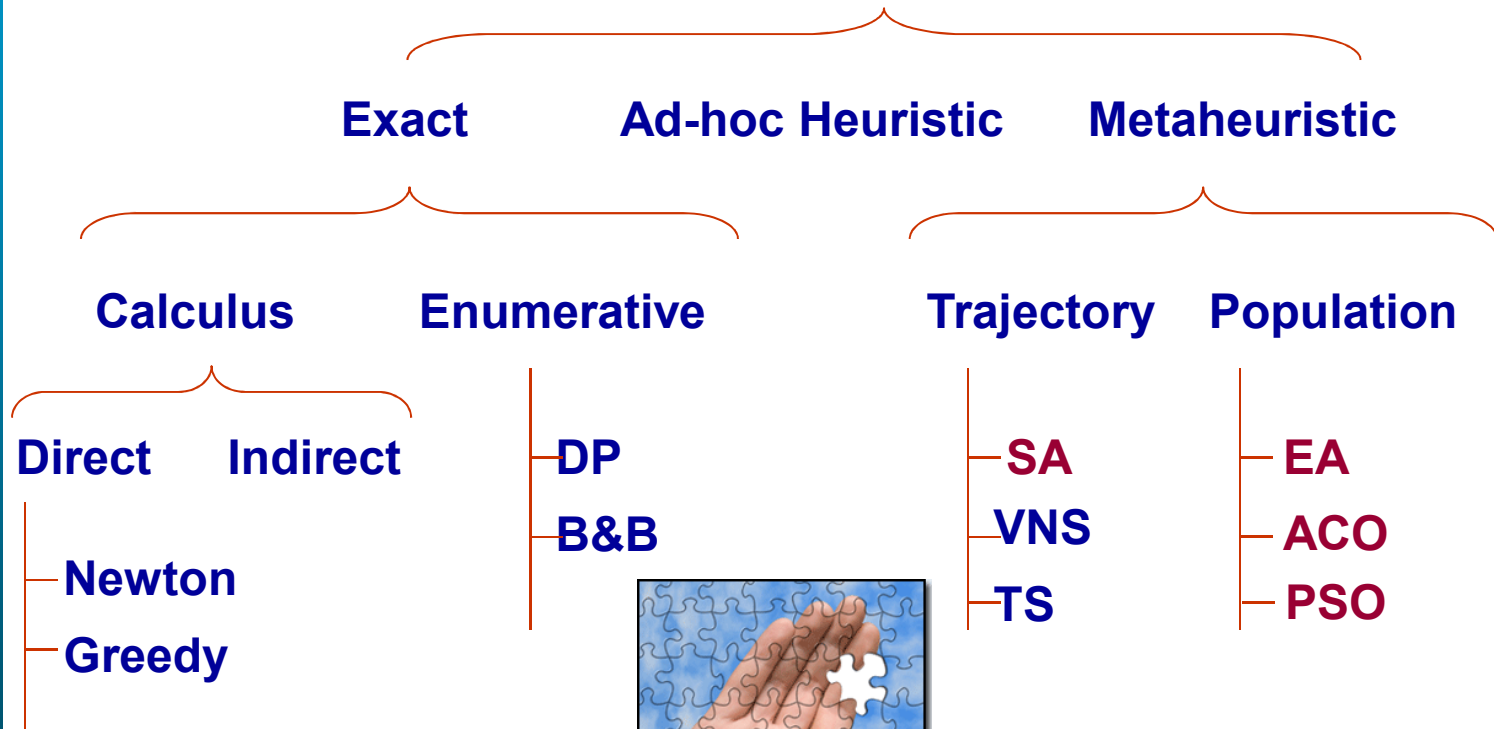
Bioinformatics

Software

Others

Introduction (III)

Optimization Algorithms



• nature inspired in red



Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

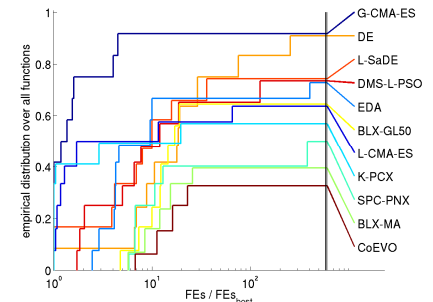
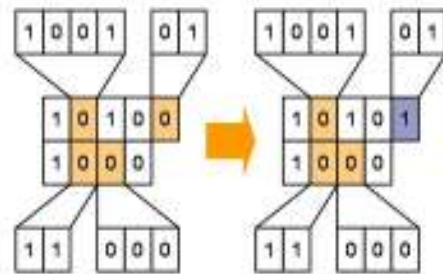
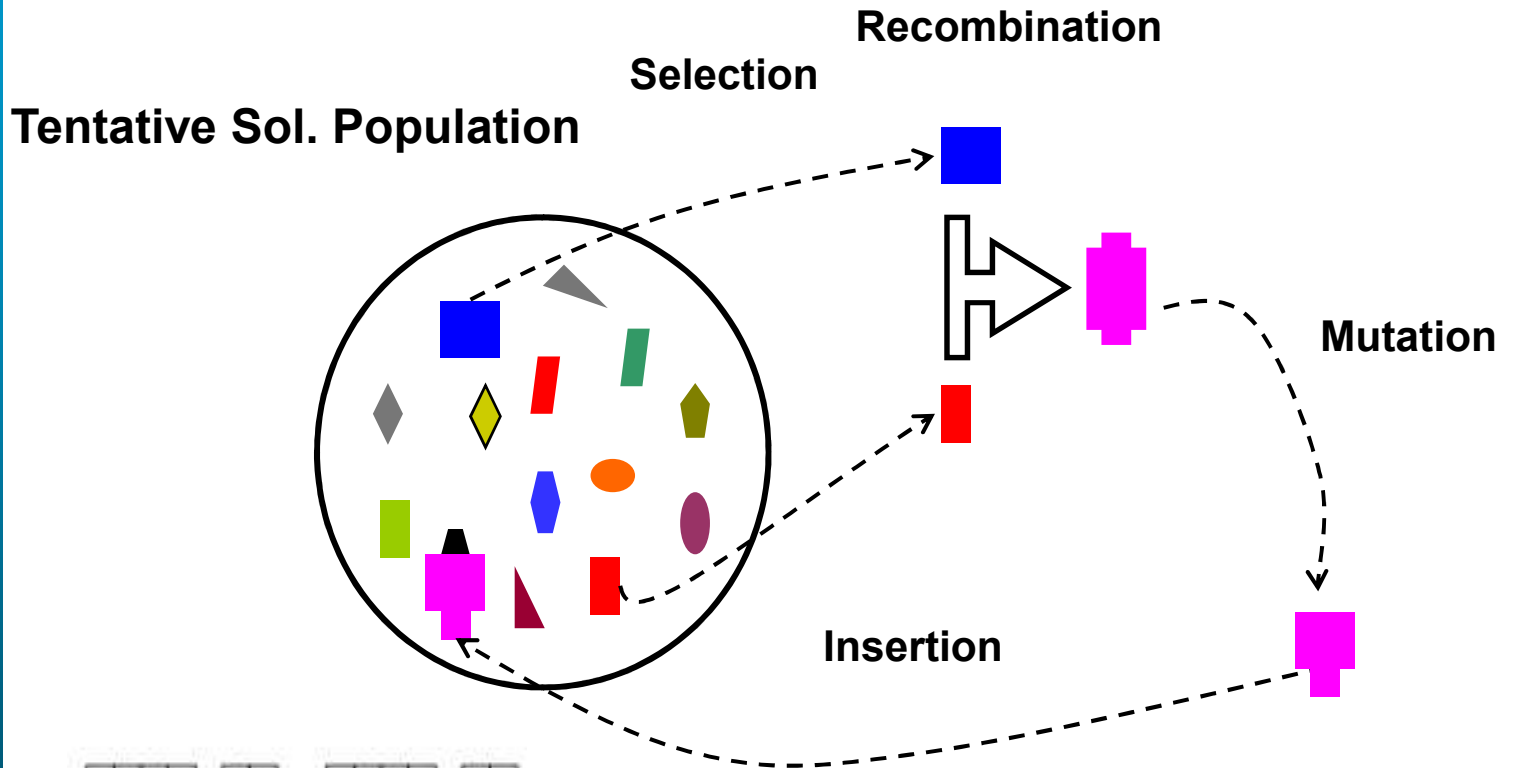
Telecoms.

Bioinformatics

Software

Others

Introduction (IV)





Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

Others

Introduction (V)

Evolutionary Algorithm

```
 $t := 0;$   
initialize  $[P(t)];$   
evaluate  $[P(t)];$   
while not end condition do  
     $P'(t) := \text{variation } [P(t)];$   
    evaluate  $[P'(t)];$   
     $P(t+1) := \text{selection } [P'(t) \cup Q];$   
     $t := t + 1 ;$   
end while
```



Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

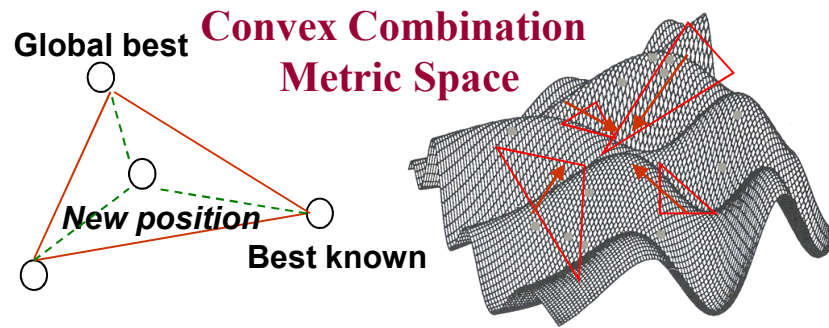
Telecoms.

Bioinformatics

Software

Others

Introduction (VI)

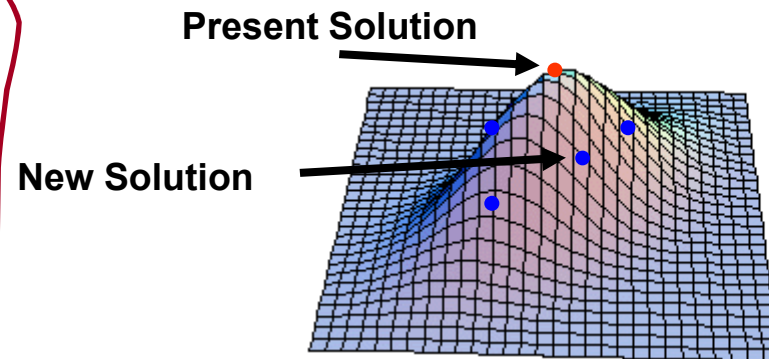


```

procedure ACOMetaheuristic
  ScheduleActivities
    ConstructAntsSolutions
    UpdatePheromones
    DaemonActions // optional
end ScheduleActivities
end procedure
    
```



- (0,2; -1,4; 3,5)** → **Solution Vector**
- (1,0; 10,3;** → **Standard Deviation**
- (1,7; 0,3; 2,1)** → **Search Angles**





Index

➔ New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

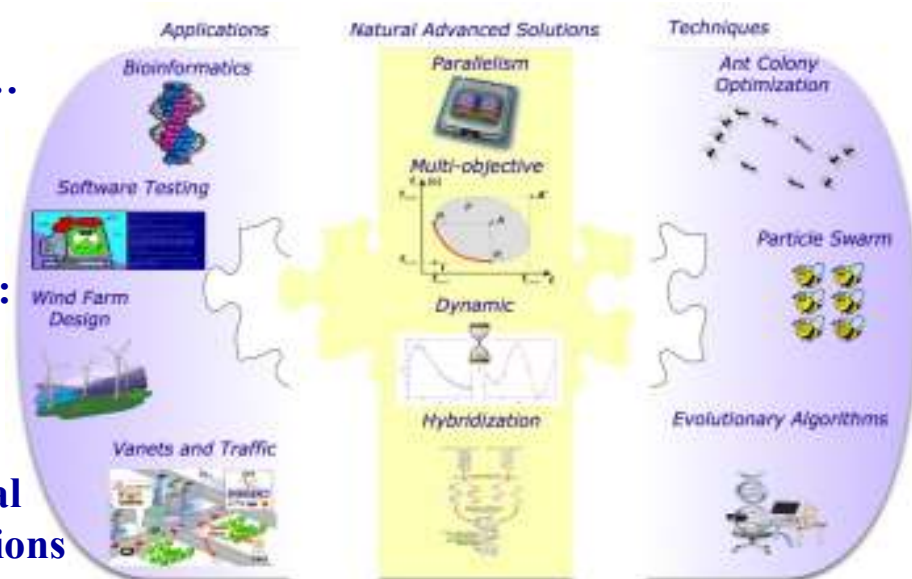
Others

Introduction (VII)

NAS: Natural Advanced Solutions

Four main ways of making an algorithm more efficient and accurate:

- Parallel:**
 Clusters, Grid computing, multicore, FPGAs, GPUs...
- Hybrid:**
 Combining algorithms, operators, representations: problem knowledge
- Multiobjective:**
 Modelling explicitly several conflicting objective functions with Pareto's concept of dominance
- Dynamic:**
 Solve a problem that changes in time and adapt previous solutions to the new scenarios





Index

New Models

➔ Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

Others

Parallel Metaheuristics

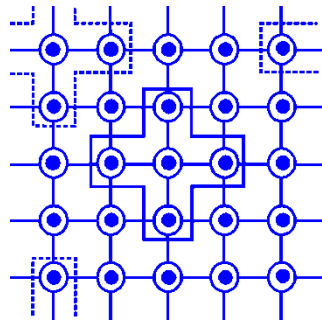
Paralellism and Metaheuristics:

The increasing availability of new kinds of CPUs and the parallel nature of metaheuristics have allowed the fast development of parallel metaheuristics

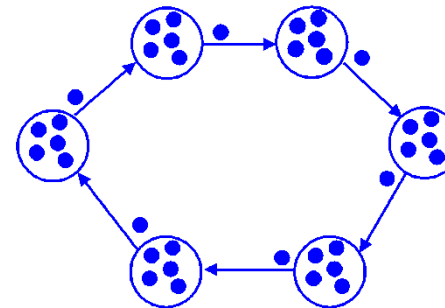


Advantages:

- Allow to tackle more complex problems and/or larger instances **GPU**
- Allow to reduce the execution time
- Allow to improve the quality of the found solutions



Examples



multicore



E. Alba (ed.), *Parallel Metaheuristics: A New Class of Algorithms*, Wiley & Sons, 2005



Index

New Models

Parallel MetaH

➔ Hybrid MetaH

Multiobjective

Telecoms.

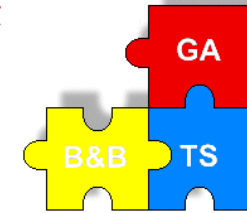
Bioinformatics

Software

Others

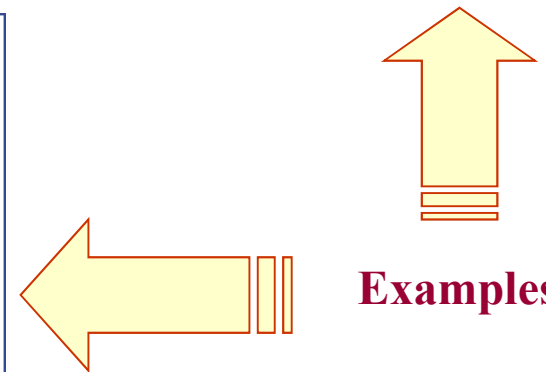
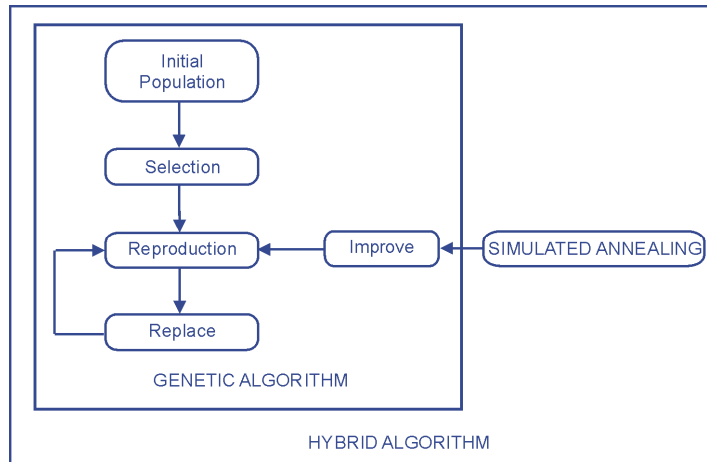
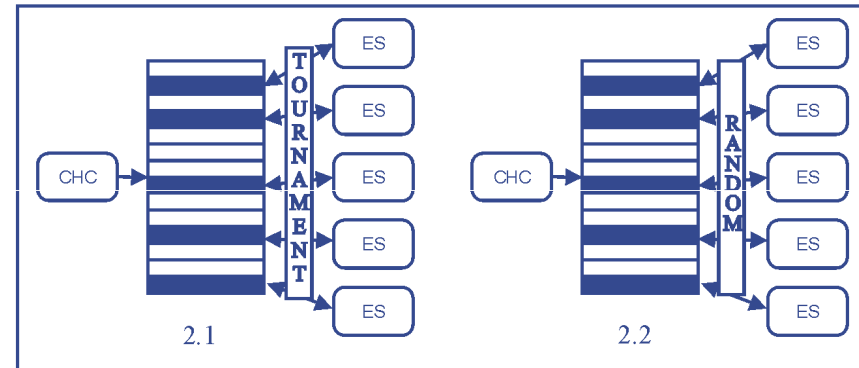
Hybrid Algorithms

Hybridization is the inclusion of problem-dependant information in the algorithm



Types:

- Strong
- Weak



Examples



Index

New Models

Parallel MetaH

Hybrid MetaH

➔ Multiobjective

Telecoms.

Bioinformatics

Software

Others

Multiobjective Optimization

Most real word optimization problems require to optimize more than one single function

- **Multiobjective Optimization Problems (MOPs)**

Multiojective optimization searches for a set of solutions

- **Pareto Optimal Set**
- **Their representation in the objective space is known as Pareto front**

Metaheuristics provide a subset of the Pareto optimal set.

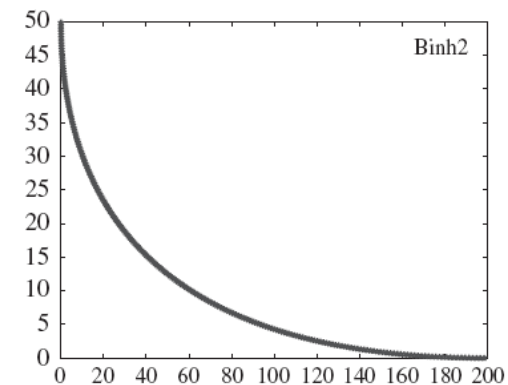
Two goals

- **Convergence to the true Pareto front**
- **Diversity of the solutions along the true Pareto front**

$$\begin{aligned} \text{Min } F &= (f_1(\vec{x}), f_2(\vec{x})) \\ f_1(\vec{x}) &= 4x_1^2 + 4x_2 \\ f_2(\vec{x}) &= (x_1 - 5)^2 + (x_2 - 5)^2 \end{aligned}$$

Subject to:

$$\begin{aligned} g_1(\vec{x}) &= (x_1 - 5)^2 + x_2^2 - 25 &\leq 0 \\ g_2(\vec{x}) &= -(x_1 - 8)^2 - (x_2 + 3)^2 + 7.7 &\leq 0 \\ 0 &\leq x_1 &\leq 5 \\ 0 &\geq x_2 &\leq 3 \end{aligned}$$





Index

New Models

Parallel MetaH

Hybrid MetaH

➔ Multiobjective

Telecoms.

Bioinformatics

Software

Others

Multiobjective Optimization: Open Issues

- Fitness assignment

- Multiobjective metaheuristics assign a unique value to the solutions used as “fitness” to compare solutions
- E.g.: Ranking in NSGA-II or strength in SPEA2

- Maintaining diversity

- Additional information is needed to know the density of solutions around a given one
- E.g.: Hypercube in PAES, crowding distance in NSGA-II

- Elitism

- The general approach uses an auxiliary population, sometimes called *archive*

- Quality indicators

- Metrics are needed to measure convergence and/or diversity
 - Hypervolume (convergence and diversity)
 - Generational Distance (convergence)
 - Spread (diversity)



Index

New Models

Parallel MetaH

Hybrid MetaH

➔ Multiobjective

Telecoms.

Bioinformatics

Software

Others

MOCe11

MOCe11

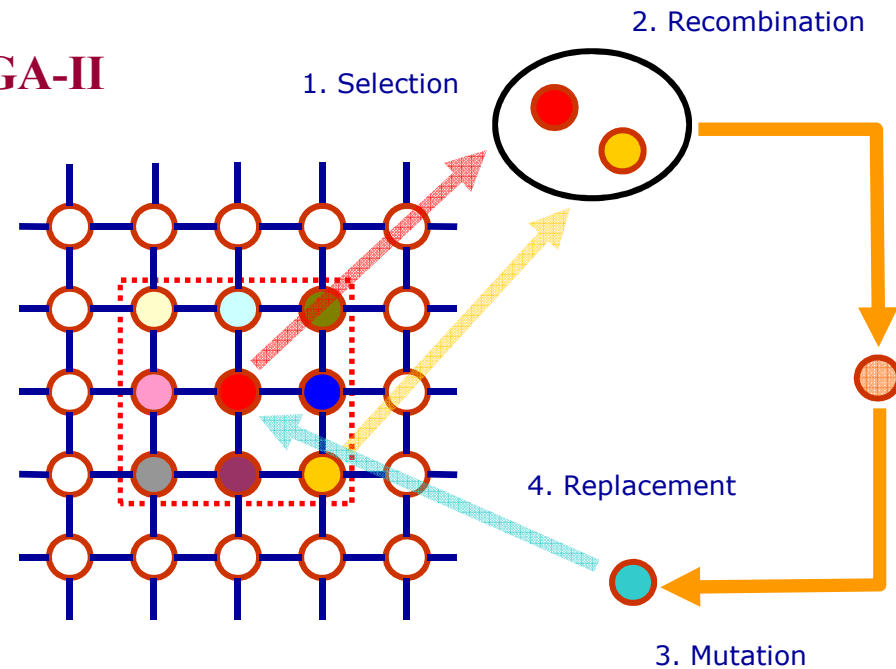
- Multiobjective cellular genetic algorithm

Main features

- Use of an external archive
 - 2-dimensions toroidal grid
 - Archive feedback

Comparison against NSGA-II and SPEA2

- Competitive results in terms of convergence and hypervolume
- Best results concerning spread





Index

New Models

Parallel MetaH

Hybrid MetaH

➔ Multiobjective

Telecoms.

Bioinformatics

Software

Others

AbYSS

AbYSS

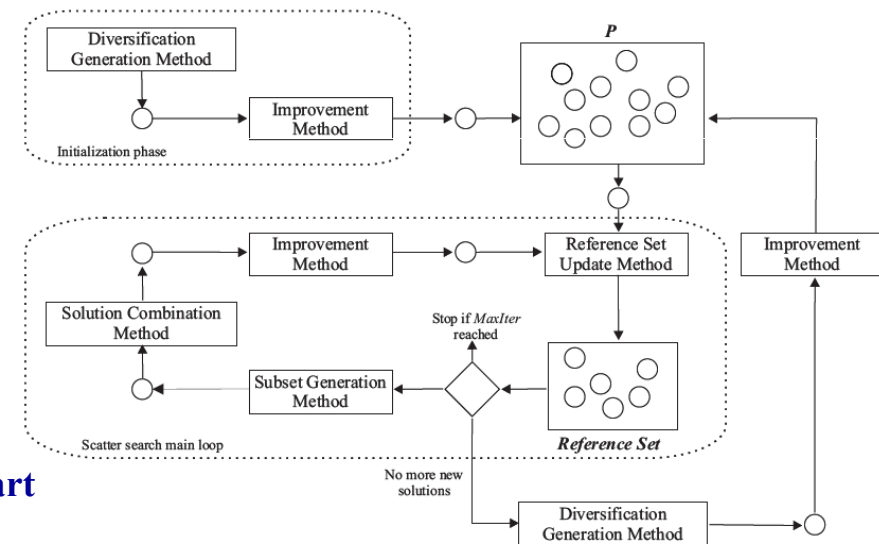
- Archive based hYbrid Scatter Search

Basic idea

- Redefining the scatter search template to adapt it to multiobjective optimization

Main features

- External Archive to maintain good solutions
- Individuals of the external archive are moved to initial set in the re-start loop



Comparison against NSGA-II and SPEA2

- Competitive results in terms of convergence and hypervolume
- Best results concerning spread



Index

New Models

Parallel MetaH

Hybrid MetaH

➔ Multiobjective

Telecoms.

Bioinformatics

Software

Others

Multiobjective Optimization and Grid Computing

Motivation:

- There are many computers in the labs of the Computer Science Department of the University of Málaga
 - Currently we directly control up to 400 processors
- Question: How can we use them together to solve multiobjective optimization problems?
- Using known message passing libraries (sockets, PVM, MPI) is not a solution
 - Machines are idle in the nights and during the weekends (and in holydays)
 - Variable availability during the day
- OUR APPROACH: using grid technologies

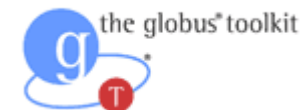
Grid computing systems used:

- Condor
- Globus
- Others: ProActive, BOINC



Issues of interest:

- Easiness of installation and administration
- Parallel programming models offered
- Programming languages available
- Use of idle CPU cycles (opportunistic computing)
- Parallel performance





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

Software

Others

Optimal Broadcasting in MANETs

- MANETs

- Stations usually are laptops, handholds, PDAs, or mobile phones
- Mobility of stations → dynamic topology of the network

- Metropolitan MANETs

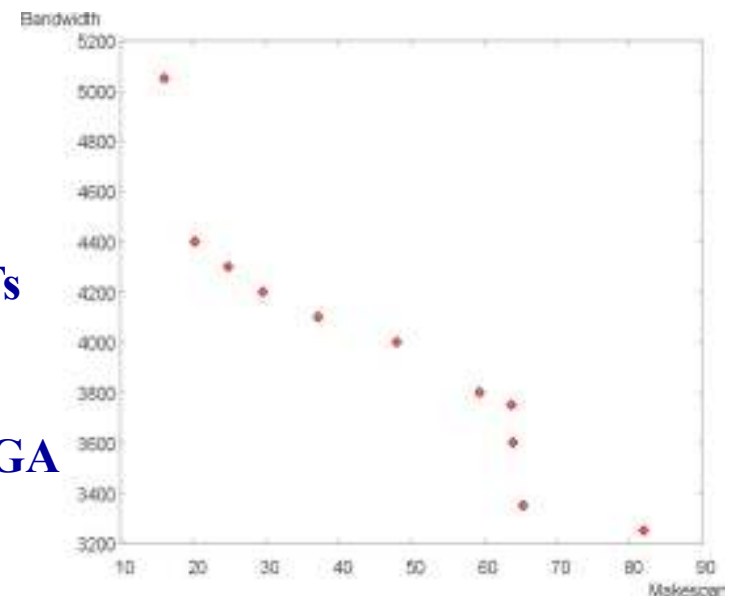
- High Density Areas (HDA): areas with high station density
- HDAs can appear and disappear from the network

- Optimization Problem

- Fine-tune of a broadcasting strategy called DFCN
- Target: metropolitan MANETs

- Multiobjective metaheuristics

- EAs: NSGA-II, SPEA2, cMOGA
- Scatter Search: AbYSS
- PSO: MOPSO





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

Software

Others

Automatic Frequency Planning in GSM Networks

- Problem definition

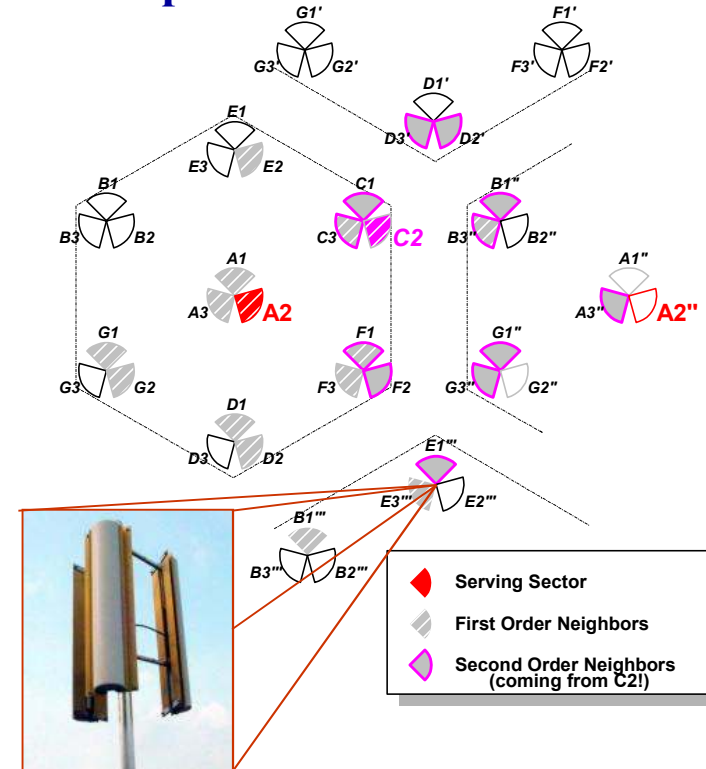
- Allocate frequencies (few dozens) to elementary transceivers (TRXs) of the network (thousands)
- Frequency reuse is mandatory → this provokes interference → QoS degradation
- Real world instances of GSM networks currently in use

- GSM architecture

- Base Transceiver Stations
- Sectors
- TRXs

- Metaheuristics used

- (1,λ) EA, ACO
- Ongoing:
 - ssGA
 - Grid computing with Condor





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

Software

Others

Automatic Cell Planning in 2G/3G Networks

- Problem definition

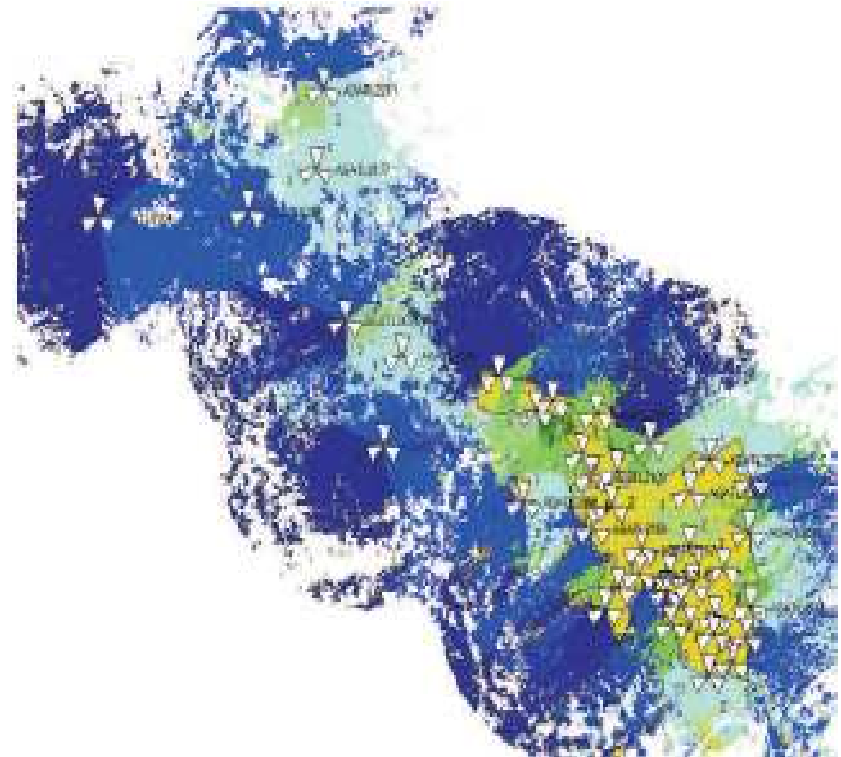
- Positioning the sites of the network
- Dimensioning a set of parameters for each antenna
- Real world instances of cellular networks
 - Highly demanding computational costs

- Objectives

- Number of sites
- Quality of service
 - Interferences
 - Traffic hold

- Metaheuristics used

- PAES, ssGA
- New work in:
 - AbYSS
 - MOCcell
 - Gridifying with
Condor/MW





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

Software

Others

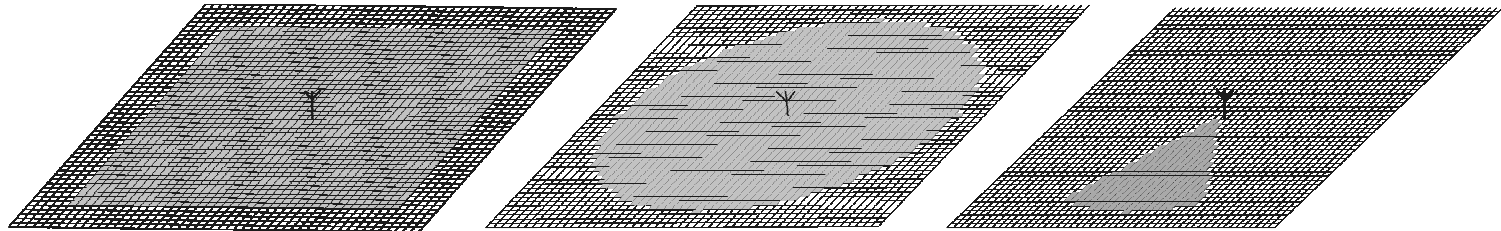
Radio Network Design: Problem

Task: Select geographical locations for cellular antennae from a list of 149 to 349 available locations

Objectives: Maximize the covered area (coverage)
Minimize the number of antennae

Fitness:
$$f(\vec{x}) = \frac{Coverage(\vec{x})^\alpha}{Number\ of\ antennae(\vec{x})} \quad \text{with } \alpha = 2$$

Models: Terrain -> Square grid with 287×287 points
Antenna coverage -> Three shapes





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

→ Telecoms.

Bioinformatics

Software

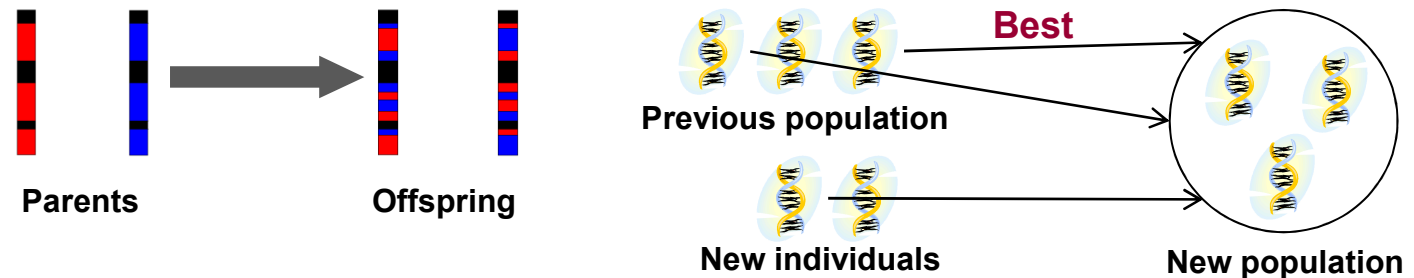
Others

Radio Network Design: Algorithms

SA: Trajectory technique
 Uses mutation to generate S_n from S_a
 Replaces S_a with S_n with probability P

$$P = \frac{2}{1 + e^{\frac{fitness(S_a) - fitness(S_n)}{T}}}$$

CHC: Evolutionary algorithm
 No mutation, HUX crossover with incest prevention
 Elitist selection, Cataclysmic mutation (restarting procedure)



GA: Evolutionary algorithm
 Mutation, two-point crossover
 Selection: Ranking (parents) + tournament (new population)



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

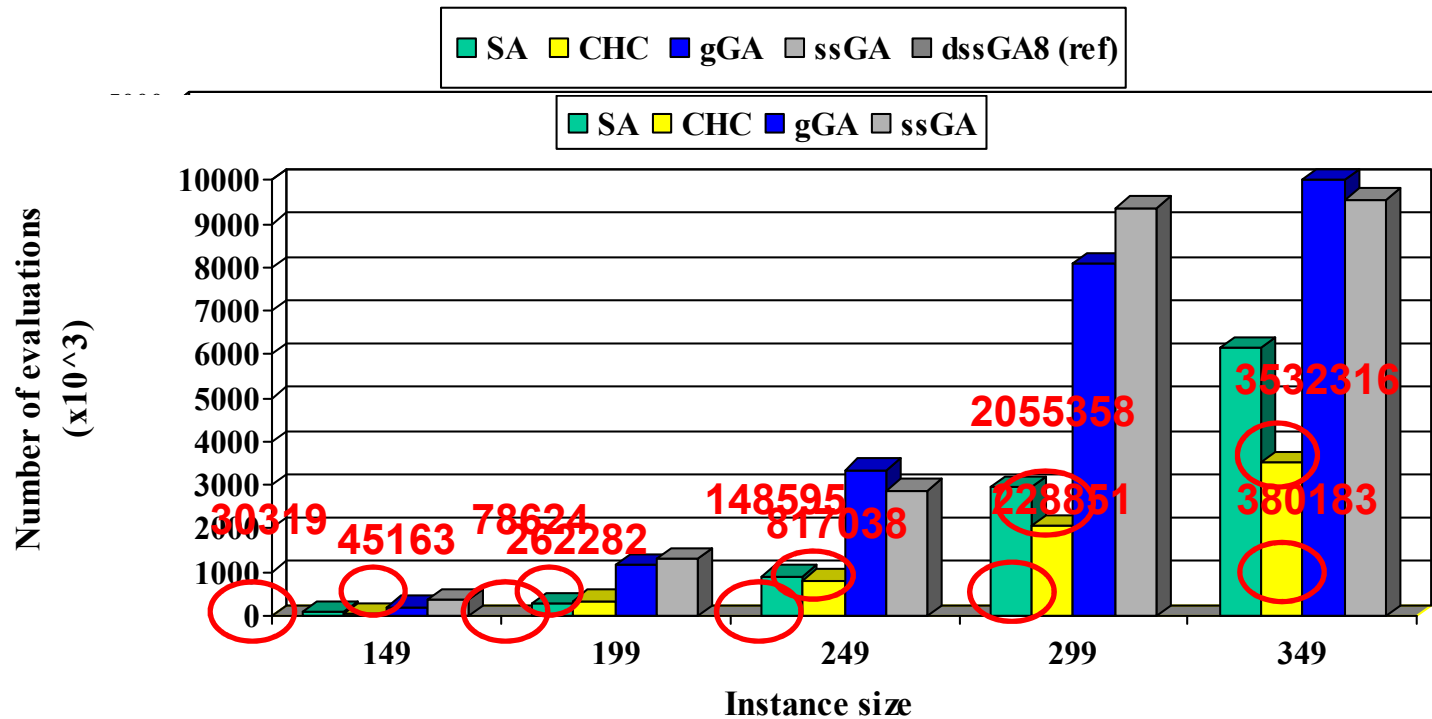
Bioinformatics

Software

Others

Radio Network Design: Results

Antenna cell: ~~Omni~~ **Quadrifunctional**



- **CHC** finds the optimal solution with less effort than SA and GA for the five instance sizes using both antenna types
- The geometry of the problem affects the complexity



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

Software

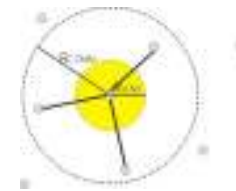
Others

Wireless Sensor Networks: Problem

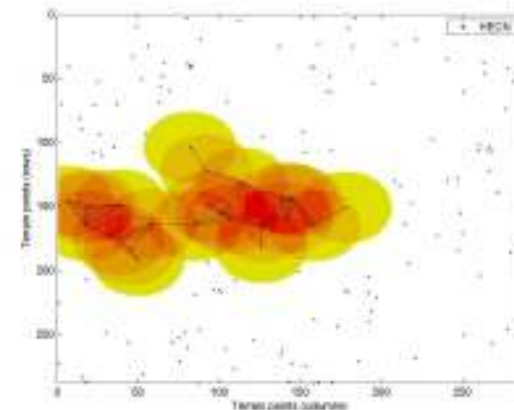
Task: Select geographical locations for sensor nodes
Locations may be chosen from a list, or freely

Objectives: Nodes must form a connected network
Maximize the coverage
Minimize the number of transmissions (energy)

Models: Node $\rightarrow (R_{\text{SENS}}, R_{\text{COMM}})$



Network \rightarrow





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

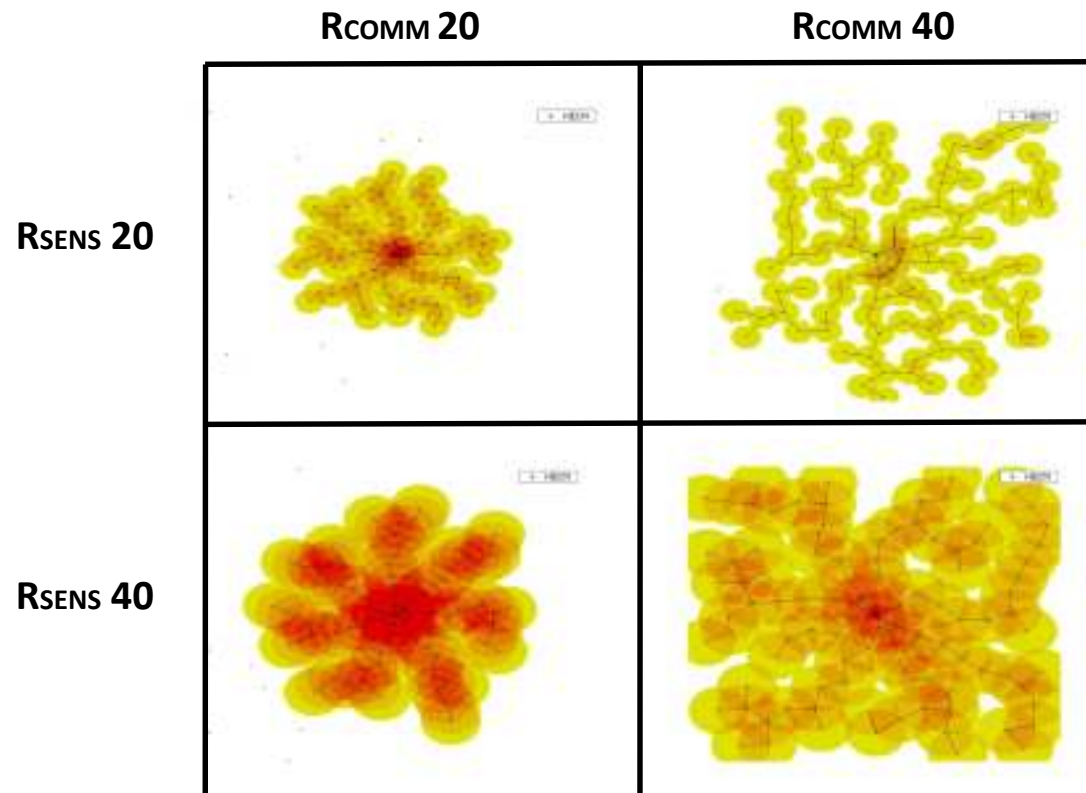
Software

Others

Wireless Sensor Networks: Results

Algorithms: **CHC** gets the best results in binary coded instances
SA gets the best results in real coded instances

Networks: Network layouts obtained for different sensor nodes





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

➔ Telecoms.

Bioinformatics

Software

Others

VANETS (I): Vehicular Ad-hoc Networks



<http://diricom.lcc.uma.es>
Information Dissemination



Website



Small Devices



Traffic Control Centre



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

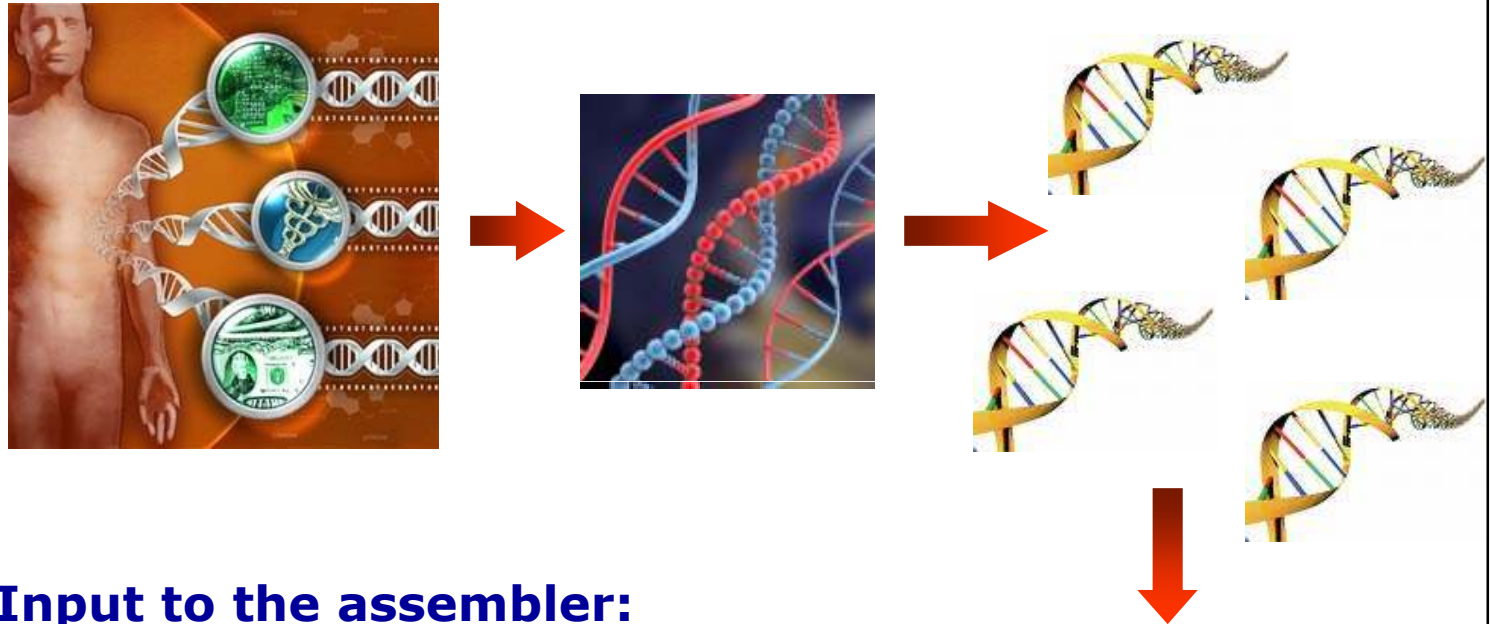
Telecoms.

➔ **Bioinformatics**

Software

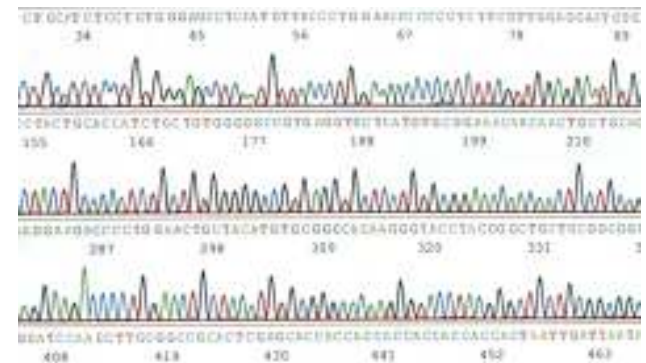
Others

Bioinformatics: Fragment Assembly Problem (I)



Input to the assembler:

CGGGATTTC AAGAAATG...
 GGATAGGTT AAAAATGC...
 CCAAAATGGAGGTCTTA...
 ...
 CATGGTACTTTTAGAGA...





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

➔ Bioinformatics

Software

Others

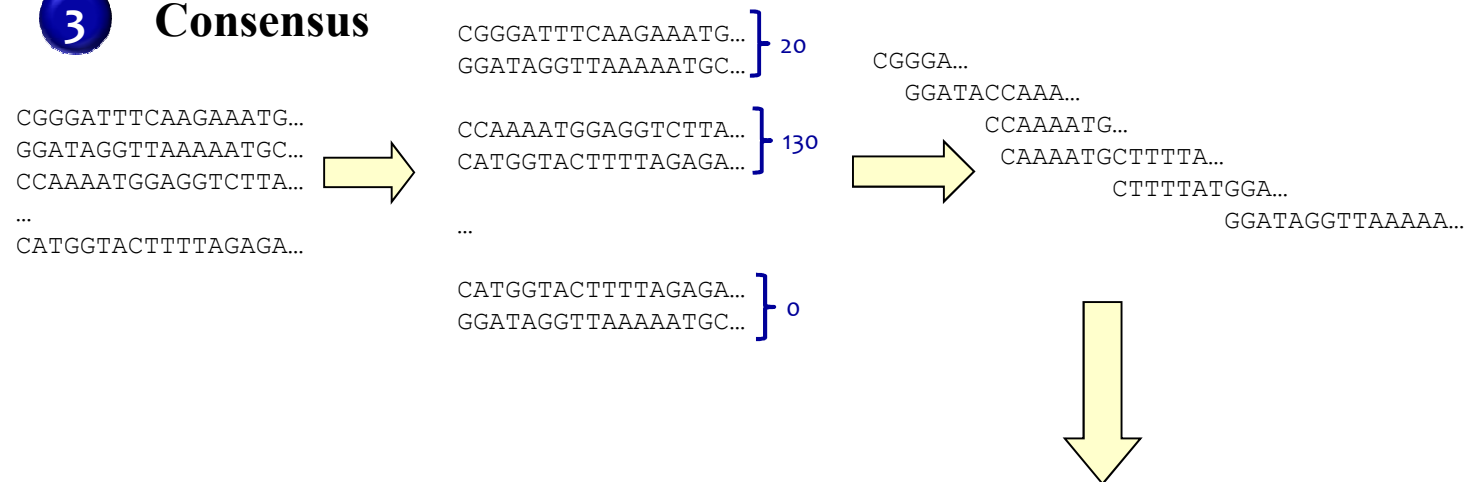
Bioinformatics: Fragment Assembly Problem (II)

The main phases performed by assemblers are:

1 **Overlap**

2 **Layout**

3 **Consensus**



Output of Assemblers: CCGGATAACCAAATGCTTTTATGGATAGGTTAAAAA ...

Our approaches:

- Parallel metaheuristics
- Specialized heuristics (PALS)

Our results:

- Able to solve very large sequences
- Large efficacy and efficiency



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

➔ Bioinformatics

Software

Others

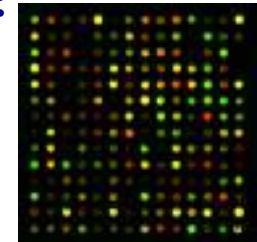
Bioinformatics: Gene Selection (I)

Problem:

Gene selection and cancer classification of DNA

Microarray, Feature selection definition:

$$F' = \operatorname{argmax}_{G \subset \Gamma} \{ \Theta(G) \}$$



Objectives:

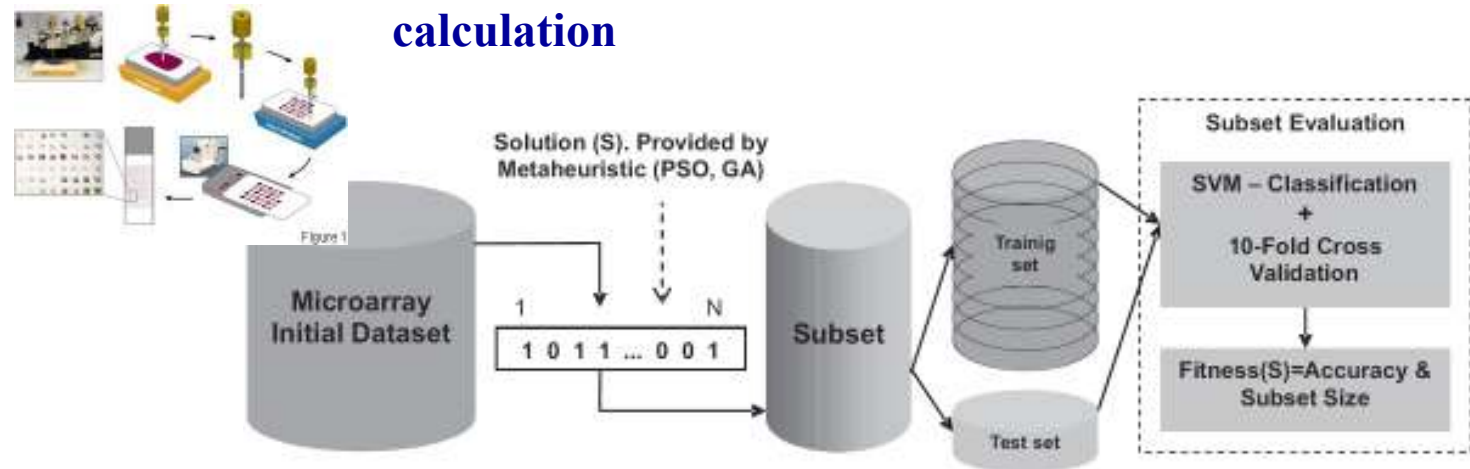
Maximize accuracy of prediction

Minimize the number of selected genes

Maximize ROC factors (sensitivity and specificity)

Phases:

Feature selection, training, validation, fitness calculation





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

➔ Bioinformatics

Software

Others

Bioinformatics: Gene Selection (II)

Fitness:

Monobjective: aggregative

$$(\alpha * 100 / \text{accuracy} + \beta * \# \text{features})$$

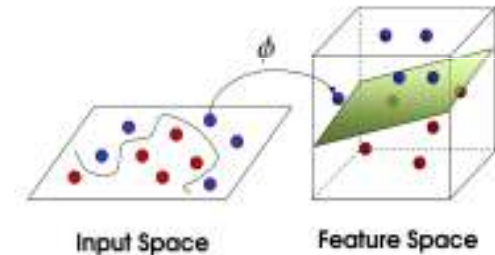
Multiobjective:

- 2 objs (*accuracy, #features*)
- 3 objs (*sensibility, specificity, #features*)

Classification:

2 Classifiers:

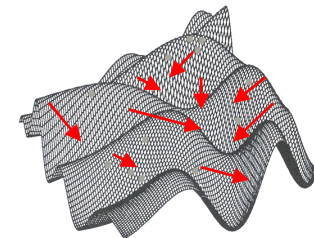
- Support Vector Machines
- *K*-Nearest Neighbors



Validation:

Cross-validation.

- Leave One Out CV
- 10-fold CV



Algorithms:

Metaheuristics

- Binary Geometric PSO for feature selection
- GA with SSO CF crossover for feature selection



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

➔ Bioinformatics

Software

Others

Bioinformatics: Gene Selection (III)

Instances: Available large scale datasets of well-known cancer DNA Microarrays: Leukemia AML-ALL, Colon, Prostate, Lung, Ovarian, Breast (e.g. breast 24481 genes and 78 patient samples)

Results: Comparison against other techniques in the literature

Dataset	PSO	GA	Huerta et al.	Juliusdotir et al.	Deb et al.	Guyon et al.	Yu et al.	Liu et al.	Shen et al.
<i>Leukemia</i>	99.33(3)	99.79(4)	100(25)	-	100(4)	100(2)	87.44(4)	-	-
<i>Breast</i>	90.64(4)	89.02(4)	-	-	-	-	79.38(67)	-	-
<i>Colon</i>	100(3)	100(4)	99.41(10)	94.12(37)	97(7)	98(4)	93.55(4)	85.48(-)	94(4)
<i>Lung</i>	100(4)	97.38(3)	-	-	-	-	98.34(6)	-	-
<i>Ovarian</i>	100(4)	100(3)	-	-	-	-	-	99.21(75)	-
<i>Prostate</i>	100(4)	100(4)	-	88.88(20)	-	-	-	-	-

Leukemia Gene Subsets

- By PSOSvm: K01383 at, U03056 at, J04130 s at
- By GASvm: L40379 at, S85963 at, U83192 at, Z49099 at



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

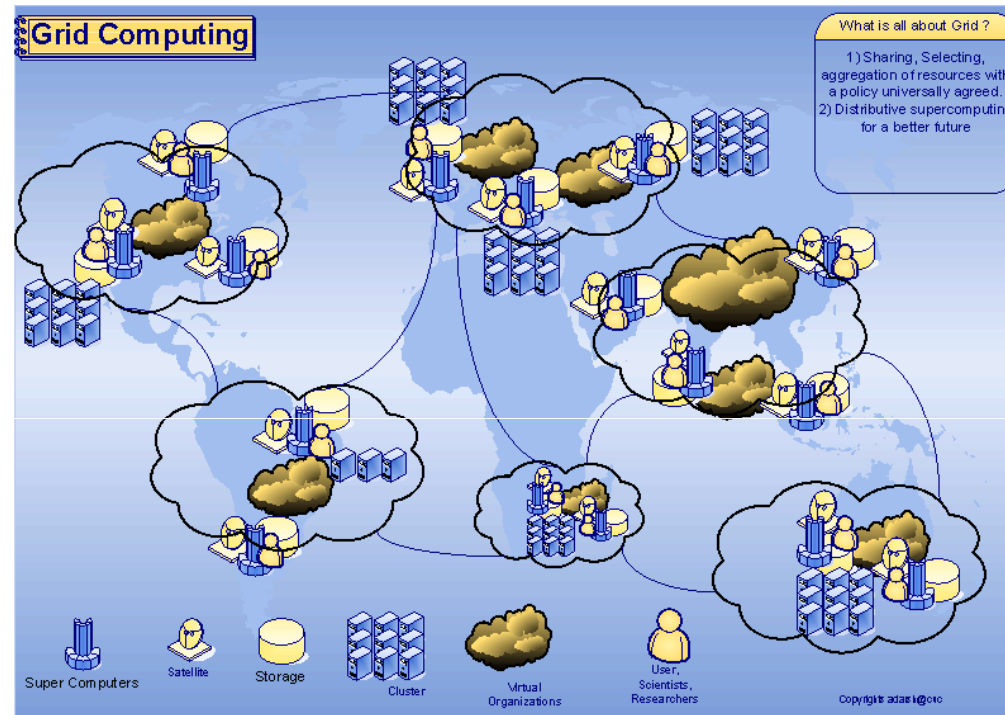
Telecoms.

➔ Bioinformatics

Software

Others

Bioinformatics and Grid Computing



At present:
In progress:
Next step:

using a stable grid with 300 computers at UMA
 stable grid of 600 computers at UMA
 connect to global grids in Europe



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

➔ Software

Others

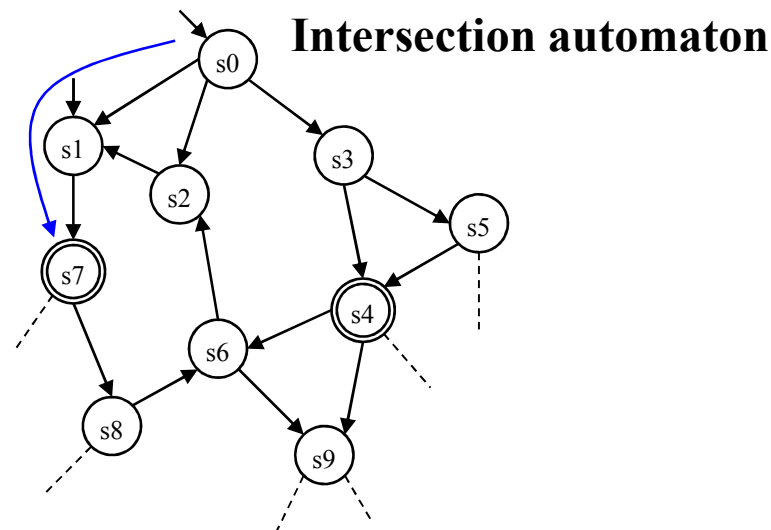
Finding Safety Property Violations in Concurrent Systems (I)

- **Objective:** find a counterexample for a safety property in a concurrent model
- **Safety properties** are those expressed by an LTL formula of the form:

$$f = \square p$$

where p is a past formula (with only past operators)

- **Finding one counterexample** \equiv finding one accepting state in the intersection Büchi automaton (graph exploration problem)



Safety Properties

Deadlocks

Invariants

Assertions

...



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

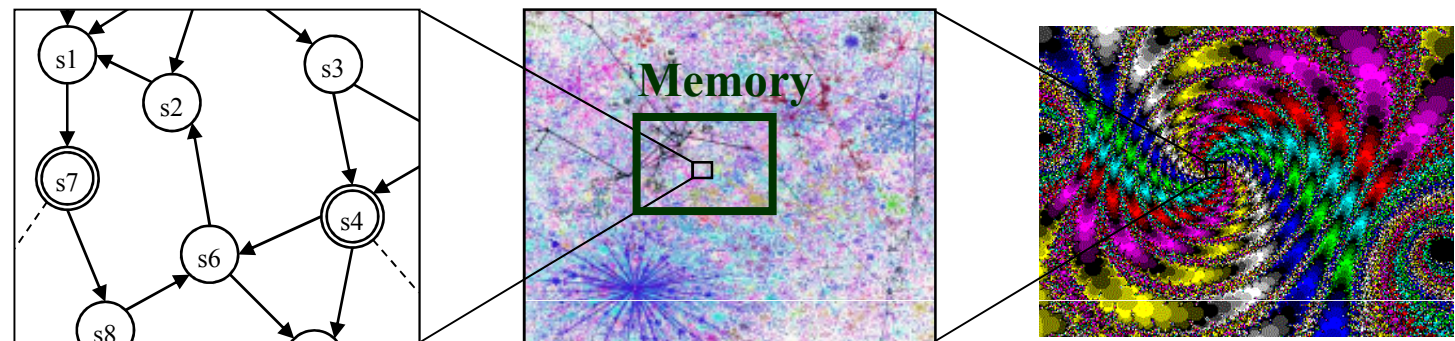
Bioinformatics

➔ Software

Others

Finding Safety Property Violations in Concurrent Systems (II)

- Number of states very large even for small models



- For example: Dijkstra Dining Philosophers

- n philosophers $\rightarrow 3^n$ states
- 20 philosophers $\rightarrow 1039$ GB for storing the states





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

➔ Software

Others

Finding Safety Property Violations in Concurrent Systems (III)

- **ACOhg** is a new Ant Colony Optimization model that can be applied to optimization problems with an unknown and/or very large construction graph

Who can really find errors?

Models	BFS	DFS	A*	BF	ACOhg
giop22		●	●	●	🐜
needham	●	●	●	●	🐜
phi16			●	●	🐜
pots	●	●	●	●	🐜
marriers4				●	🐜
marriers20					🐜

- **ACOhg** is a very robust algorithm for this problem and it outperforms traditional algorithm from the model checking domain



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

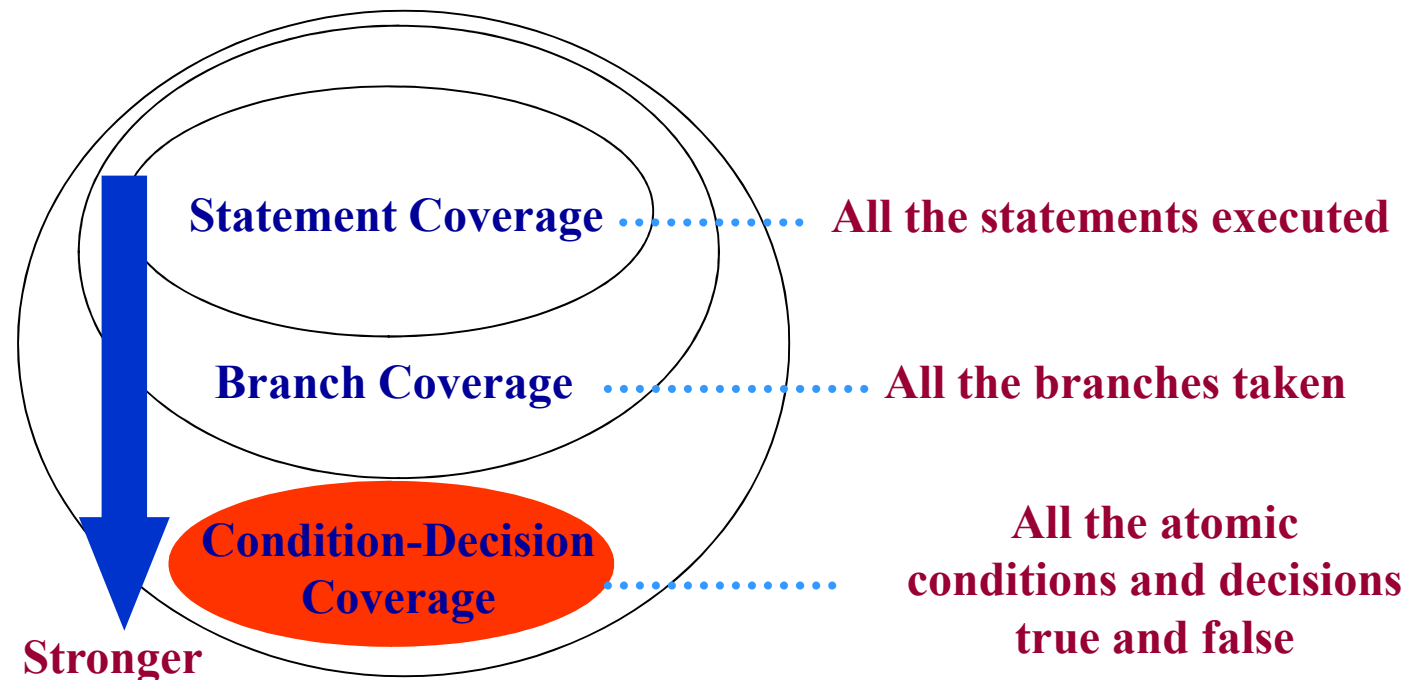
Bioinformatics

➔ Software

Others

Software Testing (I)

- **Objective:** propose a good set of test cases for a program
- The previous objective is too fuzzy, one concrete objective is: find a test case set fulfilling a test adequacy criterion
- Some examples of test adequacy criterion are:





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

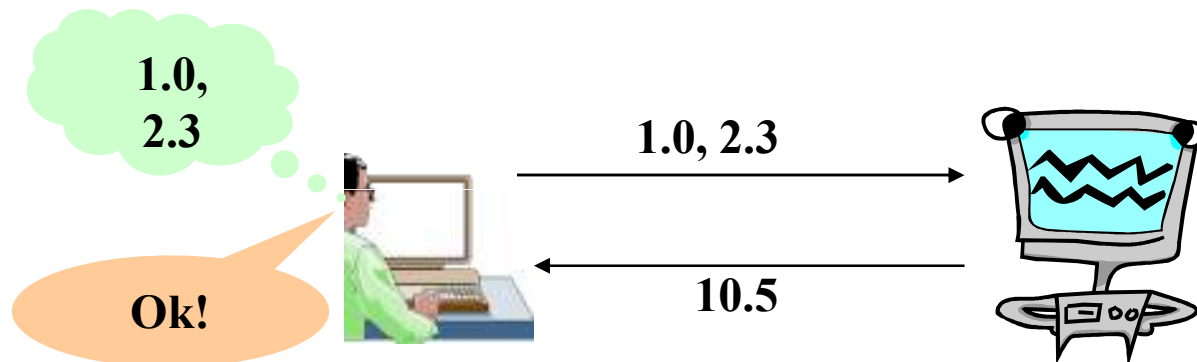
Bioinformatics

➔ Software

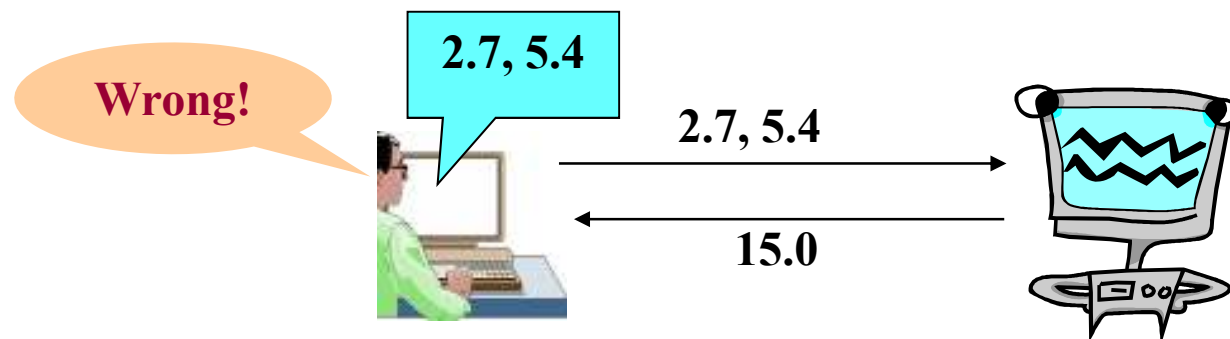
Others

Software Testing (II)

- After codification, software products require a **test phase**
- The objective is to **find errors** and to ensure **software correctness**
- Software companies dedicate **50%** of resources to this task



- We propose an automatic tool to **generate the input data for the test**





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

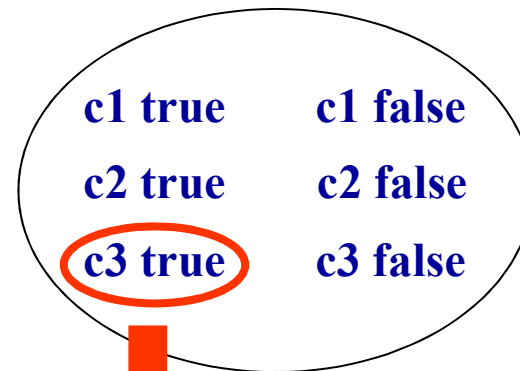
➔ Software

Others

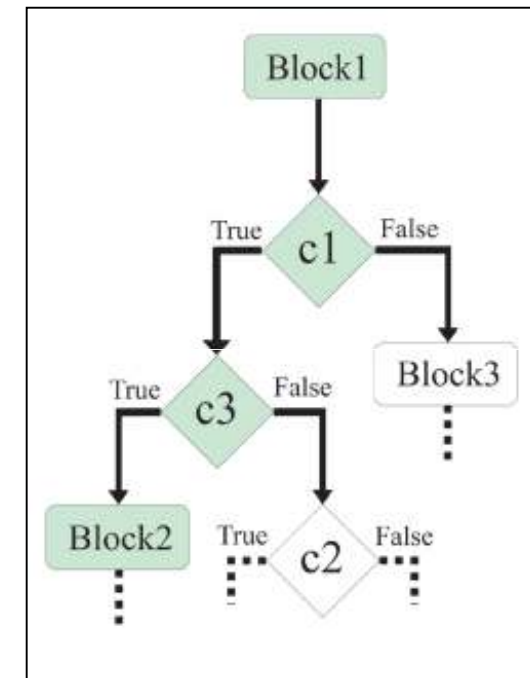
Software Testing (III)

- The global objective is broken down in small partial objectives

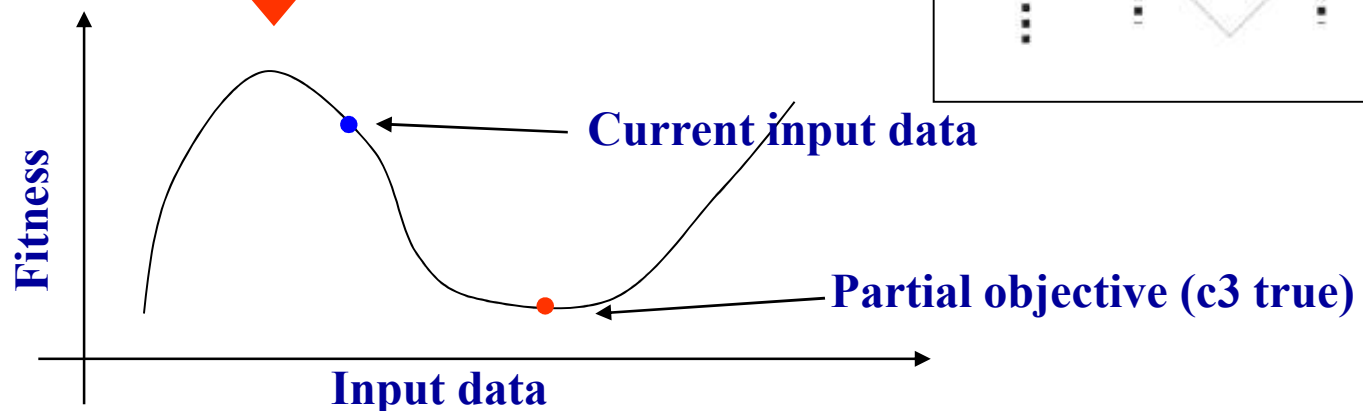
Six partial objectives



condition-
decision
coverage



Function minimization problem





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

➔ Software

Others

Software Testing (IV)

- Some results with PSO, ES, and GA (corrected condition coverage)

Programs	PSO		ES		GA	
	Cov.	Evals.	Cov.	Evals.	Cov.	Evals.
<i>triangle</i>	93.98	11295.77	99.84	2370.03	99.67	3209.47
<i>calday</i>	100.00	179.33	98.18	3166.47	90.91	75.03
<i>select</i>	88.89	380.13	83.33	13.27	83.33	83.20
<i>bessel</i>	97.56	116.90	97.56	350.63	97.56	533.03
<i>sa</i>	100.00	165.67	99.94	2337.30	96.72	176.63
<i>netflow</i>	97.77	4681.70	98.17	307.77	96.42	917.90

- PSO and ES have similar efficacy

- The coverage obtained by GA is always reached or outperformed by PSO or ES in all the cases



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

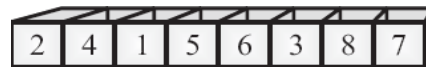
➔ Others

Others: Strip Packing (I)

- N rectangular pieces p_i with a height h_i and a width w_i and a rectangular container (the strip) with width W and unbounded height.
- **Objective:** To allocate all the pieces into the strip
 - without overlapping,
 - without rotating,
 - with their edges parallel to the edges of the strip,
 - **Bottom-up**, minimizing the height of the used strip.
- (Eq: To find a packing pattern that fulfils all these requirements)
- **Restriction:** three-stage guillotine patterns.
- **Scientific interest:** NP-hard problem.
- **Applications:** Paper, cloth, wood, and glass industries.

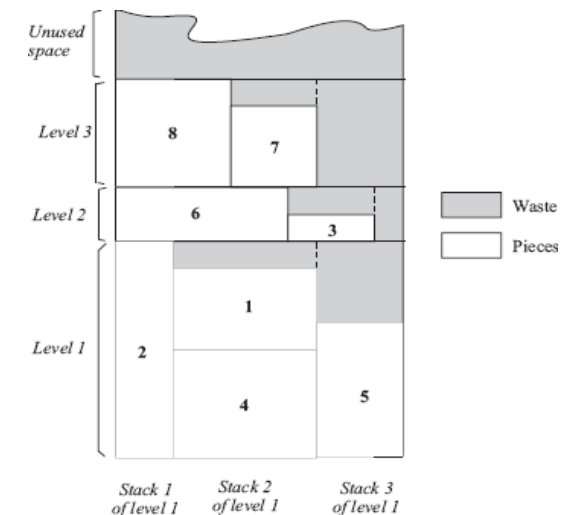
Representation

Chromosomes: sequences (permutations) of pieces which define the input for a layout algorithm.



Layout algorithm: a next fit heuristic that generates three-stage guillotine patterns.

Fitness function:
$$F(\pi) = \text{strip.length} - \frac{l.\text{waste}}{l.\text{area}}$$





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

➔ Others

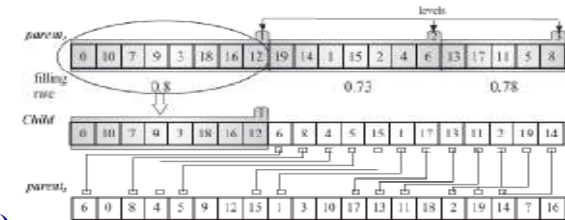
Others: Strip Packing (II)

Evolution Step

Best Inherited Level Recombination: Transmits the levels with the highest filling rate from one parent to the child.

Mutation: Best and Worst Stripe Exchange (BW_SE). Pieces of the best level are allocated in the first positions while the pieces of the worst level are assigned to the last positions.

Adjustment Operator: Applies a First Fit heuristic and the obtained layout is codified in a chromosome.



Initial Seeding

#	Rule Description	#	Rule Description
1	sorts pieces by decreasing width.	2	sorts pieces by increasing width.
3	sorts pieces by decreasing length.	4	sorts pieces by increasing length.
5	sorts pieces by decreasing area.	6	sorts pieces by increasing area.
#	Rule Description		
7	sorts pieces by alternating between decreasing width and height.		
8	sorts pieces by alternating between decreasing width and increasing height.		
9	sorts pieces by alternating between increasing width and height.		
10	sorts pieces by alternating between increasing width and decreasing height.		
11	the pieces are reorganized following the BFDH heuristic.		
12	the pieces are reorganized following the FFDH heuristic.		
13	The packing pattern remains without modifications, so here the rule preserves the original piece position (random generation).		



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

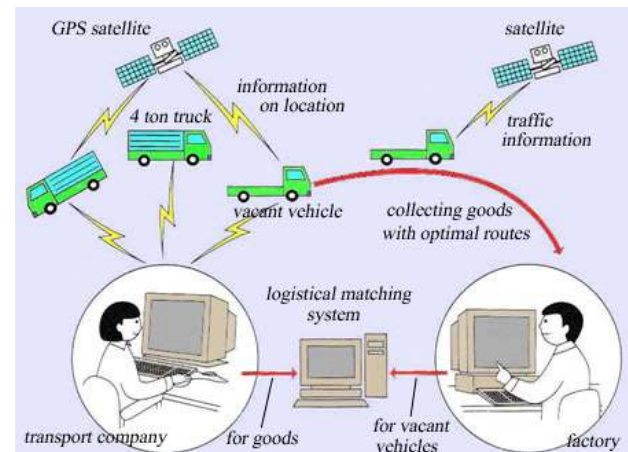
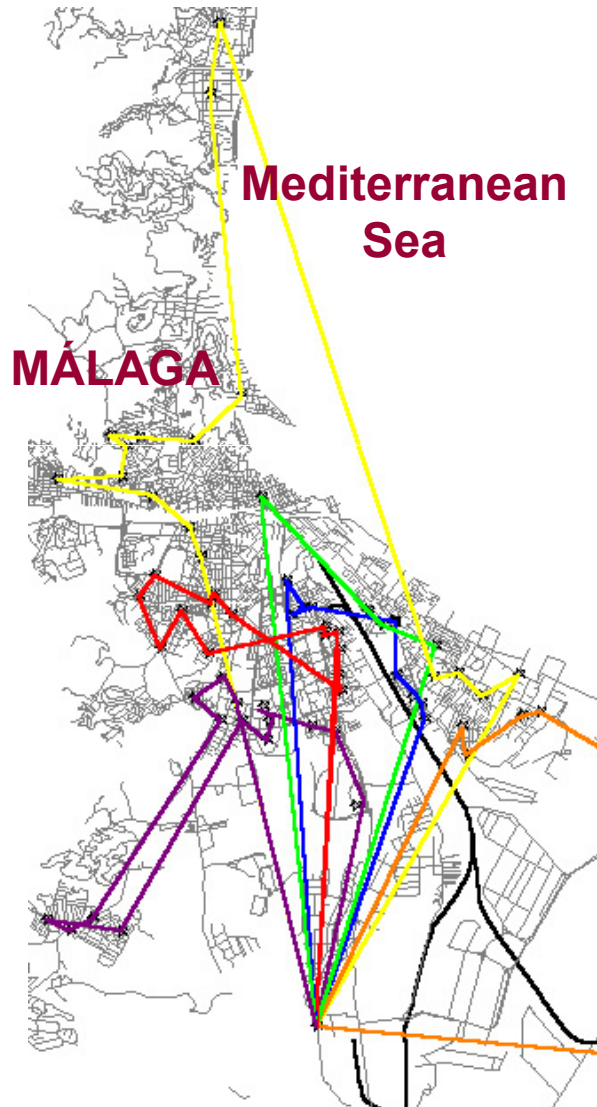
Telecoms.

Bioinformatics

Software

➔ Others

Others: Logistics





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

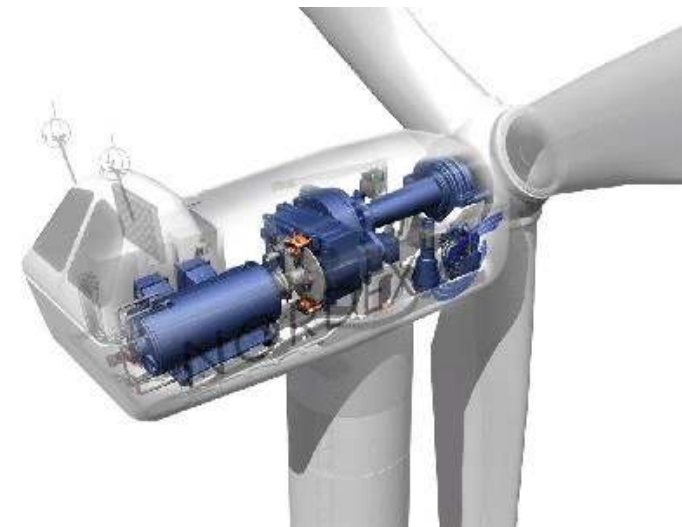
➔ **Others**

Others: Dynamic Problems

Optimal location of aerogenerators



Wind farm design



Aerogenerator design



Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

➔ **Others**

Others: Dynamic Problems

DYNAMIC OPTIMIZATION

Traffic Light
Controllers



Elevator
Systems



Gas Engine
Control





Index

New Models

Parallel MetaH

Hybrid MetaH

Multiobjective

Telecoms.

Bioinformatics

Software

➔ Others

End of Presentation

Málaga

<http://neo.lcc.uma.es>

