

# Evolutionary Optimization in Electromagnetics

COST IC 1407 Workshop,  
Bratislava, 5.4.2017

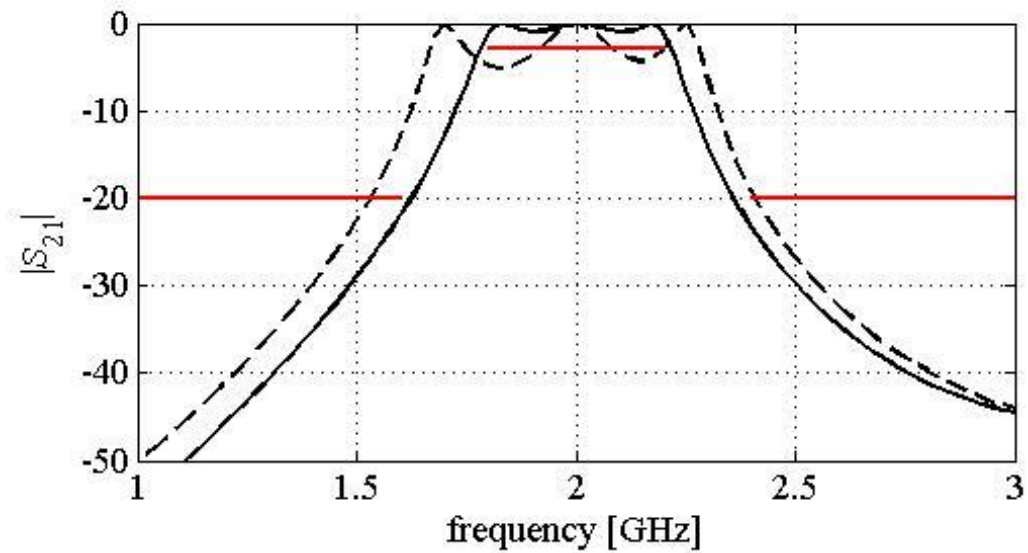
# Outline

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- Optimization fundamentals
- Local vs. global methods
- Evolutionary algorithms
- Genetic Algorithms
- Particle Swarm Optimization
- Multi-objective optimization
- Examples in FOPS

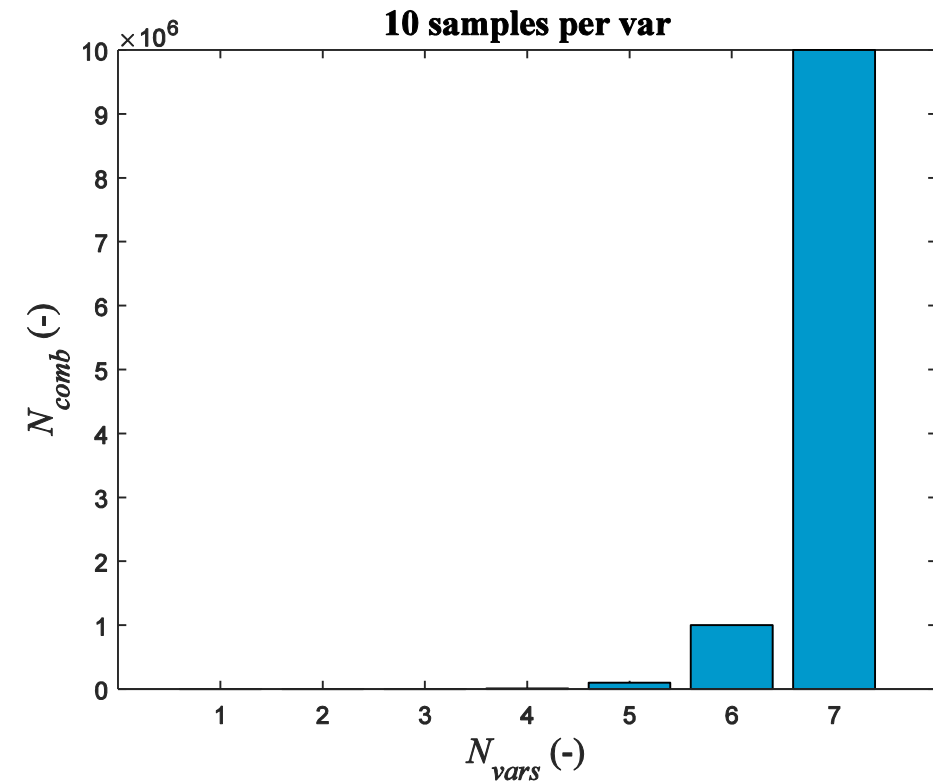
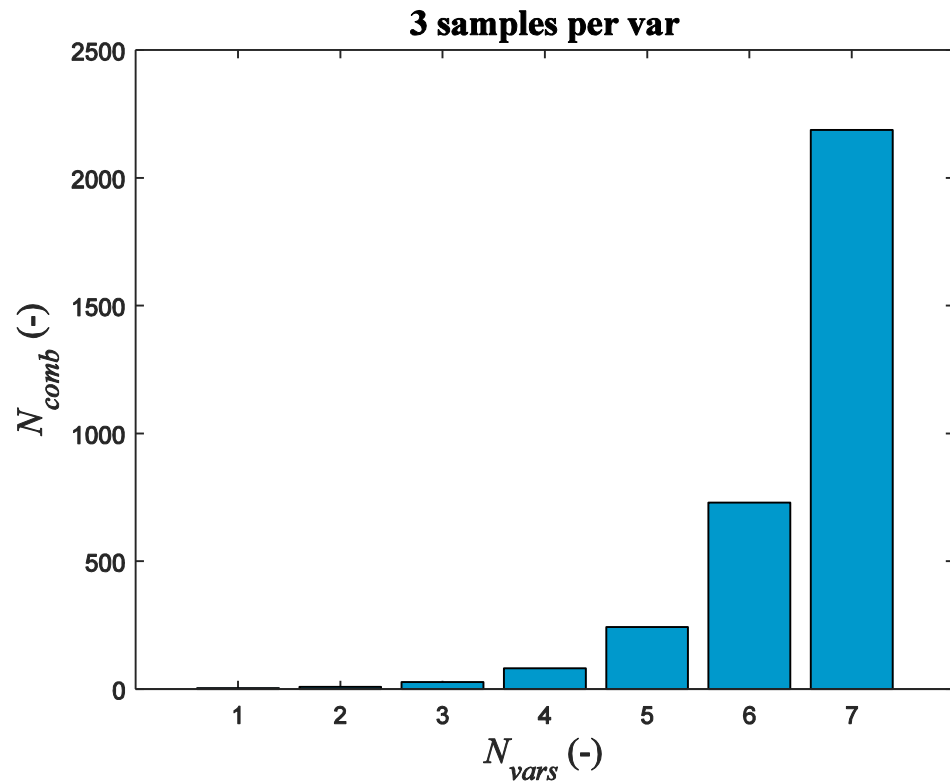
# Motivation

*Always try to do the best...*



# Motivation

*Is parametric analysis the way?*



# Optimization

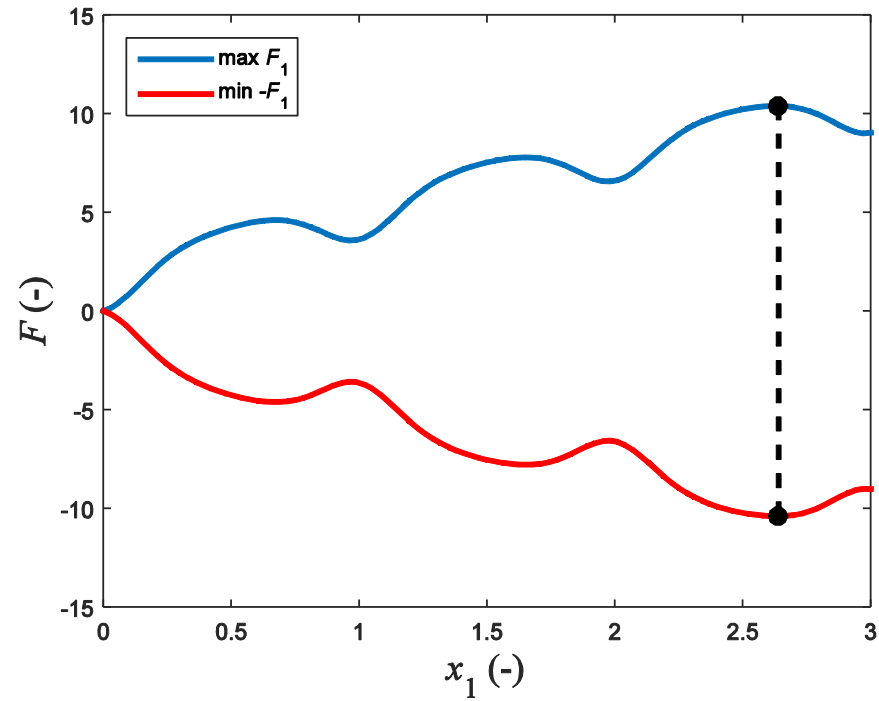
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- Choice of the best variant from available options
- In mathematics:

$$\begin{aligned} & \textit{Minimize} \quad F_m(\mathbf{x}), \quad m = 1, 2, \dots, M, \\ & \textit{subject to} \quad g_j(\mathbf{x}) \geq 0, \quad j = 1, 2, \dots, J, \\ & \quad \quad \quad x_{n,\min} \leq x_n \leq x_{n,\max}, \quad n = 1, 2, \dots, N. \end{aligned}$$

# Min vs. Max

$$\text{Min } F_m(\mathbf{x}) \approx \text{Max } -F_m(\mathbf{x})$$



# Objective formulation

## Problem:

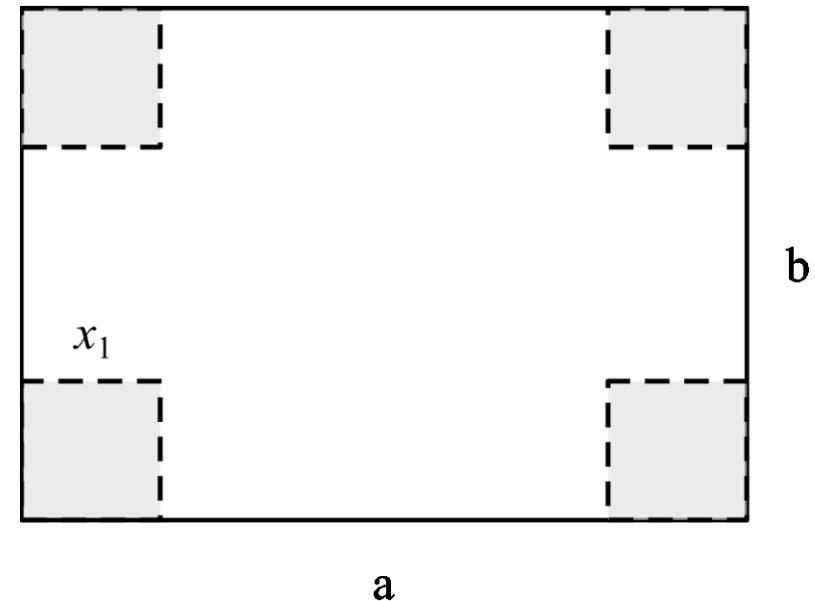
Having sheet of paper with dimensions  $a$  and  $b$ , what is the size of squares to be cut from the corners of the sheet to build a box with the highest volume?

Decision space:

$$\mathbf{x} = [x_1]$$

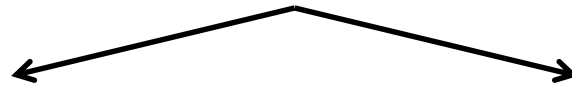
Objective space

$$f_1 = 4x_1^3 - (2a + 2b)x_1^2 + abx_1$$



# Optimization taxonomy

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**SOO (Single-Objective)**

**Single variable**

**Global**

**Continuous dec. space**

**Unconstrained**

**Static**

**MOO (Multi-Objective)**

**Multiple variables**

**Local**

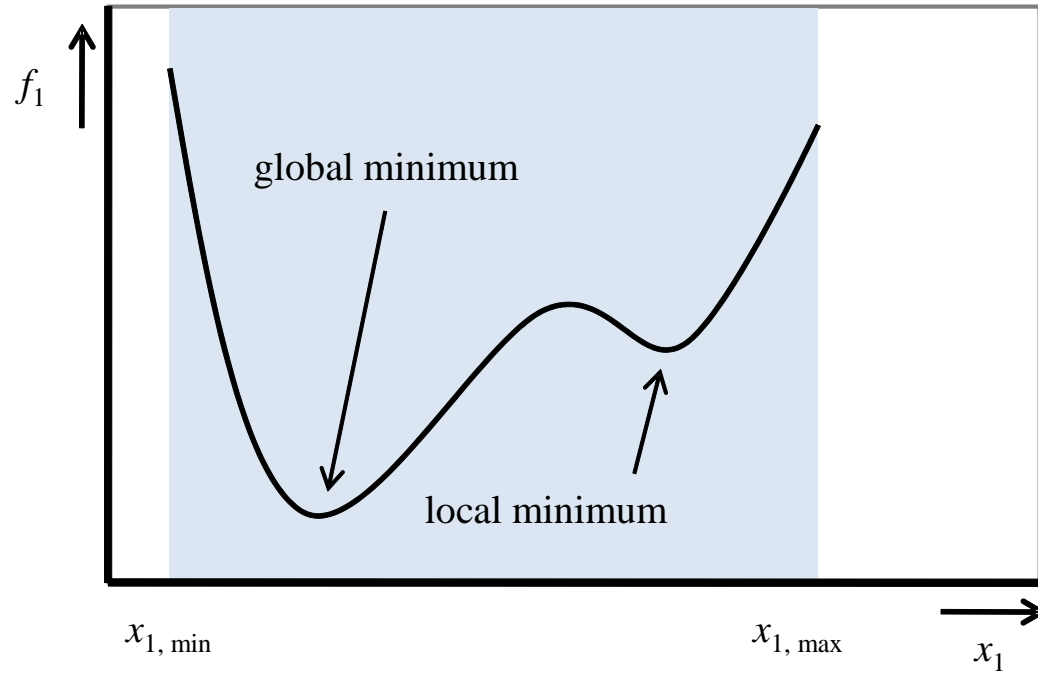
**Discrete dec. space**

**Constrained**

**Dynamic**



# Global vs. local minimum



# Global vs. Local

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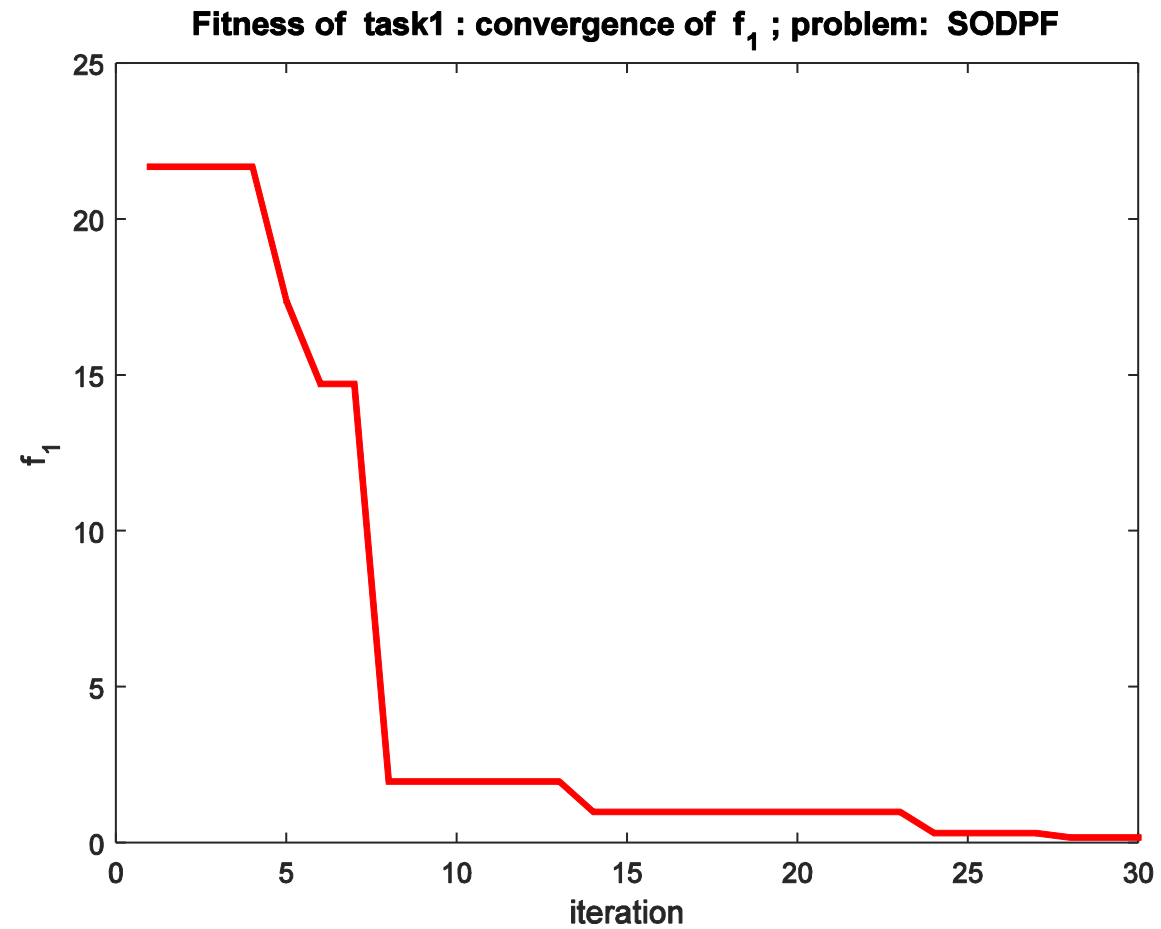
- local methods are faster and more efficient, but global are more robust, no dependence on initial guess
- derivatives can not be evaluated,
- objective functions cannot be formulated in closed form (e.g. use of solver output),
- initial guess is too far,
- user is lazy!

# Evolutionary methods

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- random start – choice of the initial guess is not so important
- agent (individual, ...) updates its position within the decision space
- agents are able to escape from local minimum (maximum)
- stochastic – different runs different results (at least development)

# Convergence plot



# Genetic algorithms

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- Holland 1962
- Inspired by Darwin – only good properties of genome are maintained for next generations
- **GAs work with discrete decision space!!!**

# GA - taxonomy

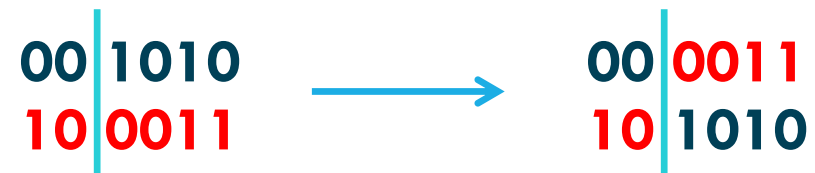
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- Gene – variable coded in binary form
- Gene length – number of bits used for decoding
- Chromosome – genes from all variables
- Generation – multiple chromosomes
- Decimation – reduction of the worst Chromosomes in Generation
- Mating pool – set of Chromosomes selected for reproduction
- Crossover – combination of two Chromosomes
- Mutation – bit change in one Chromosome

# GA - reproduction

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- **Crossover:**



- **Mutation:**



# GA - reproduction

- **Decimation:**

$g_1$	$g_2$	$g_3$	$f$
110	0101	00010	0 , 15
101	1101	00101	0 , 36
011	1011	00110	1 , 53
100	1010	10001	6 , 27
011	0011	11010	11 , 83
101	1011	10010	13 , 21
100	0101	00101	20 , 89
011	0101	10101	56 , 12

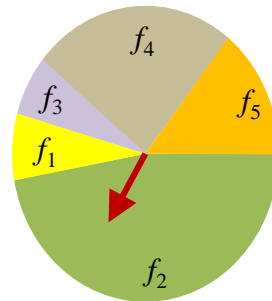
- **Elitism:**

- Solution with the best value of objective function is automatically considered also for the next generation.



# GA - reproduction

- **Roulette selection:**



#	$f(-)$	$1/f(-)$	%
1	6.82	0.15	7.80
2	1.11	0.90	47.94
3	8.48	0.12	6.28
4	2.57	0.39	20.71
5	3.08	0.32	17.28
total	22.06	1.88	100.00

- **Tournament selection:**

$f$	Chrom.	Round	Mating pool
1.25	C1	C1 vs. C2	C1
7.37	C2	C4 vs. C2	C4
2.42	C3	C1 vs. C3	C1
6.12	C4	C4 vs. C3	C3

# Particle Swarm Optimization

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- 1995 - Eberhart and Kennedy
- cooperation of swarm over the meadow
- cognitive learning: personal experience of agent
- social learning: experience of the whole swarm

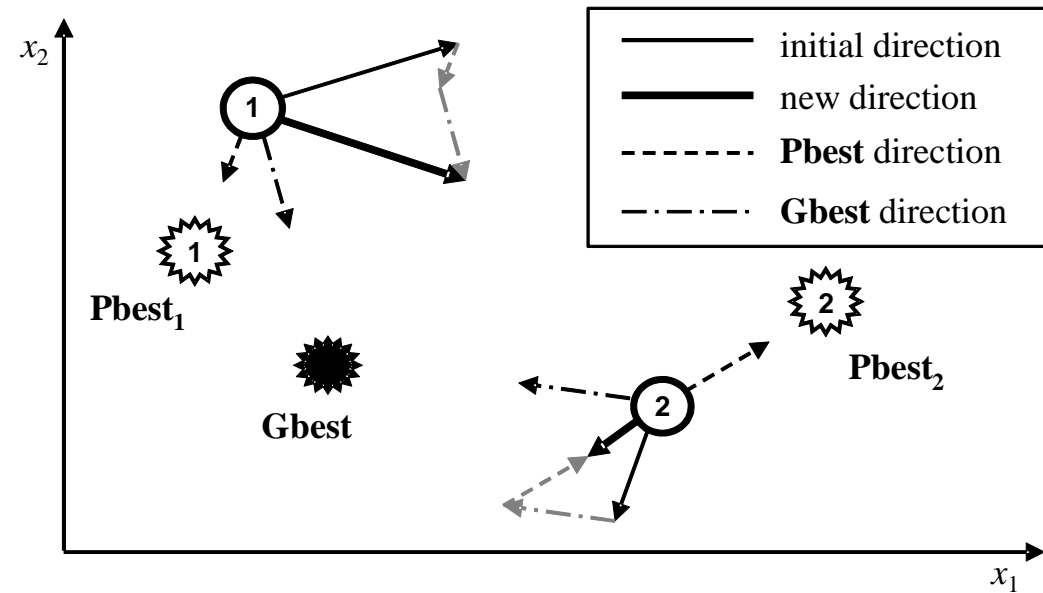
# Swarm update

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$$\mathbf{x}_q(i) = \mathbf{x}_q(i-1) + \mathbf{v}_q(i-1)$$

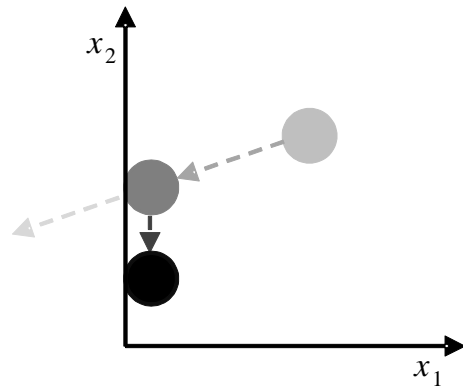
$$\begin{aligned} \mathbf{v}_q(i) = & w \cdot \mathbf{v}_q(i-1) + c_1 \cdot \mathbf{rnd}_q \left[ \mathbf{Pbest}_q(i-1) - \mathbf{x}_q(i-1) \right] \\ & + c_2 \cdot \mathbf{rnd}_q \left[ \mathbf{Gbest}(i-1) - \mathbf{x}_q(i-1) \right] \end{aligned}$$

# Swarm update

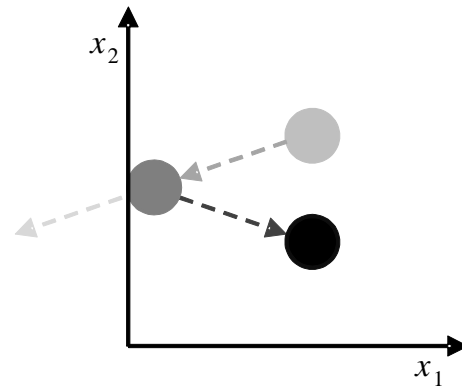


# Wall types

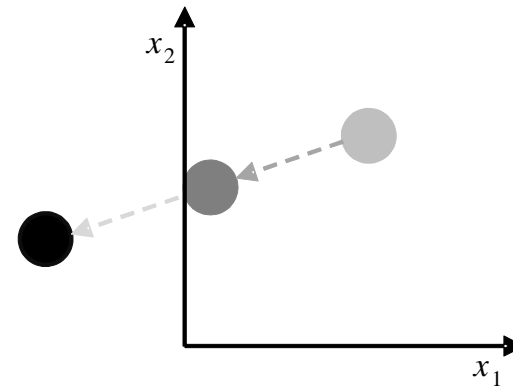
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Absorbing

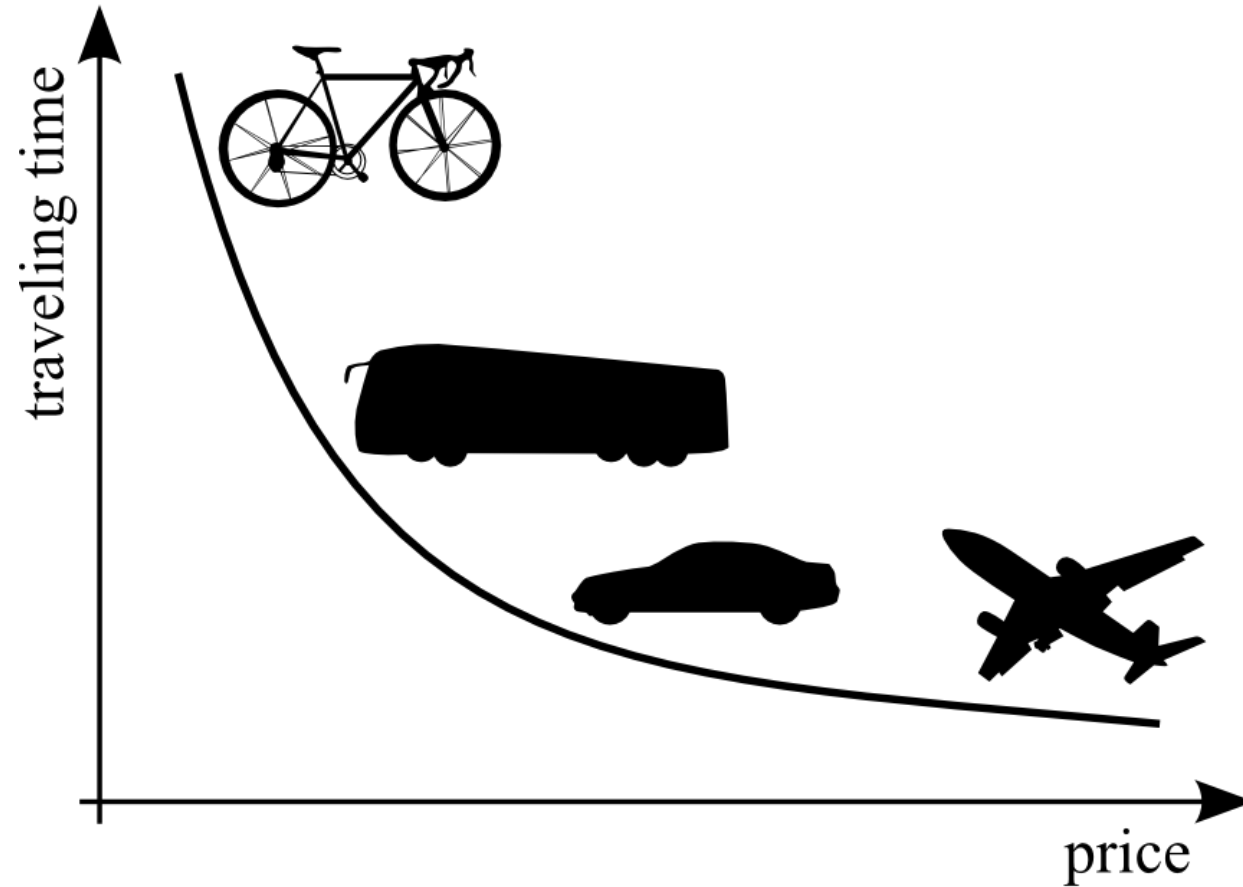


Reflecting



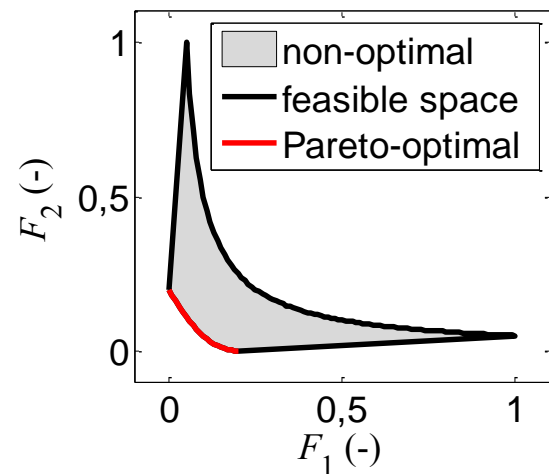
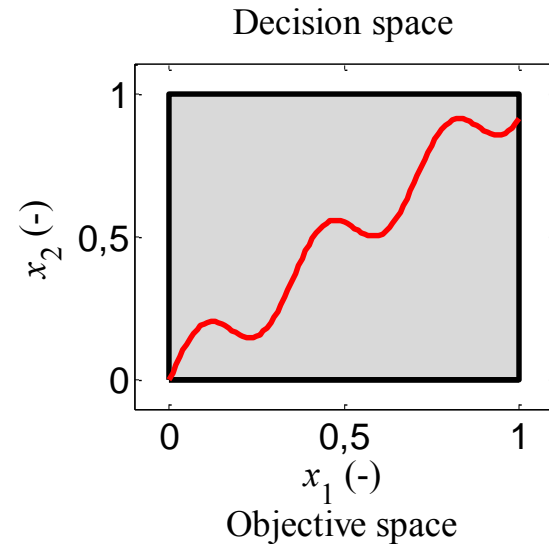
Invisible

# Multi-objective optimization



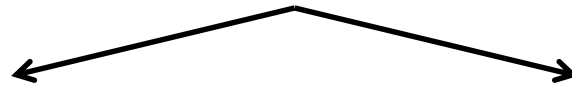
# Pareto front

- Trade-off solution
- Decision space vs. Objective space
- Two goals – accuracy vs. Distribution
- Choice of final solution
- Pareto front shapes



# Multi-objective strategies

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## **Aggregating methods**

Transform to SOOP

Single solution

How to choose  $w$ ?

## **MO Optimization**

More complex routine

Set of Pareto optimal

Extra information

Decision making

$$F = \sum_{m=1}^M w_m F_m, \sum_{m=1}^M w_m = 1.$$



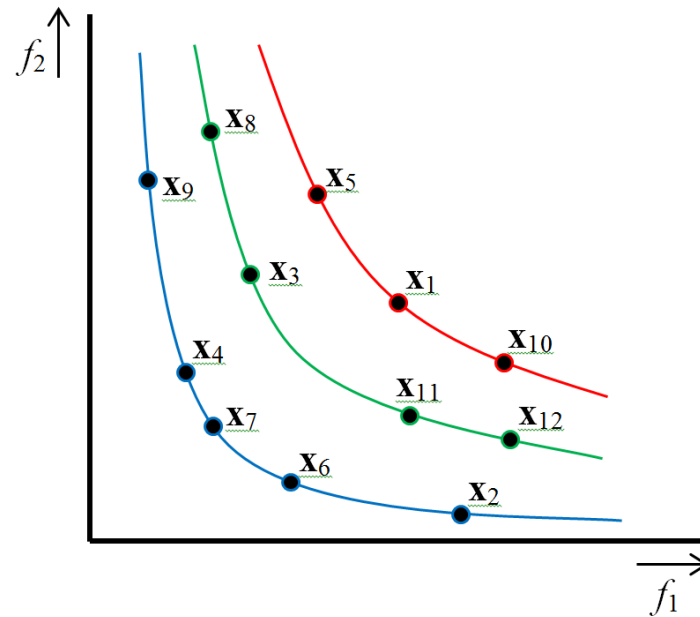
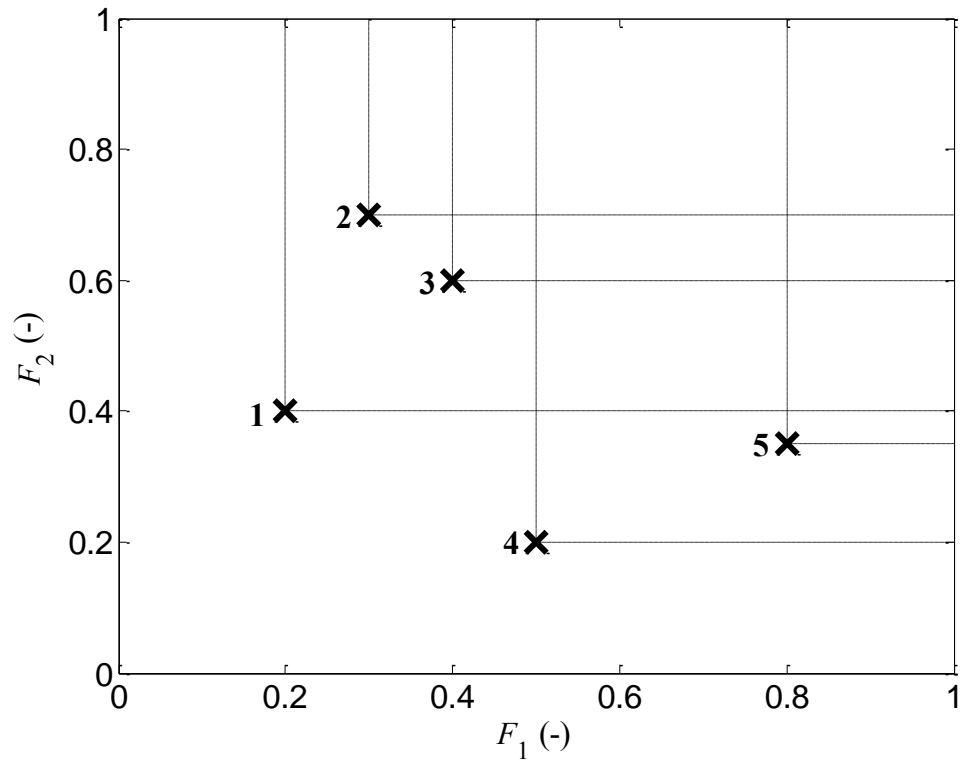
# Dominance concept

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*Solution  $\mathbf{x}_1$  is said to dominate the other solution  $\mathbf{x}_2$ , if both conditions 1 and 2 are true:*

- 1. Solution  $\mathbf{x}_1$  is no worse than  $\mathbf{x}_2$  in all objectives.*
- 2. Solution  $\mathbf{x}_1$  is strictly better than  $\mathbf{x}_2$  in at least one objective.*

# Dominance concept



Set Q

- $x_9$
- $x_4$
- $x_7$
- $x_6$
- $x_2$
- $x_8$
- $x_3$
- $x_{11}$
- $x_{12}$
- $x_5$
- $x_1$
- $x_{10}$

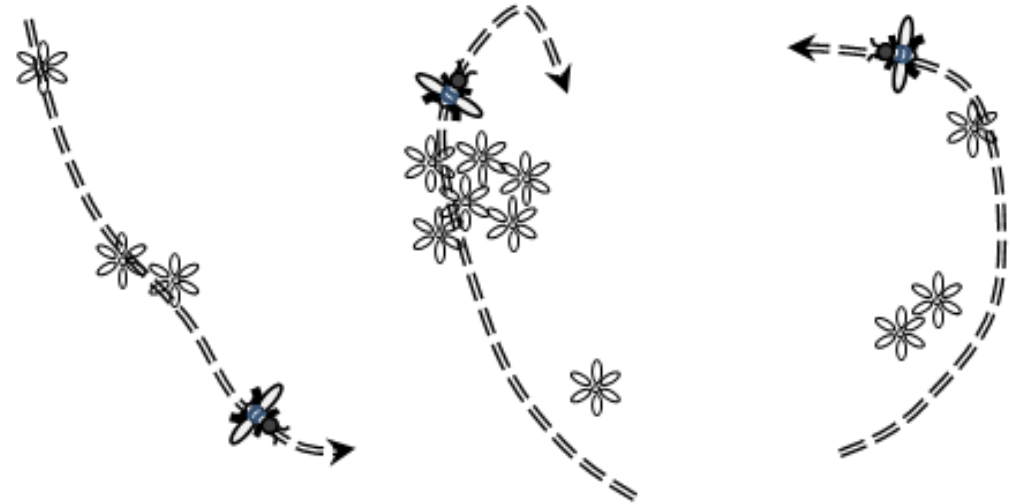
best

worst

# MOPSO

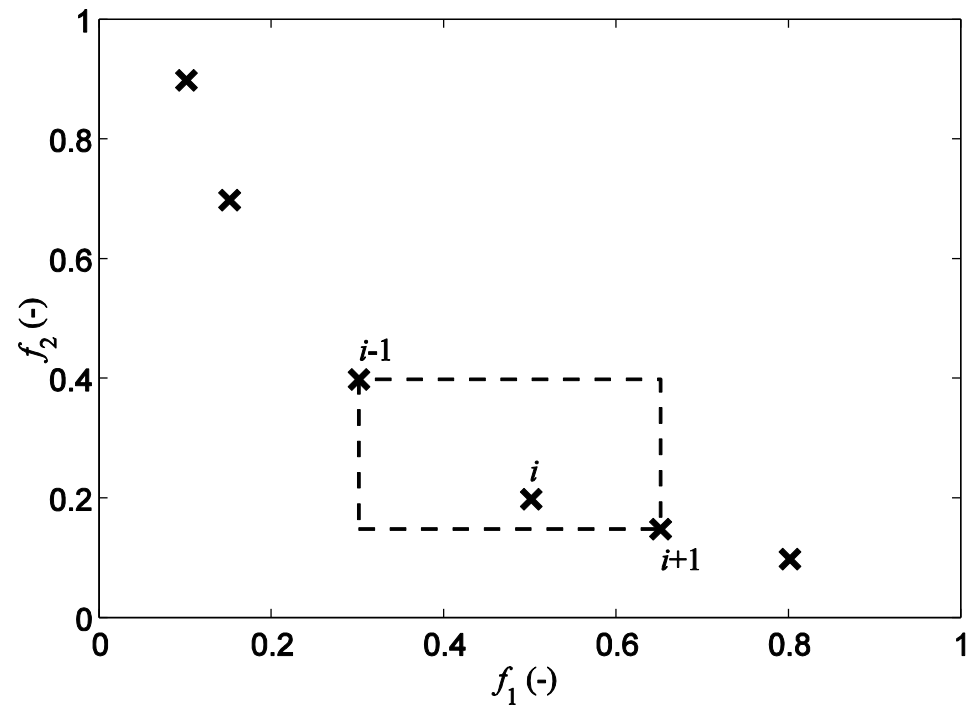
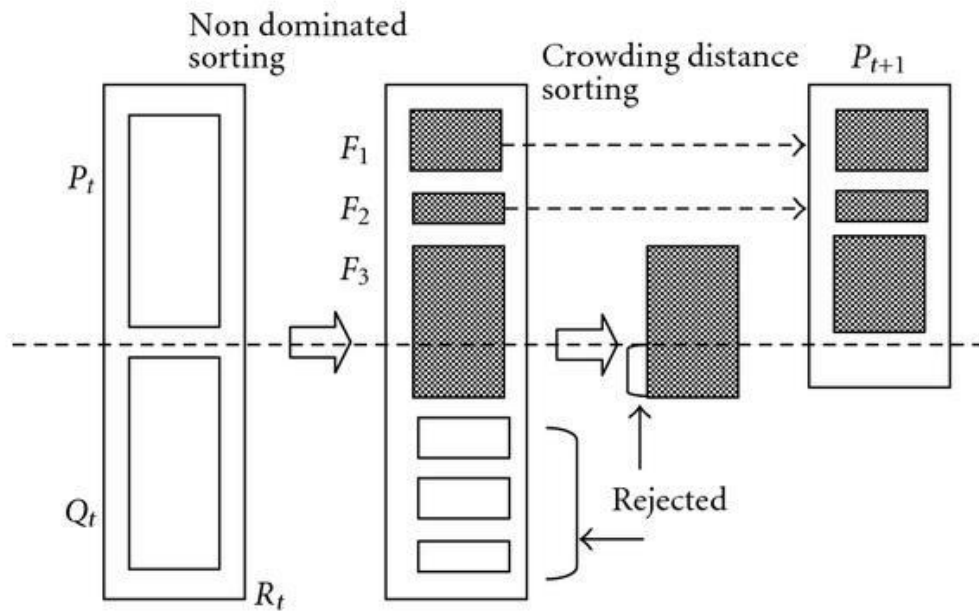
Nanbo, J., Rahmat-Samii, Y. "Advances in Particle Swarm Optimization for Antenna Designs: Real-Number, Binary, Single-Objective and Multiobjective Implementations," IEEE Transactions on Ant. Propag., vol. 55, no. 3, pp. 556-567, 2007.

- **gbest** – closest solution from external archive
- **pbest** – first non-dominated solution found by particle



# NSGA-II

Deb, K., Pratap, A., Agarwal, S., Meyarivan, T. "A fast and elitist multiobjective genetic algorithm: NSGA-II," IEEE Transactions on Evol. Comput., vol. 6, no. 2, pp. 182-197, 2002.



# FOPS

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- Fast Optimization ProcedureS
- <http://antennatoolbox.com/fops-about.php>
- single- and multi-objective codes
- chains from individual methods
- local methods: steepest descent, Newton method
- global methods: Nelder Mead, GA, PSO, DE, SOMA ...

# Feeding Point

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## Problem:

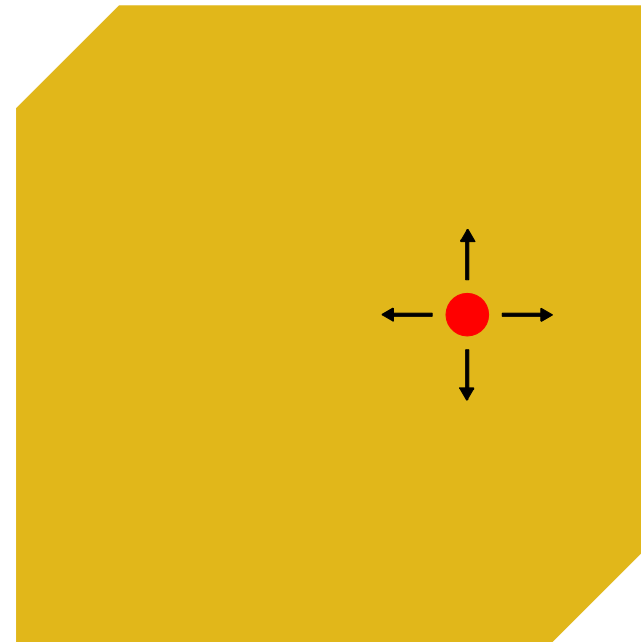
Find proper point on antenna for coaxial port.

Decision space:

$$\mathbf{x} = [x_1, x_2]$$

Objective space

$$f = \left( \operatorname{Re}\{Z_{inp}(\mathbf{x})\} - 50 \right)^2 + \left( \operatorname{Im}\{Z_{inp}(\mathbf{x})\} - 0 \right)^2$$



# Feeding Point

Algorithm settings

Load algorithm settings from file

Number of iterations:  
30

Number of agents:  
20

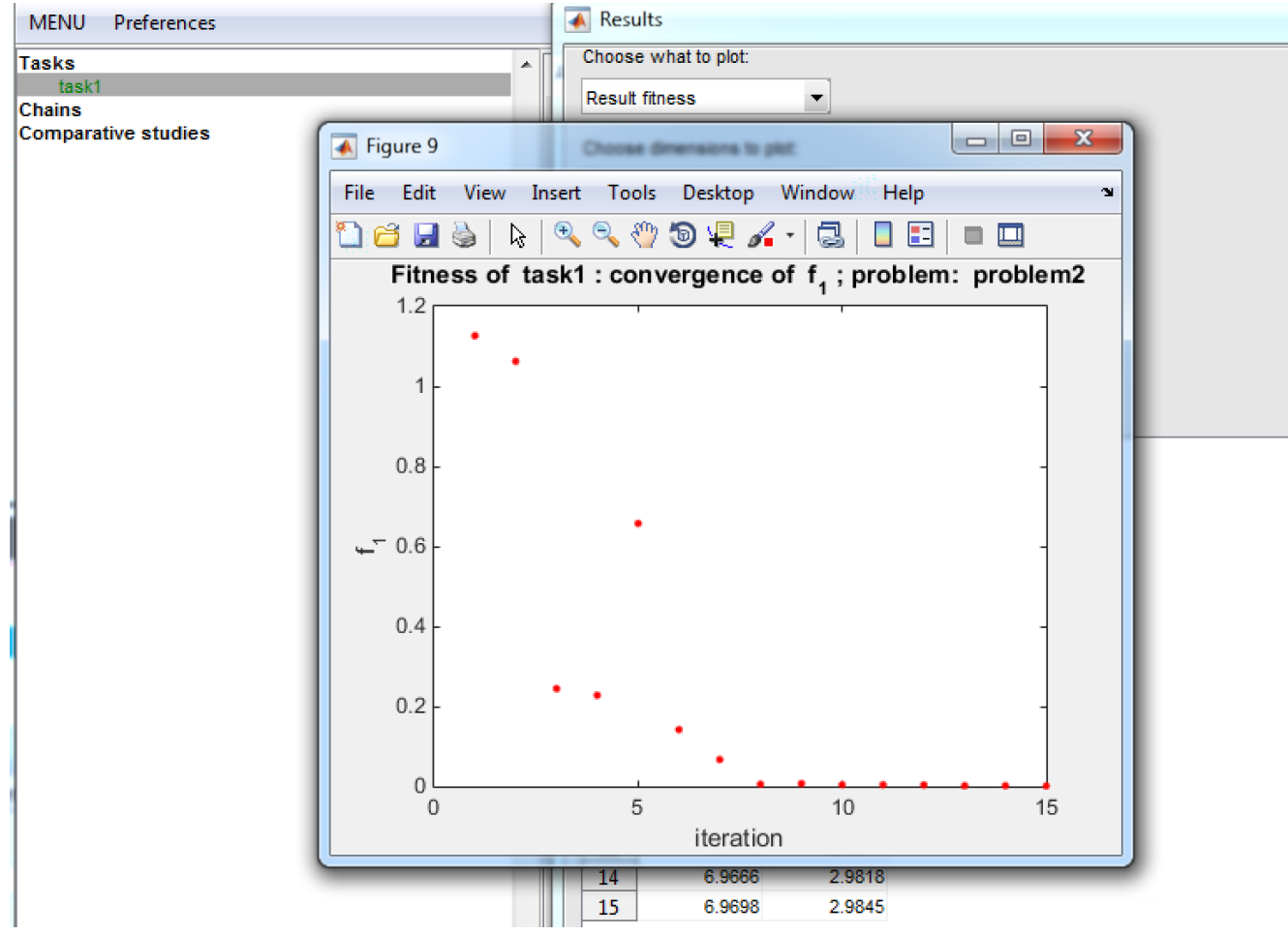
Inertia weight (W):  
0.6 0.4

Cognitive learning factor (C1):  
1.5

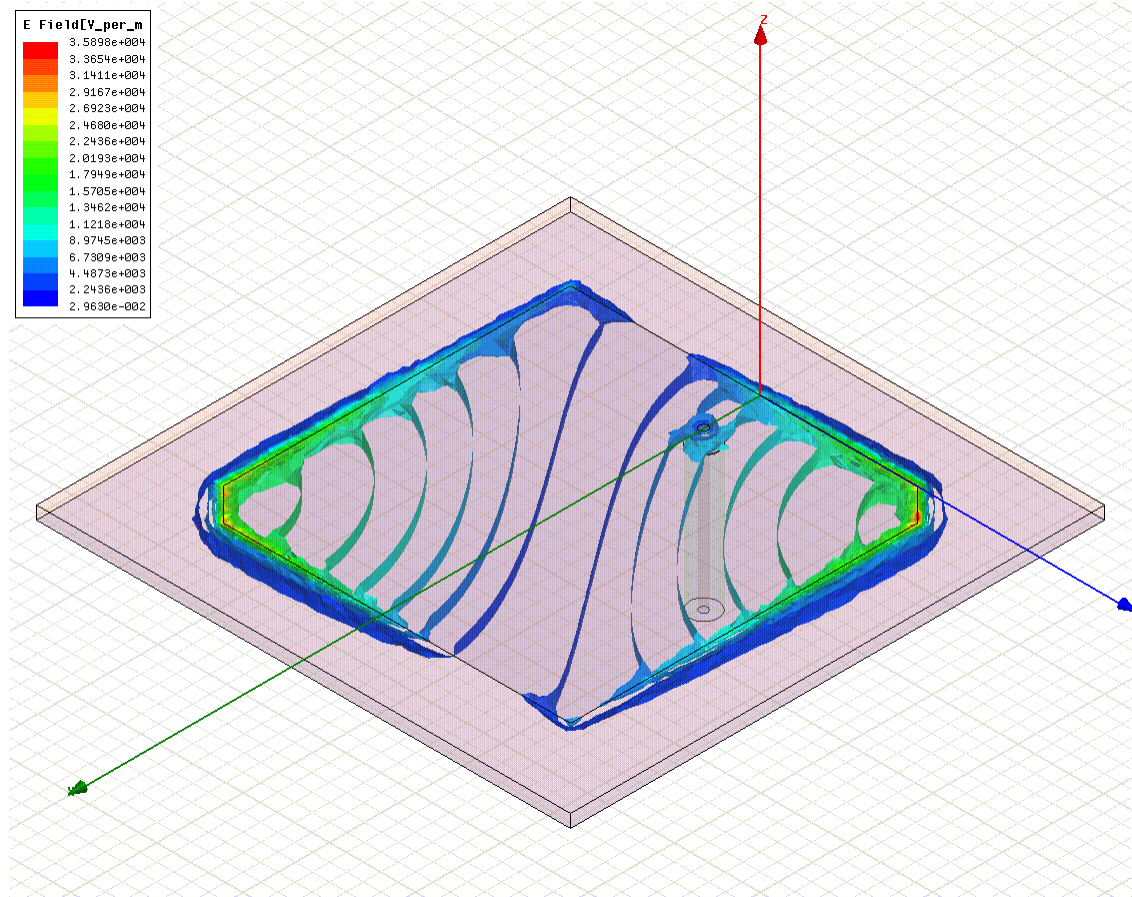
Social learning factor (C2):  
1.5

Boundary type:  
reflecting

Confirm settings



# Feeding Point





# Filter design

## Problem:

Find proper material and width of layers to design band pass filter.

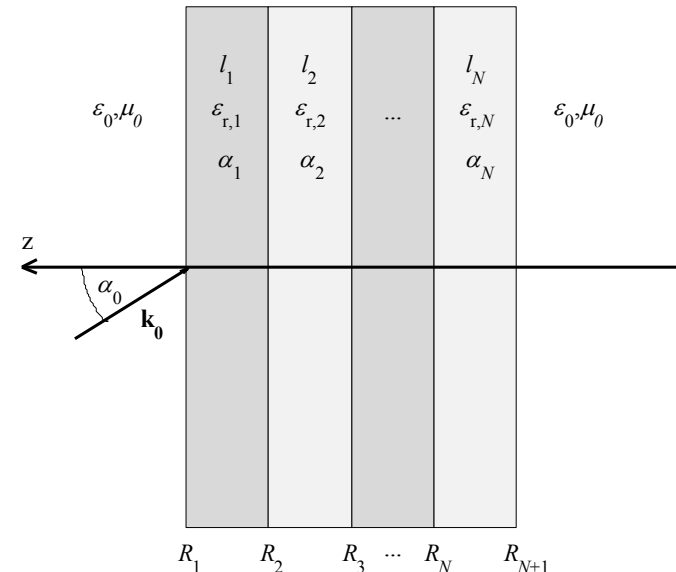
## Decision space:

$$\mathbf{x} = [l_1, \varepsilon_1, l_2, \varepsilon_2, \dots, l_N, \varepsilon_N]$$

## Objective space

$$F_1 = \frac{1}{P} \sum_{p=1}^P \left[ |R_{1,TE}(f_p)|^2 + |R_{1,TM}(f_p)|^2 \right]$$

$$F_2 = \frac{1}{S} \sum_{s=1}^S \left[ 2 - |R_{1,TE}(f_s)|^2 - |R_{1,TM}(f_s)|^2 \right]$$



# Filter design

The image shows a software interface for configuring optimization tasks. The main window has a menu bar with "MENU" and "Preferences". On the left, there is a sidebar with "Tasks", "Chains", and "Comparative studies". The main area has three buttons: "Add task", "Add chain", and "Add comparative study".

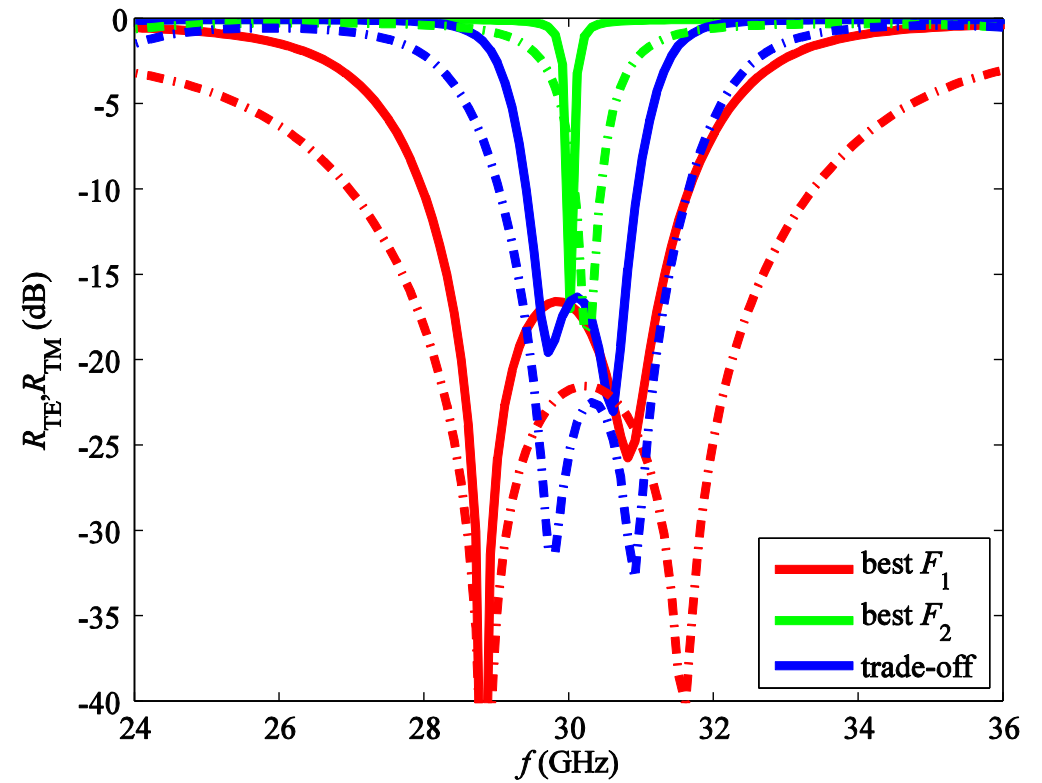
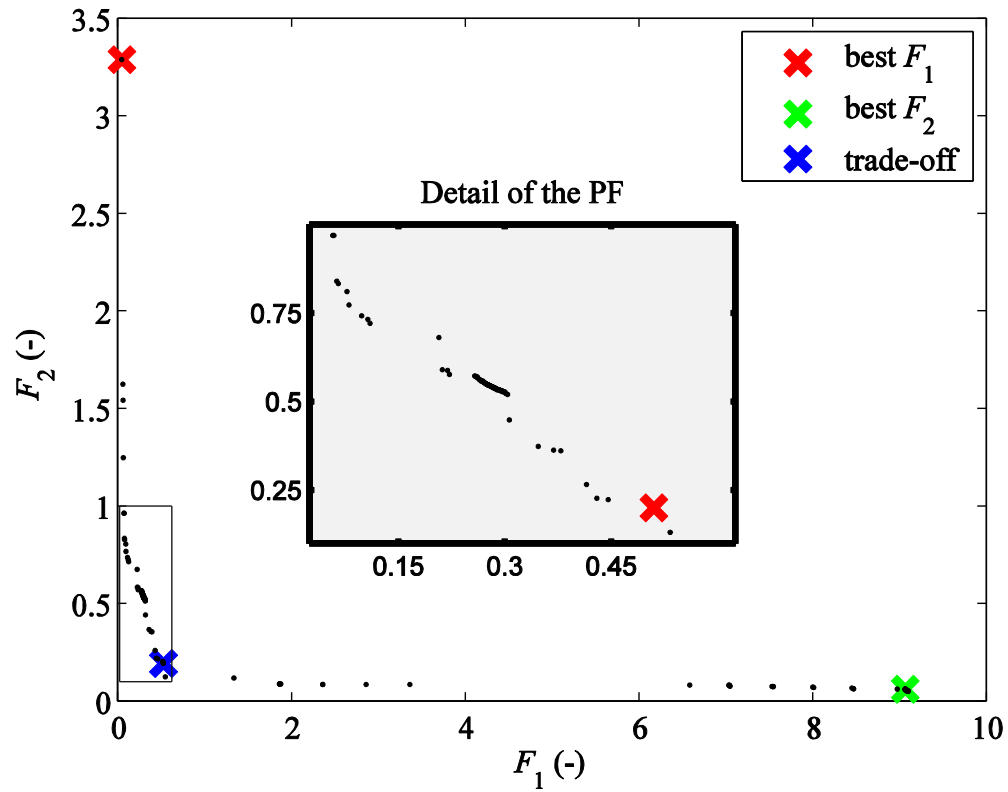
The "Add task" configuration is shown with the following fields:

- Name: task1
- Algorithm: MO\_NSGA-II
- Settings: Change algorithm settings
- Problem: user defined
- Stop conditions: 1 @ (x, y) x > niterations

A "Problem definition" dialog box is open, showing the following configuration:

- Problem name: FilterProblem
- Limits: [0.5\*ones(1,7), ones(1,7)] and [3\*ones(1,7), 10\*ones(1,7)]
- Fitness function: @ (x) {1/P\*sum(rTE(fP)^2); 1/S\*sum(1-rTE(fS)^2)}
- Constraints function (optional):
- Problem vectorized:
- Advanced problem settings
- Discrete variables (optional): {[0.568, 1.526, ...], [2.28, 3.38, ...], ...}

# Filter design



# Chain

MENU Preferences

**Tasks**

- Chains
  - myChain
    - task1
    - task2
- Comparative studies

**Add task** **Add chain** **Add comparative study**

Name: chain1

Algorithm: MO\_GDE3 >> MO\_NSGA-II  
MO\_GDE3 <<

Settings: MO\_GDE3 task will be modified.

Change algorithm settings

Problem: MODTLZ1 user defined problem

Same stop conditions for all tasks:

Stop conditions of MO\_GDE3:

1 @(x, y)x > niterations

Add another stop condition

# Comparative Study

Name:

Algorithm:

	Task 1	Task 2	
Chain 1	MO_GDE3	MO_NSGA-II	>>
Chain 2	MO_GDE3		

\

Settings: MO\_GDE3 task will be modified.

Problem:

>>

<<

Number of runs:

Request:

>>

<<

# Comparative Study

The interface shows a task tree on the left with the following structure:

- Tasks
- Chains
  - myChain
    - task1
    - task2
- Comparative studies
  - comparativeStudy1
    - chain1
      - task1
      - task2
    - chain2
      - task1
    - chain3
      - task1
      - task2
    - chain4
      - task1
    - chain5
      - task1
      - task2
    - chain6
      - task1

The configuration window for a new comparative study is titled "Name:" and contains the text "comparativeStudy2".

The Results window displays a table with the following data:

	chain1, MOFON	chain2, MOFON	chain3, MOKUR	chain4, MOKUR	chain5, MOPOL	chain6, MOPOL
Run1	0.0999	0.1091	0.1749	0.2018	0.1585	0.2310
Run2	0.0844	0.1074	0.1764	0.2193	0.3772	0.2155
Run3	0.1024	0.1157	0.1822	0.2418	0.1873	0.2259
Run4	0.1051	0.1077	0.1910	0.1816	0.1849	0.4289
Run5	0.0958	0.1228	0.2030	0.1870	0.1712	0.2003
Run6	0.1250	0.1098	0.1720	0.2179	0.1812	0.2469
Run7	0.1069	0.1163	0.1901	0.1958	0.1901	0.2525
Run8	0.1026	0.1253	0.2117	0.2099	0.2038	0.4875
Run9	0.0907	0.1199	0.1930	0.1886	0.1879	0.2247
Run10	0.0978	0.1157	0.1948	0.1949	0.1948	0.2197
Mean	0.1010	0.1150	0.1889	0.2039	0.2037	0.2733
Standard deviation	0.0108	0.0064	0.0127	0.0185	0.0622	0.0995
Minimum	0.0844	0.1074	0.1720	0.1816	0.1585	0.2003
Maximum	0.1250	0.1253	0.2117	0.2418	0.3772	0.4875

# Comparative Study

The screenshot shows a software interface for managing comparative studies. On the left, a tree view lists the following structure:

- Tasks
- Chains
  - myChain
    - task1
    - task2
- Comparative studies
  - comparativeStudy1
    - chain1
      - task1
      - task2
    - chain2
      - task1
    - chain3
      - task1
      - task2
    - chain4
      - task1
    - chain5
      - task1
      - task2
    - chain6
      - task1

At the top right, there are three buttons: "Add task", "Add chain", and "Add comparative study". Below them is a "Name:" label and a text input field containing "comparativeStudy2".

The "Results" window is open, showing a table of metrics. The "Choose metric to show:" dropdown is set to "Spread". The table has the following data:

	chain1, MOFON	chain2, MOFON	chain3, MOKUR	chain4, MOKUR	chain5, MOPOL	chain6, MOPOL
Run1	0.0999	0.1091	0.1749	0.2018	0.1585	0.2310
Run2	0.0844	0.1074	0.1764	0.2193	0.3772	0.2155
Run3	0.1024	0.1157	0.1822	0.2418	0.1873	0.2259
Run4	0.1051	0.1077	0.1910	0.1816	0.1849	0.4289
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Run6	0.1250	0.1098	0.1720	0.2179	0.1812	0.2469
Run7	0.1069	0.1163	0.1901	0.1958	0.1901	0.2525
Run8	0.1026	0.1253	0.2117	0.2099	0.2038	0.4875
Run9	0.0907	0.1199	0.1930	0.1886	0.1879	0.2247
Run10	0.0978	0.1157	0.1948	0.1949	0.1948	0.2197
Mean	0.1010	0.1150	0.1889	0.2039	0.2037	0.2733
Standard deviation	0.0108	0.0064	0.0127	0.0185	0.0622	0.0995
Minimum	0.0844	0.1074	0.1720	0.1816	0.1585	0.2003
Maximum	0.1250	0.1253	0.2117	0.2418	0.3772	0.4875

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Thank you for your attention!



<http://antennatoolbox.com/>

[kadlecp@feec.vutbr.cz](mailto:kadlecp@feec.vutbr.cz)

+420 541146552