

Evolutionary Optimization in Electromagnetics

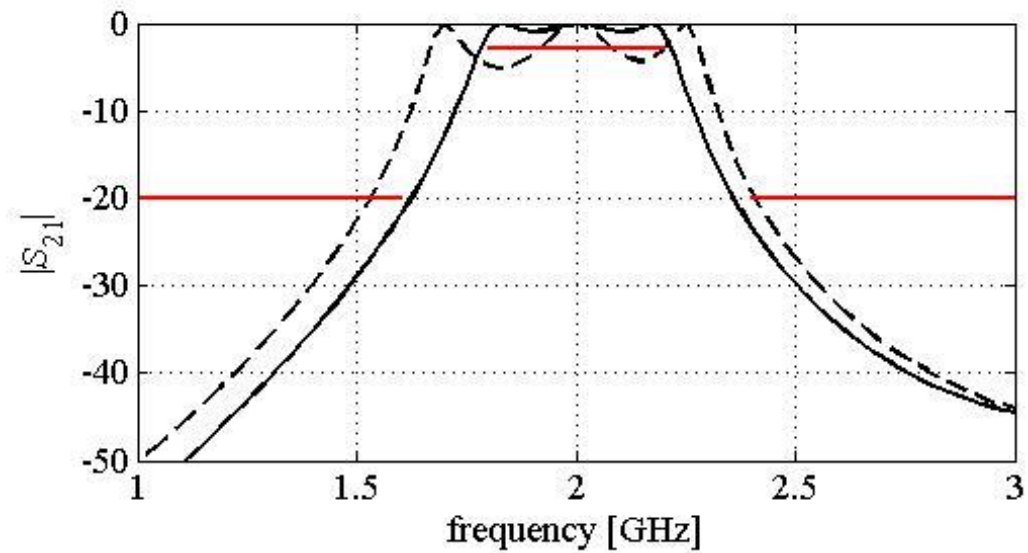
COST IC 1407 Workshop,
Bratislava, 5.4.2017

Outline

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

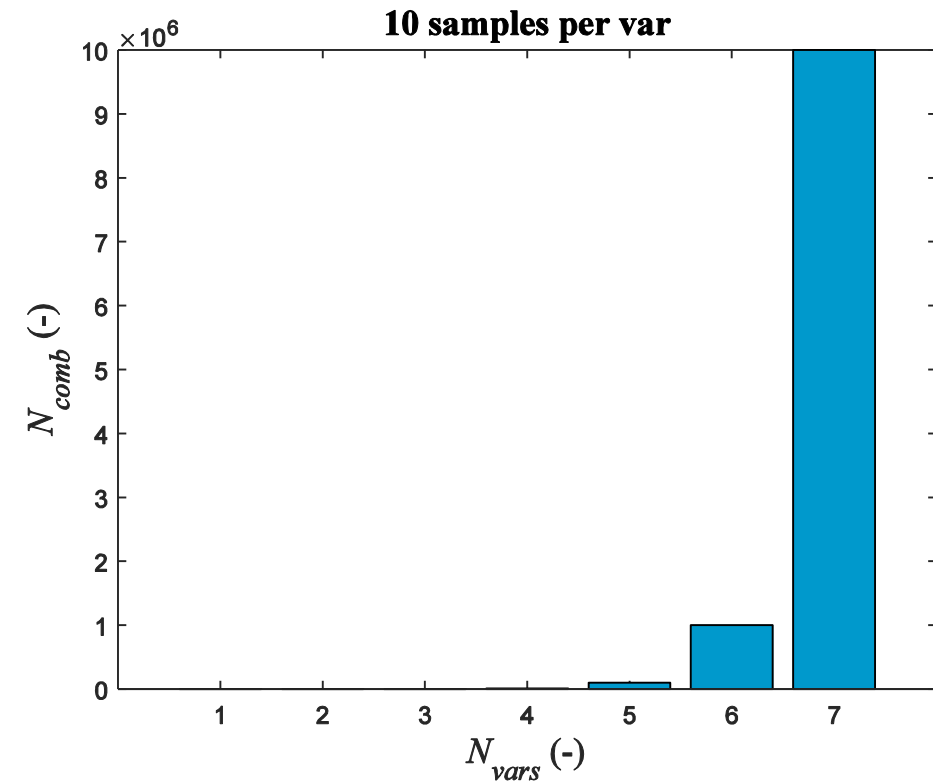
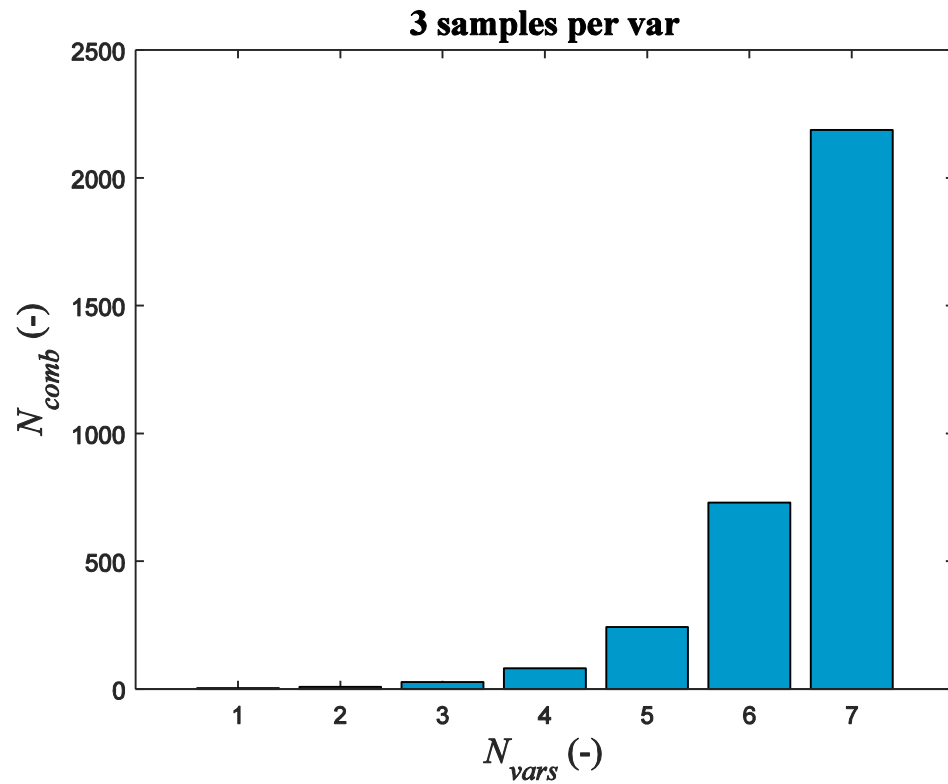
Motivation

Always try to do the best...



Motivation

Is parametric analysis the way?



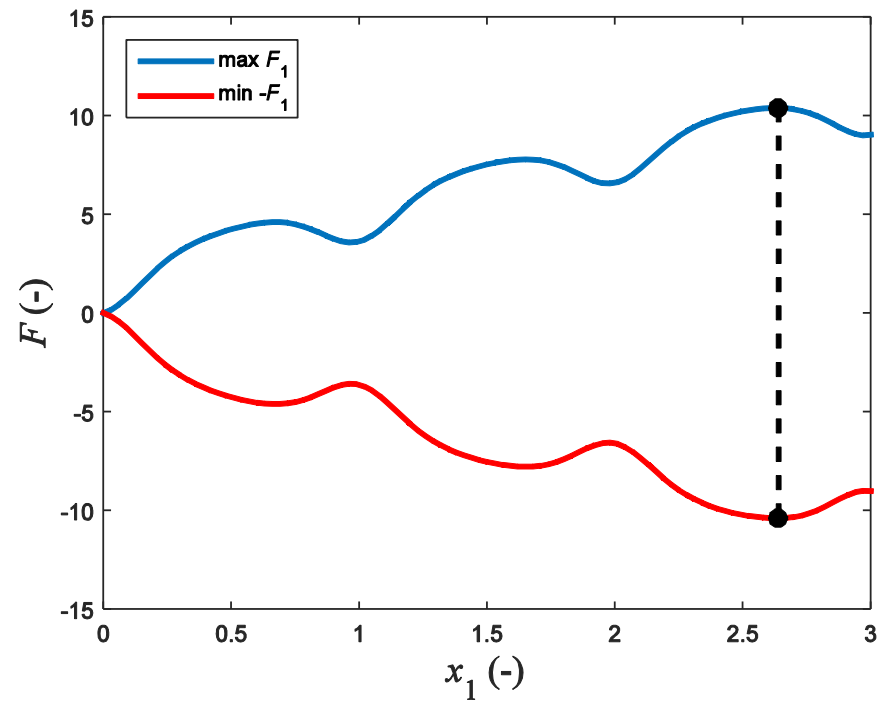
Optimization

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

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

Min vs. Max

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



Objective formulation

Problem:

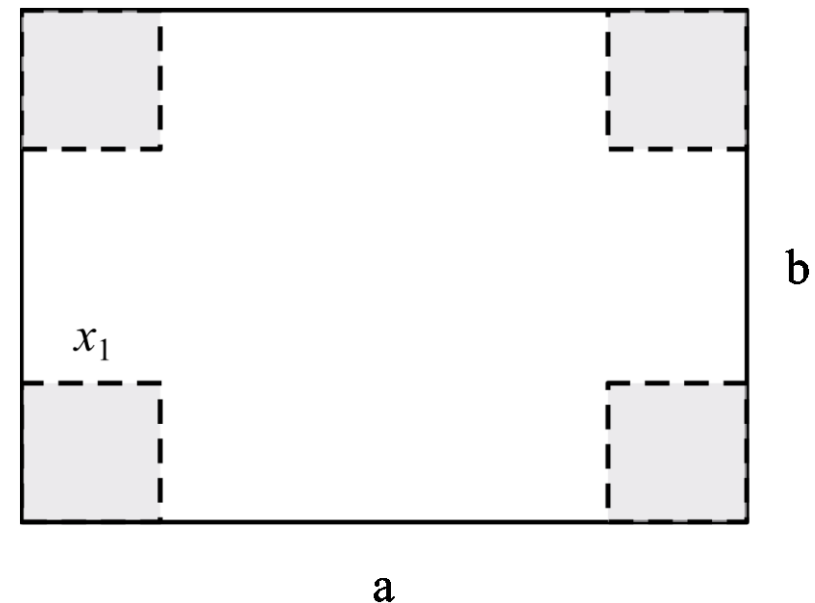
Having sheet of paper with dimensions a and b , what is the size of squares to be cut from the corners of the sheet to build a box with the highest volume?

Decision space:

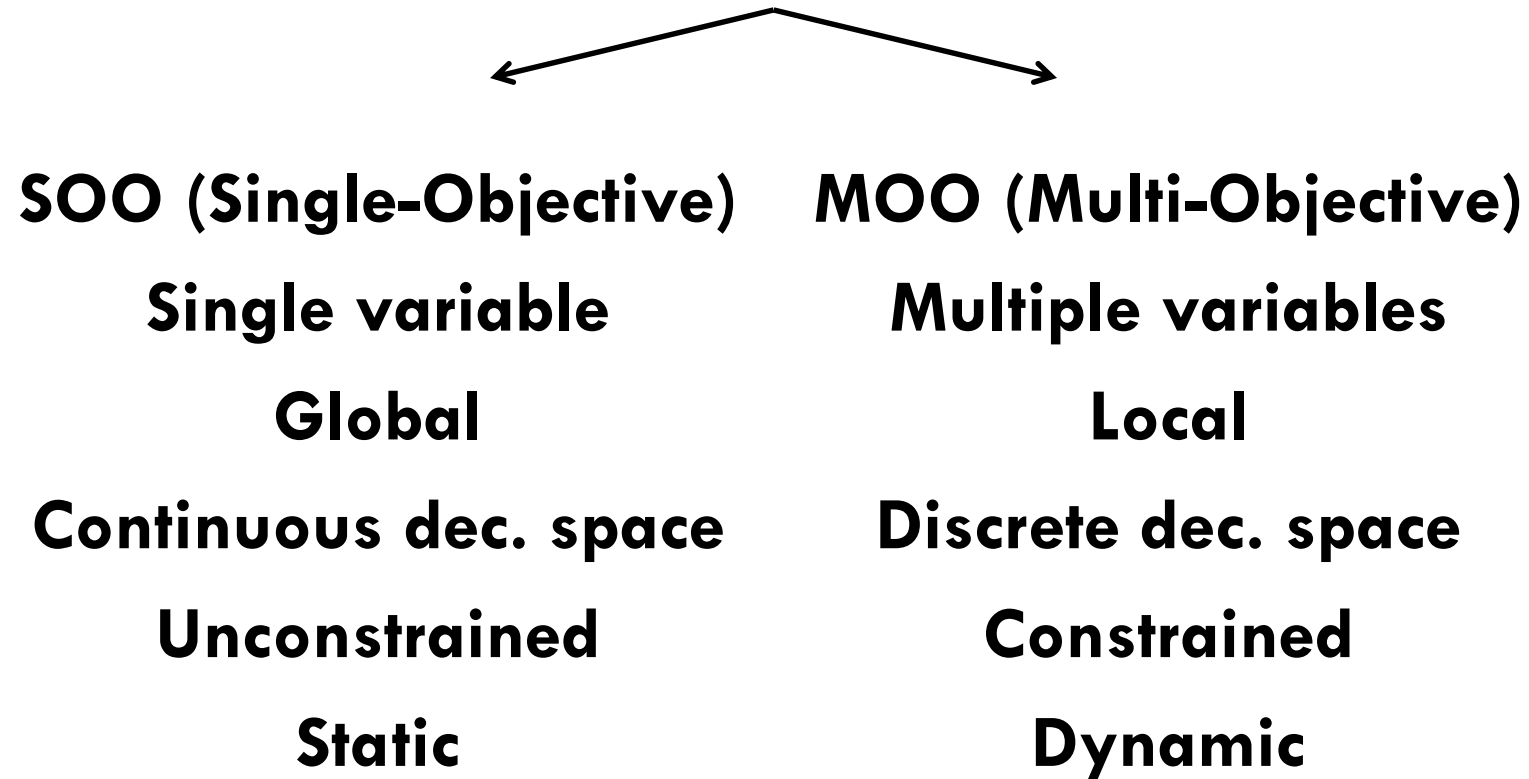
$$\mathbf{x} = [x_1]$$

Objective space

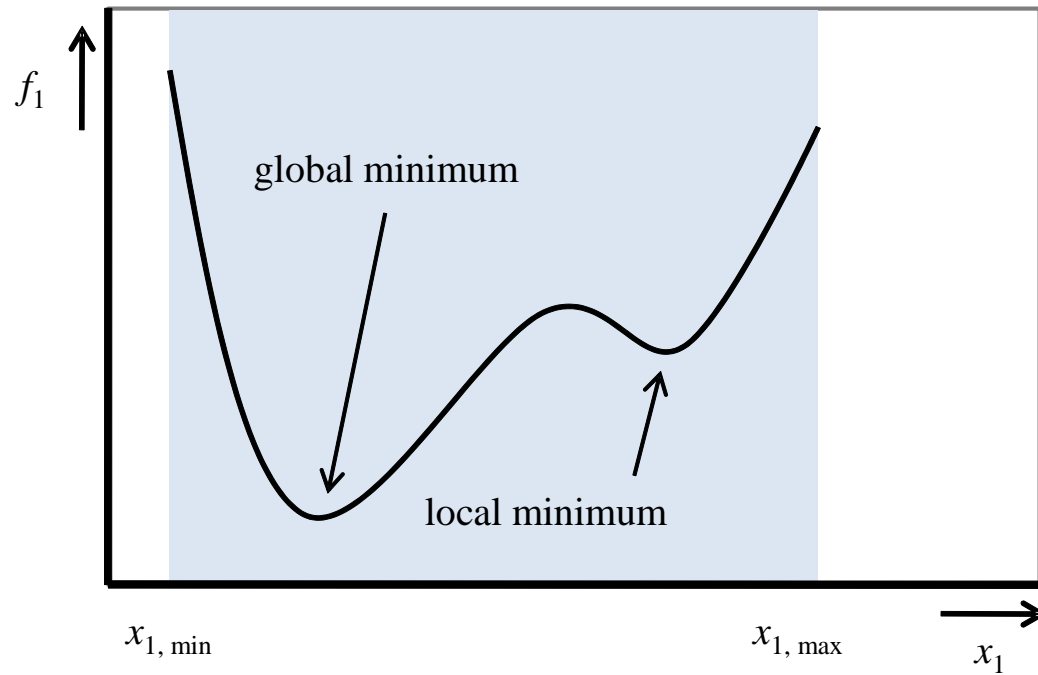
$$f_1 = 4x_1^3 - (2a + 2b)x_1^2 + abx_1$$



Optimization taxonomy



Global vs. local minimum



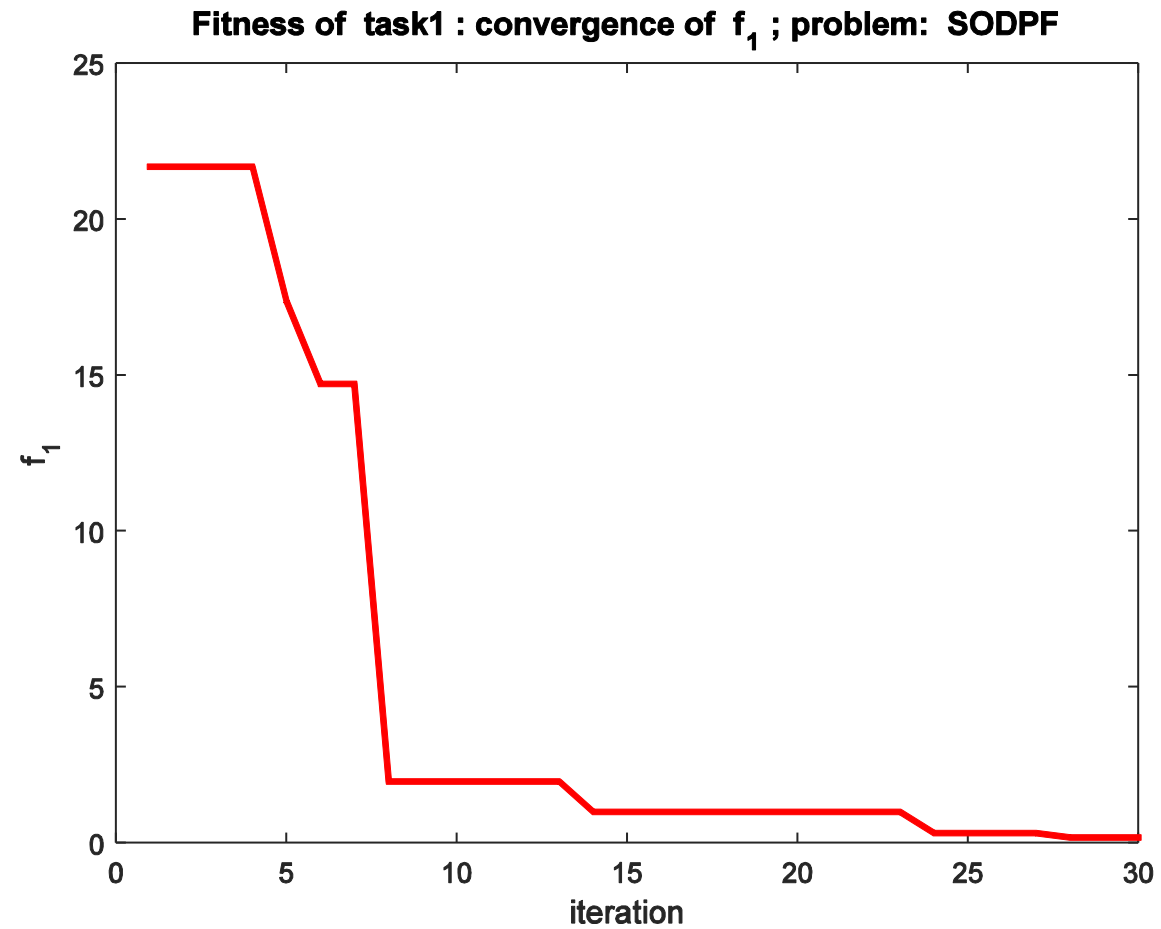
Global vs. Local

- 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

- 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

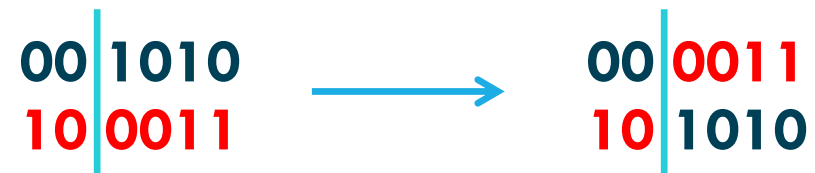
- Holland 1962
- Inspired by Darwin – only good properties of genome are maintained for next generations
- **GAs work with discrete decision space!!!**

GA - taxonomy

- 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

- **Crossover:**



- **Mutation:**



GA - reproduction

- **Decimation:**

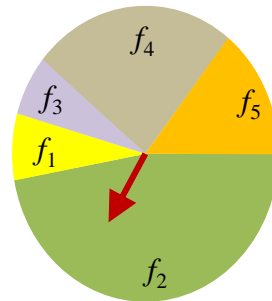
g_1	g_2	g_3	f
110	0101	00010	0 , 15
101	1101	00101	0 , 36
011	1011	00110	1 , 53
100	1010	10001	6 , 27
011	0011	11010	11 , 83
101	1011	10010	13 , 21
100	0101	00101	20 , 89
011	0101	10101	56 , 12

- **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

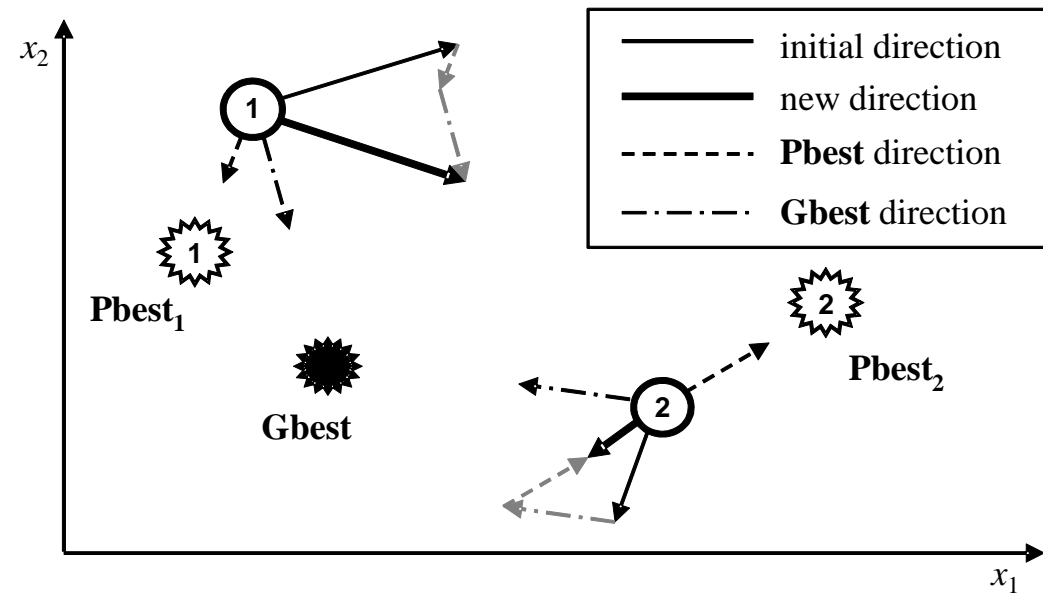
- 1995 - Eberhart and Kennedy
- cooperation of swarm over the meadow
- cognitive learning: personal experience of agent
- social learning: experience of the whole swarm

Swarm update

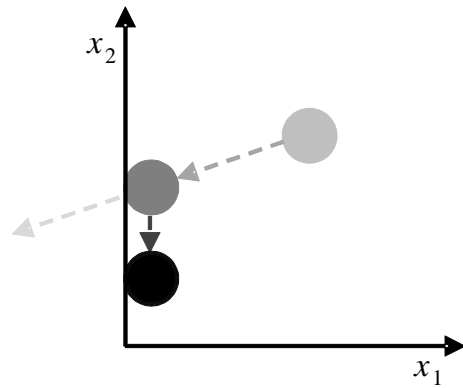
$$\mathbf{x}_q(i) = \mathbf{x}_q(i-1) + \mathbf{v}_q(i-1)$$

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

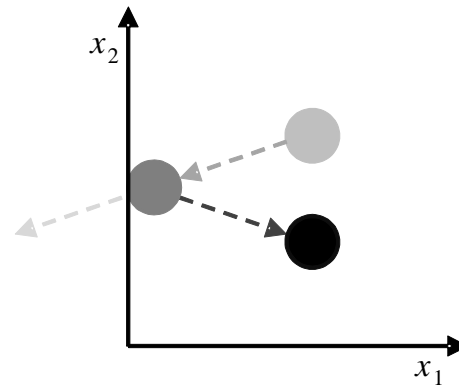
Swarm update



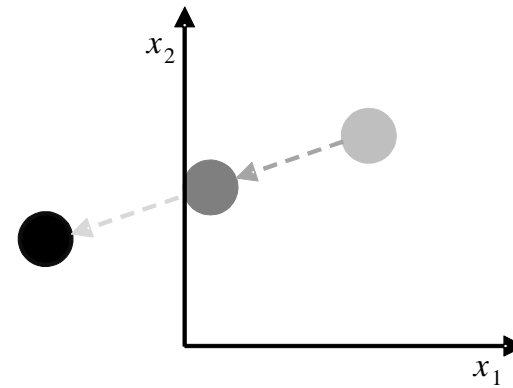
Wall types



Absorbing

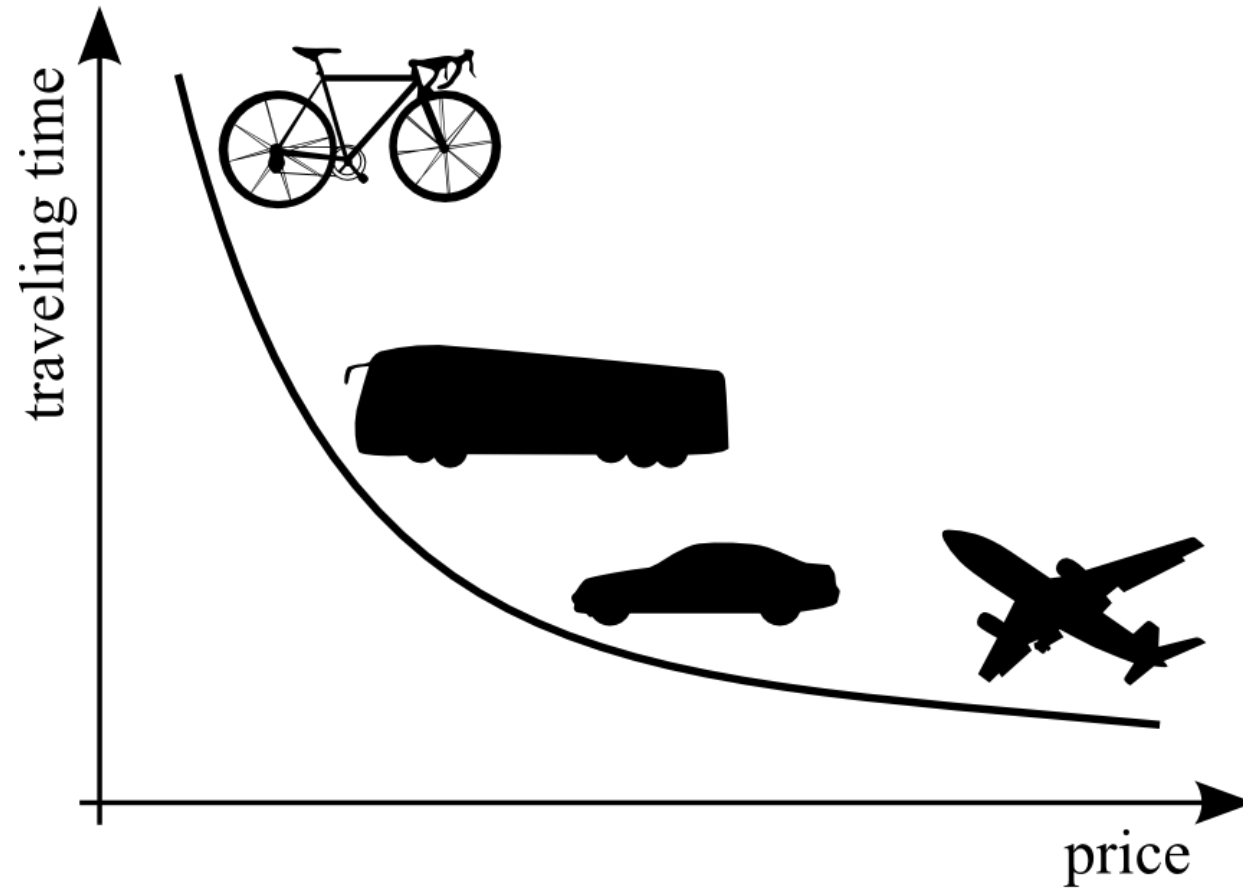


Reflecting



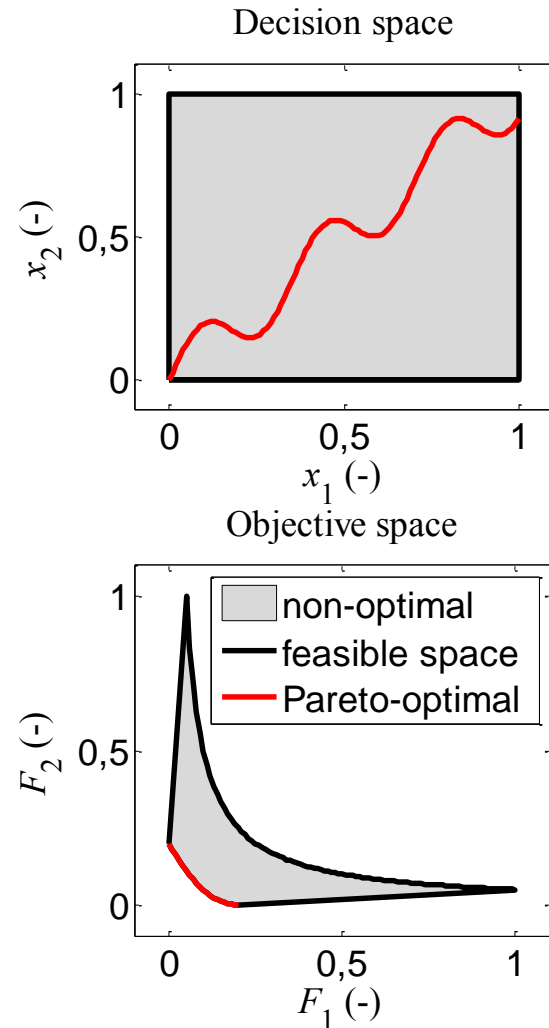
Invisible

Multi-objective optimization

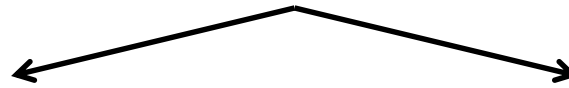


Pareto front

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



Multi-objective strategies



Aggregating methods

Transform to SOOP

Single solution

How to choose w ?

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

MO Optimization

More complex routine

Set of Pareto optimal

Extra information

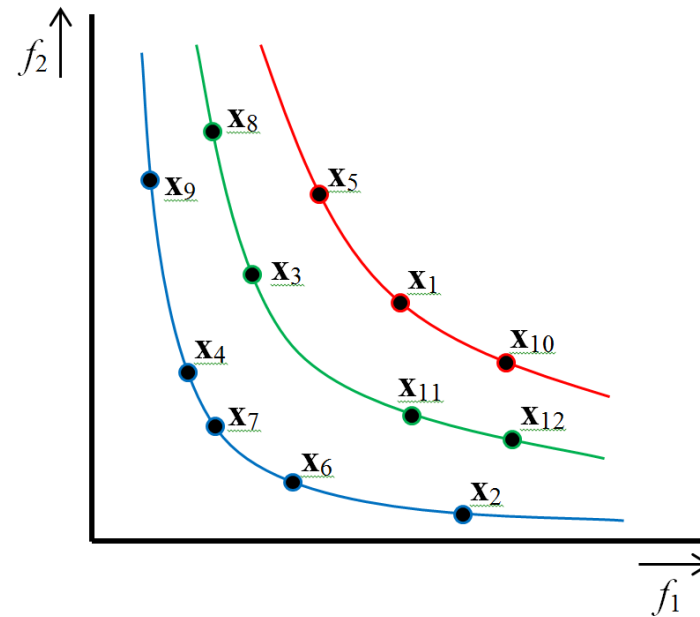
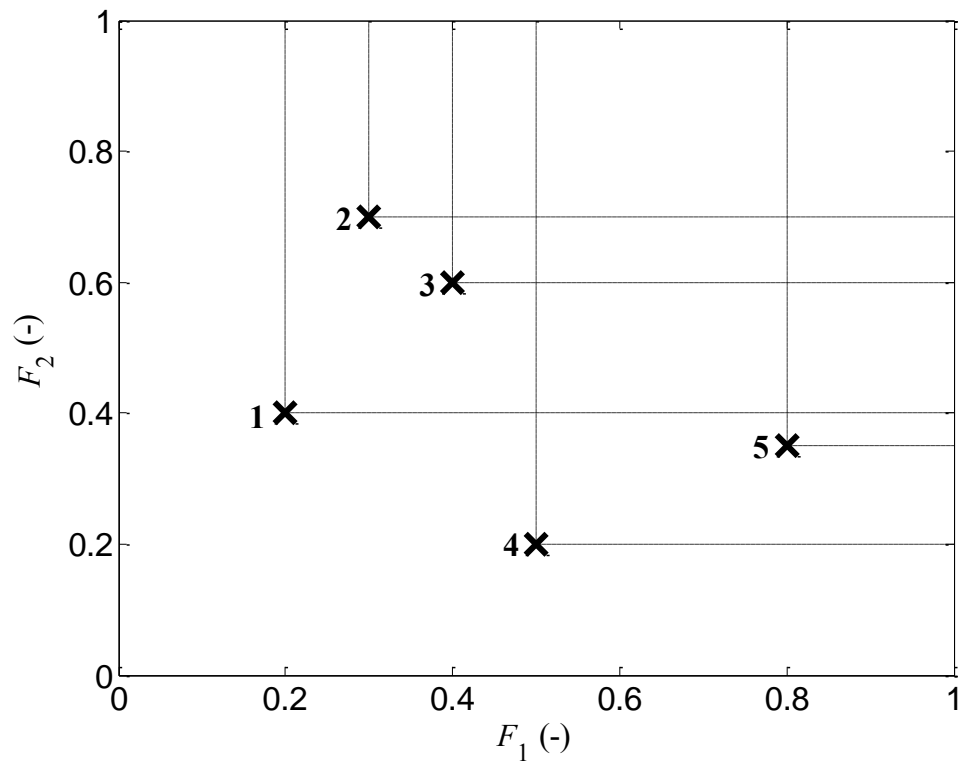
Decision making

Dominance concept

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}

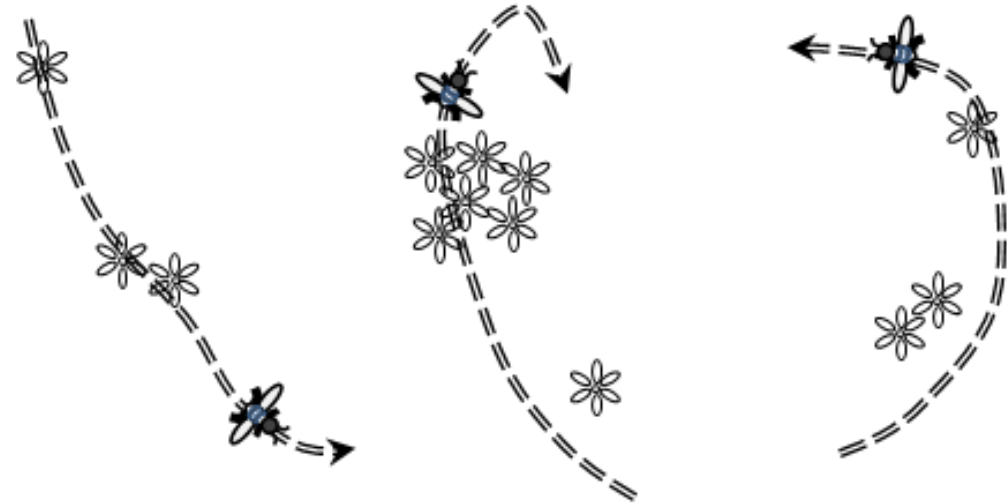
best

worst

MOPSO

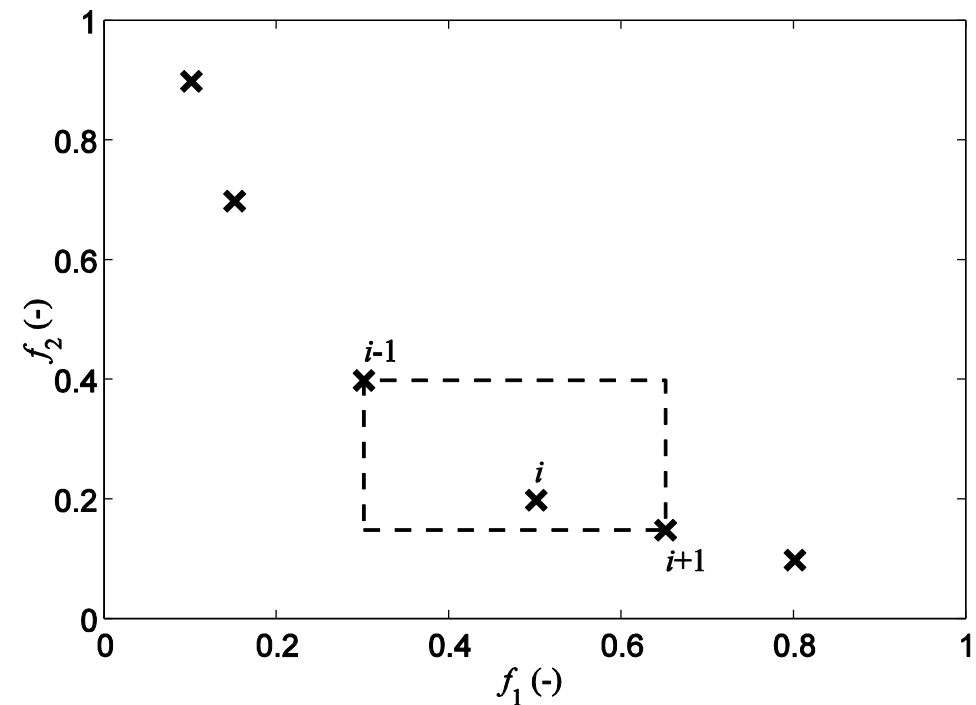
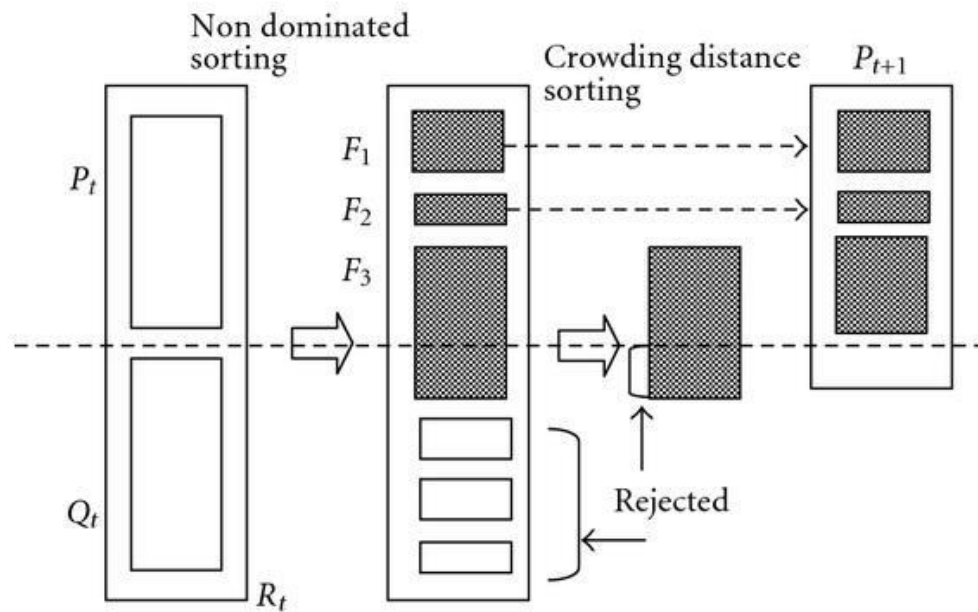
Nanbo, J., Rahmat-Samii, Y. "Advances in Particle Swarm Optimization for Antenna Designs: Real-Number, Binary, Single-Objective and Multiobjective Implementations," IEEE Transactions on Ant. Propag., vol. 55, no. 3, pp. 556-567, 2007.

- **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

- 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

Problem:

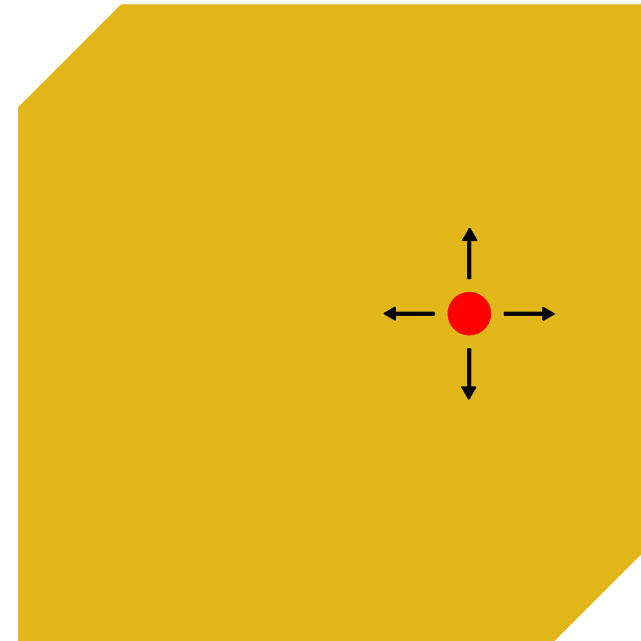
Find proper point on antenna for coaxial port.

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

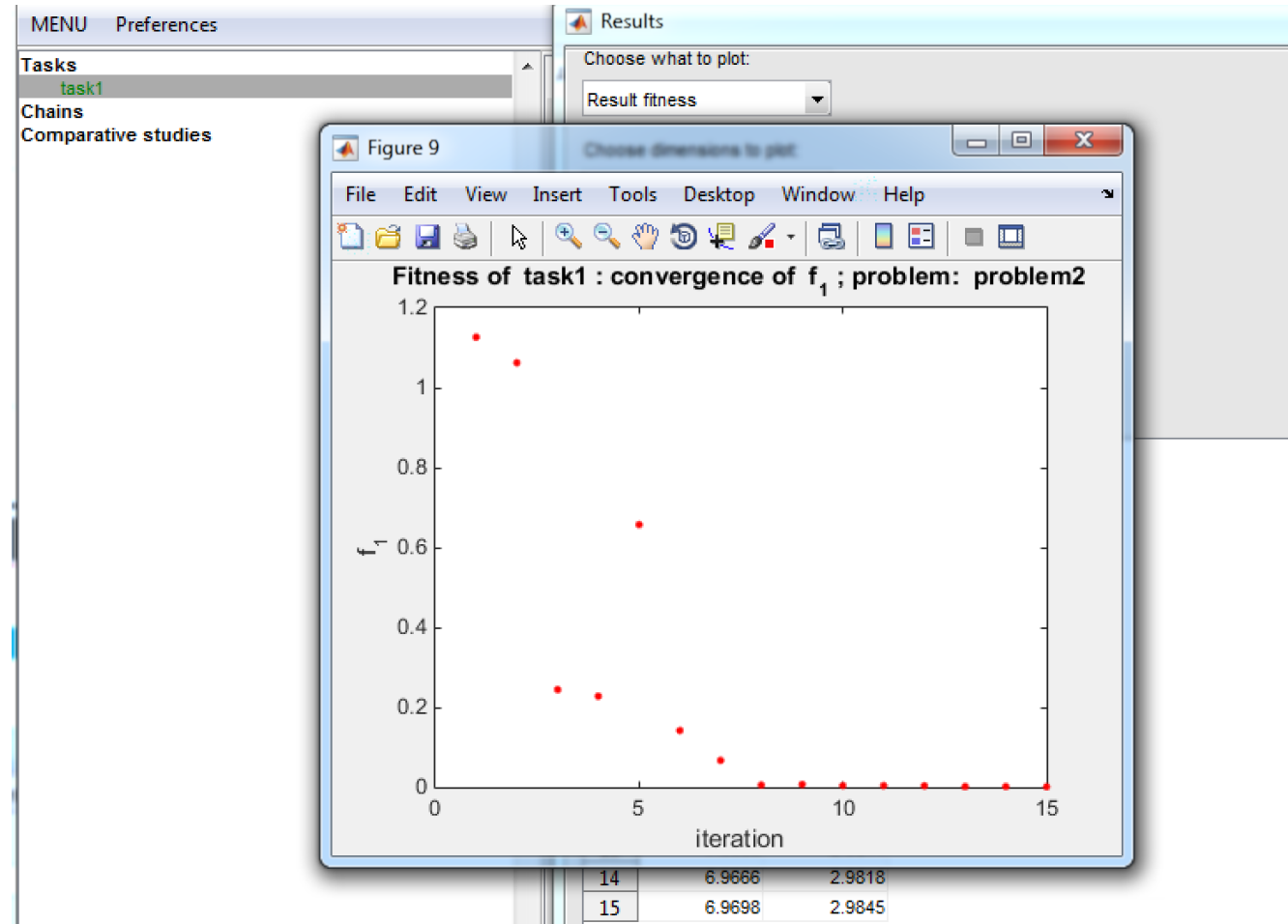
Inertia weight (W):
0.6 0.4

Cognitive learning factor (C1):
1.5

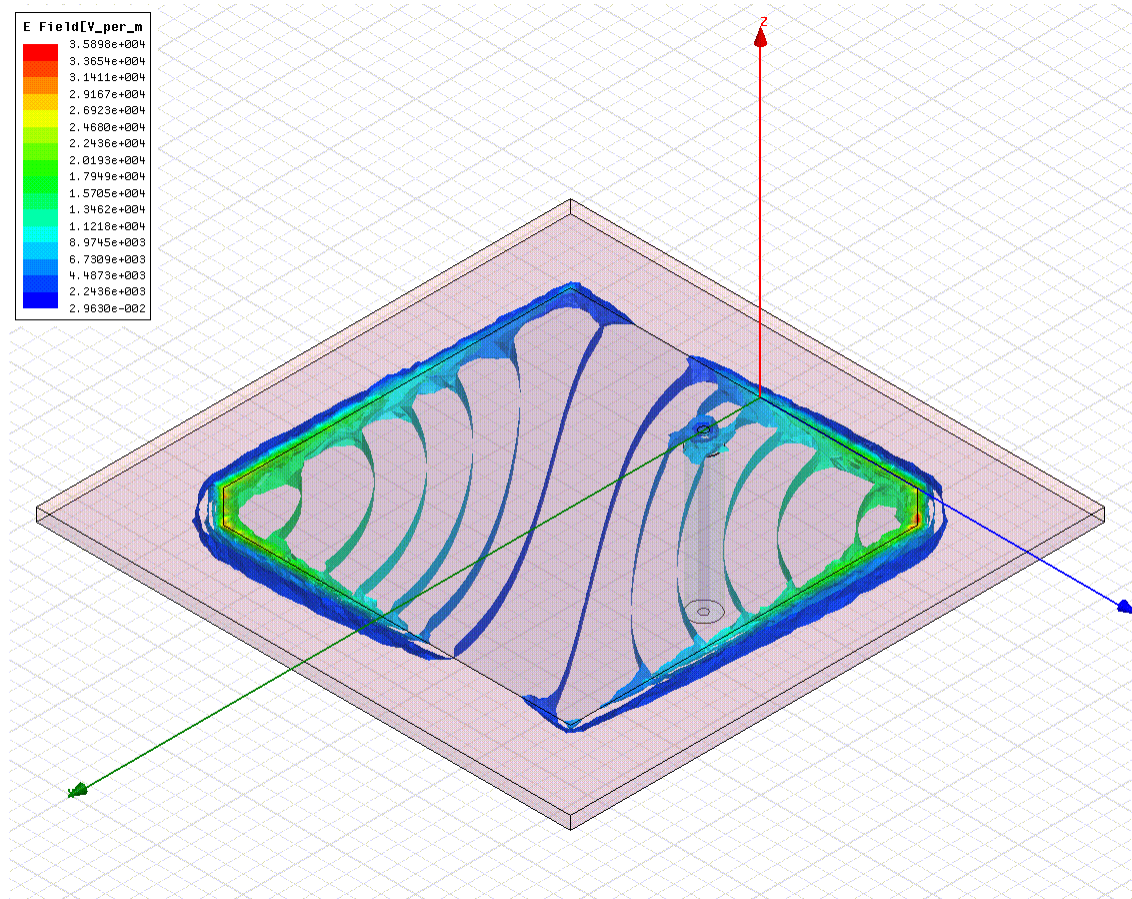
Social learning factor (C2):
1.5

Boundary type:
reflecting

Confirm settings



Feeding Point



Filter design

Problem:

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

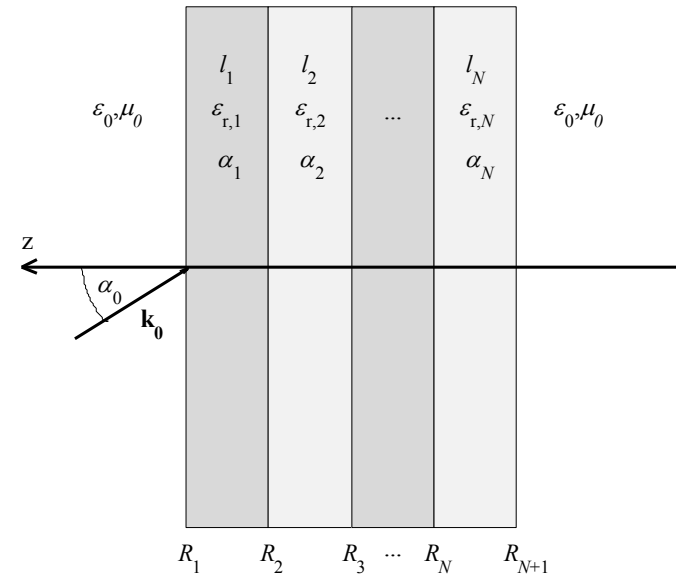
Decision space:

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

Objective space

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

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



Filter design

MENU Preferences

Tasks
Chains
Comparative studies

Add task Add chain Add comparative study

Name:
task1

Algorithm:
MO_NSGA-II

Settings:
Change algorithm settings

Problem:
user defined

Stop conditions:
1 @ (x, y) x > niterations
Add another condition

Problem definition

Problem name:
FilterProblem

Limits:
[0.5*ones(1,7), ones(1,7)]
[3*ones(1,7), 10*ones(1,7)]

Fitness function:
@(x) {1/P*sum(rTE(fP)^2); 1/S*sum(1-rTE(fS)^2)}

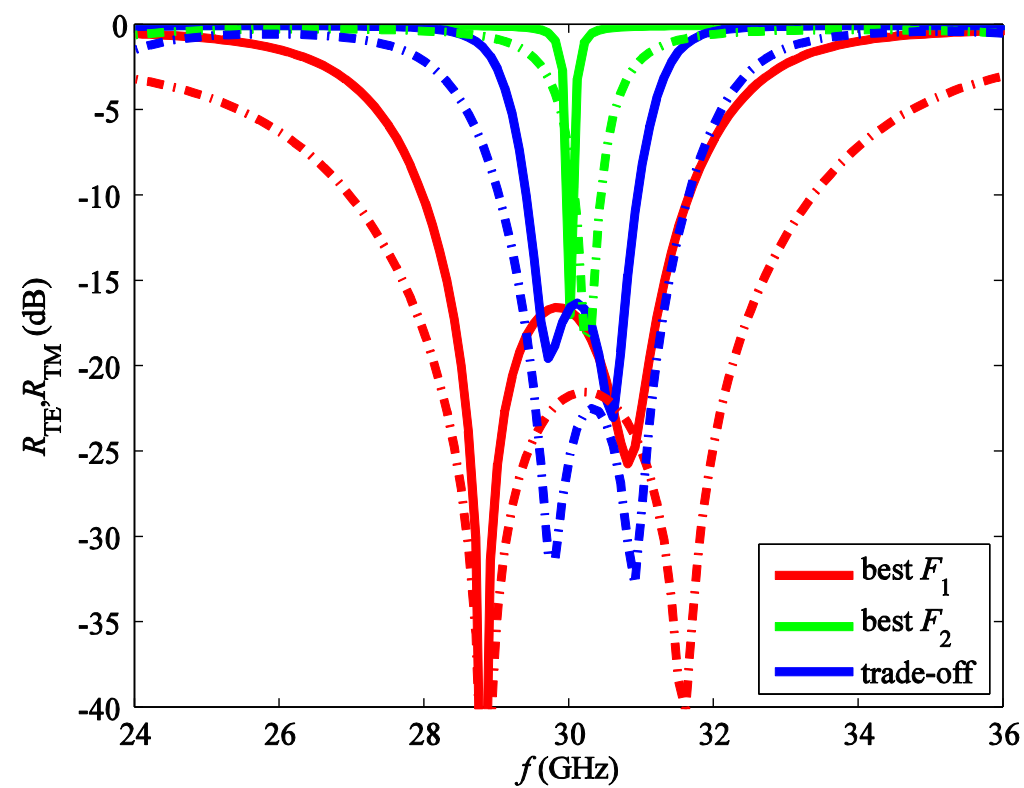
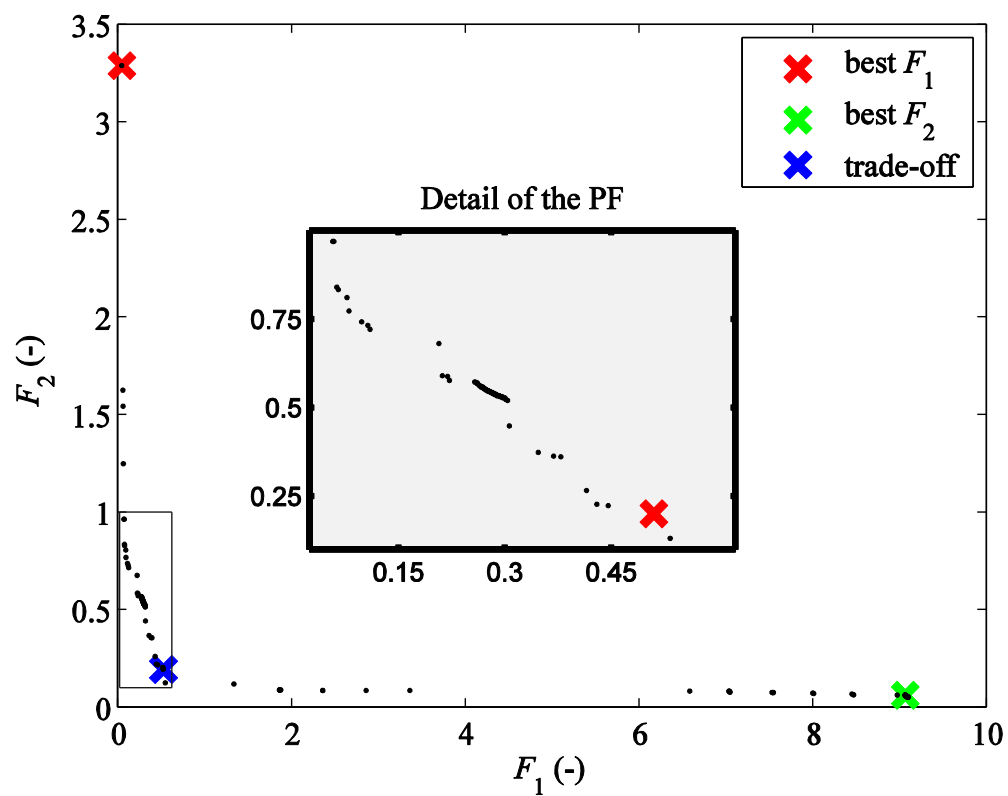
Constraints function (optional):

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:

>>

<<

MOFON

MOKUR

MOPOL

Number of runs:

Request:

>>

<<

GenerationalDistance

Spread

Comparative Study

The interface shows a tree view 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 top panel contains buttons: Add task, Add chain, Add comparative study. Below these is a 'Name:' field with the value 'comparativeStudy2'.

The 'Results' window is open, showing a table of metrics. The 'Choose metric to show:' dropdown is set to 'Spread'.

	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

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 - chain6
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Thank you for your attention!



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