

Multi-Objective Evolutionary Algorithms

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Classical Approaches: ε -Constraint Method

Method: Minimize a primary objective while expressing all the other objectives in the form of inequality constraints

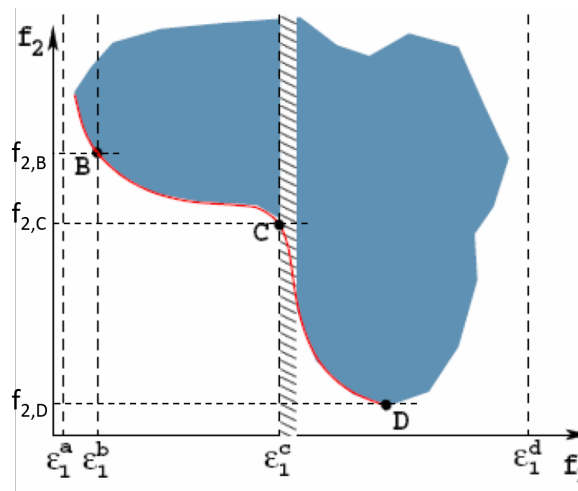
$$\text{Minimize } f_p(x)$$

$$\text{subject to } f_i(x) \leq \varepsilon_i; \quad i = 1, \dots, m; \quad i \neq p$$

Remarks:

- Need to know relevant ε vectors to ensure a feasible solution
- Non-uniformity in Pareto-optimal solutions
- However, any Pareto-optimal solution can be found with this method

Ex.: Minimize $f_2(x)$
subject to $f_1(x) \leq \varepsilon_i$



Difficulties with Most Classical Approaches

- Need to run a single-objective optimizer many times
- Expect a lot of problem knowledge
- Even then, good distribution of solutions is not guaranteed
- Multi-objective optimization as an application of single-objective optimization

- **NSGA**

Srinivas, N., and Deb, K.: Multi-objective function optimization using non-dominated sorting genetic algorithms, Evolutionary Computation Journal 2(3), pp. 221-248, 1994

- **NSGA-II**

Kalyanmoy Deb, Samir Agrawal, Amrit Pratap, and T Meyarivan: A Fast Elitist Non-Dominated Sorting Genetic Algorithm for Multi-Objective Optimization: NSGA-II, In Proceedings of the Parallel Problem Solving from Nature VI Conference, 2000

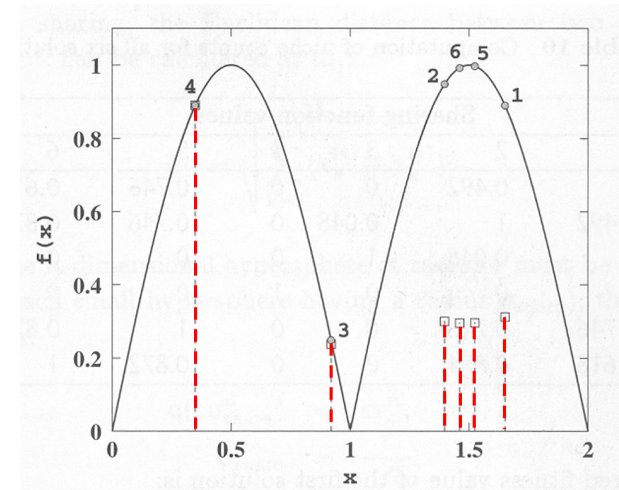
- ...

Fitness Sharing: Example

Bimodal function - six solutions and corresponding shared fitness functions

- $\sigma_{share} = 0.5, \alpha = 1.$

Sol. i	String	Decoded value	$x^{(i)}$	f_i	nc_i	f'_i
1	110100	52	1.651	0.890	2.856	0.312
2	101100	44	1.397	0.948	3.160	0.300
3	011101	29	0.921	0.246	1.048	0.235
4	001011	11	0.349	0.890	1.000	0.890
5	110000	48	1.524	0.997	3.364	0.296
6	101110	46	1.460	0.992	3.364	0.295



©Kalyanmoy Deb: Multi-Objective Optimization using Evolutionary Algorithms.

Let's take the first solution

- $d_{11} = 0.0, d_{12} = 0.254, d_{13} = 0.731, d_{14} = 1.302, d_{15} = 0.127, d_{16} = 0.191$
- $Sh(d_{11}) = 1, Sh(d_{12}) = 0.492, Sh(d_{13}) = 0, Sh(d_{14}) = 0, Sh(d_{15}) = 0.746, Sh(d_{16}) = 0.618.$
- $nc_1 = 1 + 0.492 + 0 + 0 + 0.746 + 0.618 = 2.856$
- $f'(1) = f(1)/nc_1 = 0.890/2.856 = 0.312$

NSGA-II

Fast non-dominated sorting approach

- Computational complexity of $O(MN^2)$.

Diversity preservation

- the sharing function method is replaced with a **crowded comparison approach**,
- parameterless approach.

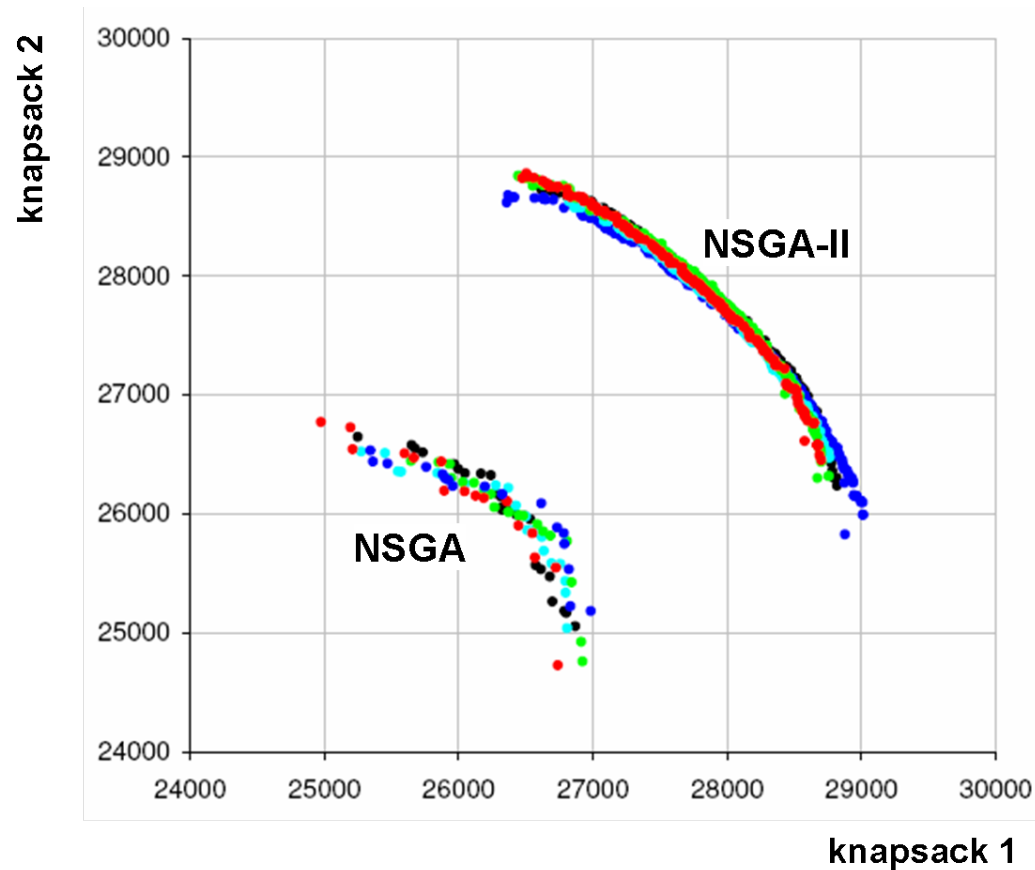
Elitist evolutionary model

- Only the best solutions survive to subsequent generations.

Simulation Results: NSGA vs. NSGA-II

Comparison of NSGA and NSGA-II on bi-objective 0/1 Knapsack Problem with 750 items.

NSGA-II outperforms NSGA with respect to both performance measures.



NSGA-II: Constraint Handling Approach

Binary tournament selection with modified domination concept is used to choose the better solution out of the two solutions i and j , randomly picked up from the population.

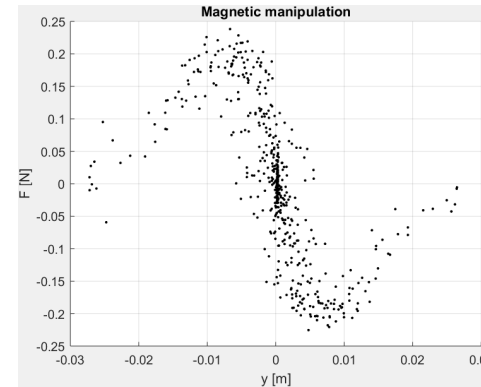
In the presence of constraints each solution in the population can be either **feasible** or **infeasible**, so that there are the following three possible situations:

1. both solutions are feasible,
2. one is feasible and other is not,
3. both are infeasible.

NSGA-II: Bi-objective Symbolic Regression

Optimization objectives:

- Minimize MSE on the training data set
- Minimize deviation of the symbolic models from the desired properties

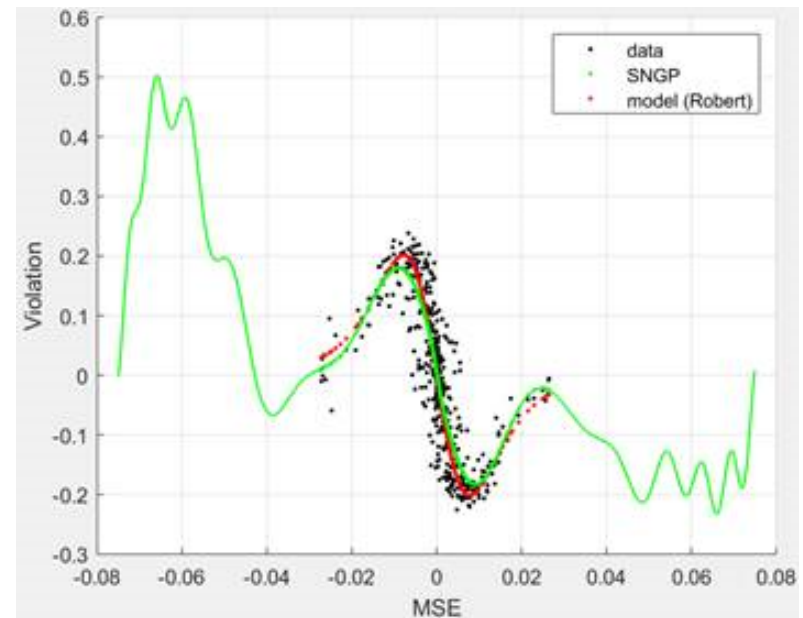
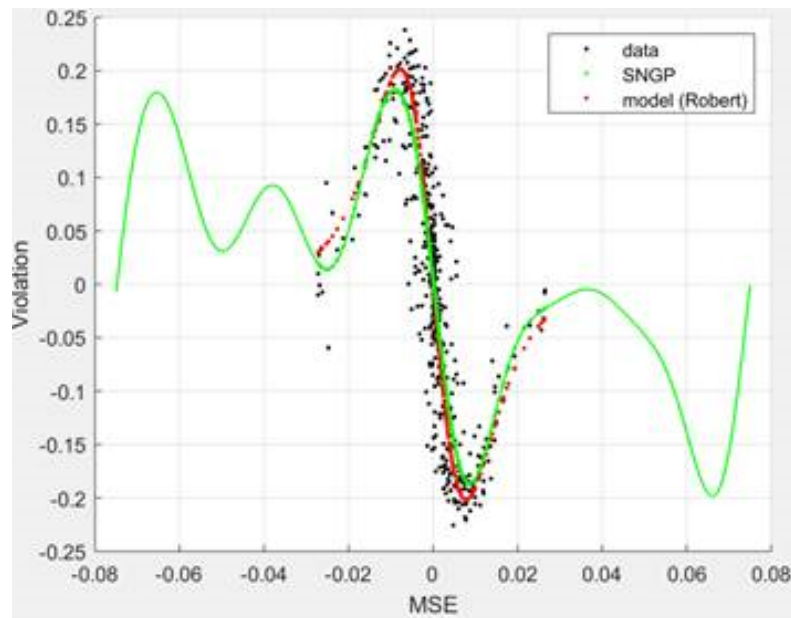


Desired properties:

- Monotonically increasing in the intervals $y = \langle -0.075, -0.01 \rangle$ and $y = \langle 0.01, 0.075 \rangle$
- Monotonically decreasing in the interval $y = \langle -0.07, 0.07 \rangle$
- $F(y) \geq$, for $y \in \langle -0.075, 0.0 \rangle$
- $F(y) \leq$, for $y \in \langle 0.0, 0.075 \rangle$
- $|F(0.0)| < 0.005$
- $|F(-0.075) - 0.001| < 0.0005$
- $|F(0.075) + 0.001| < 0.0005$

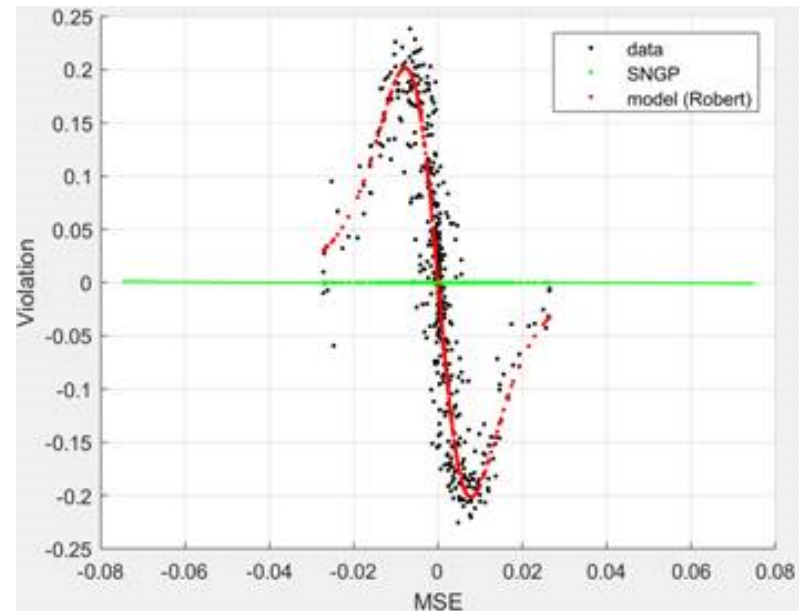
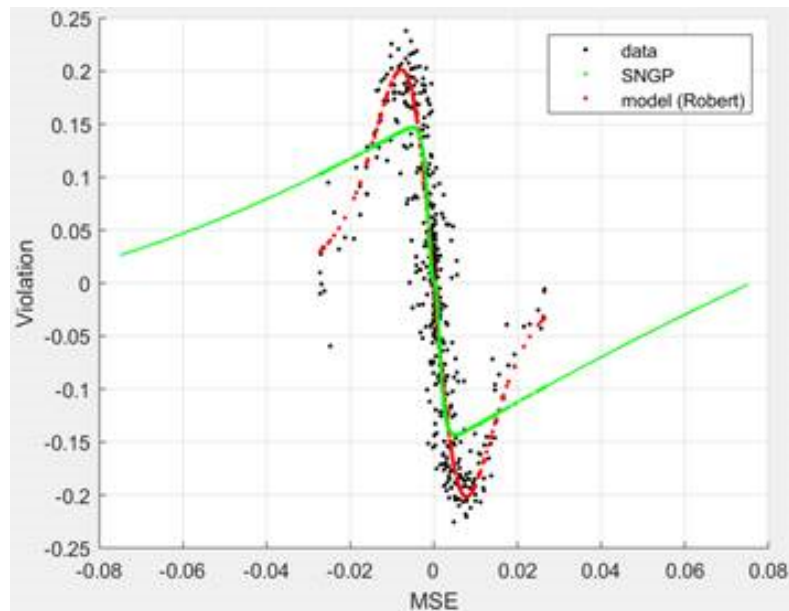
NSGA-II: Bi-objective Symbolic Regression

Well-fit models w.r.t. the MSE on training data only



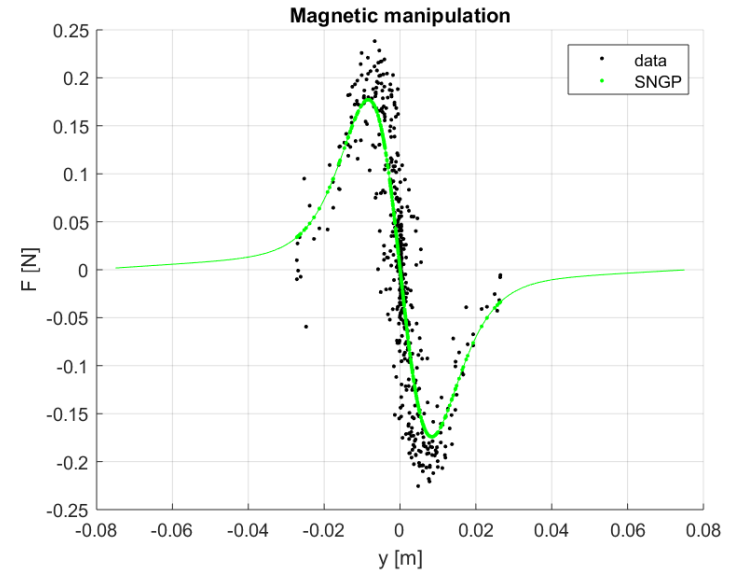
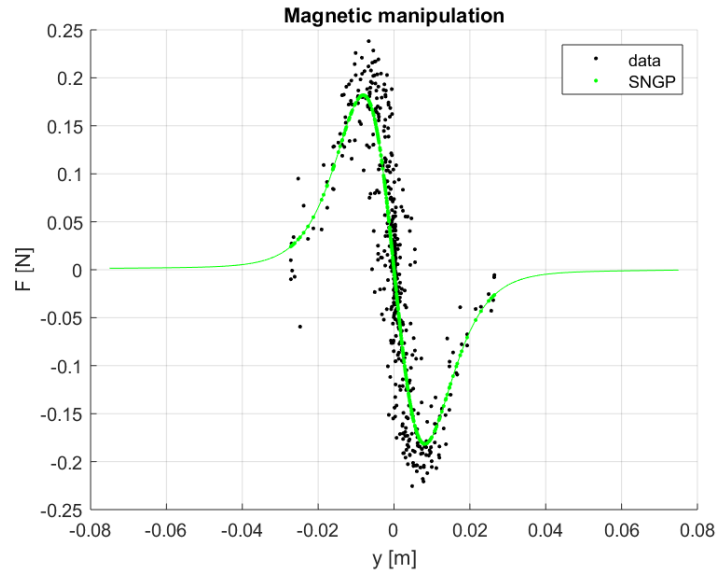
NSGA-II: Bi-objective Symbolic Regression

Well-fit models w.r.t. the constraint violations



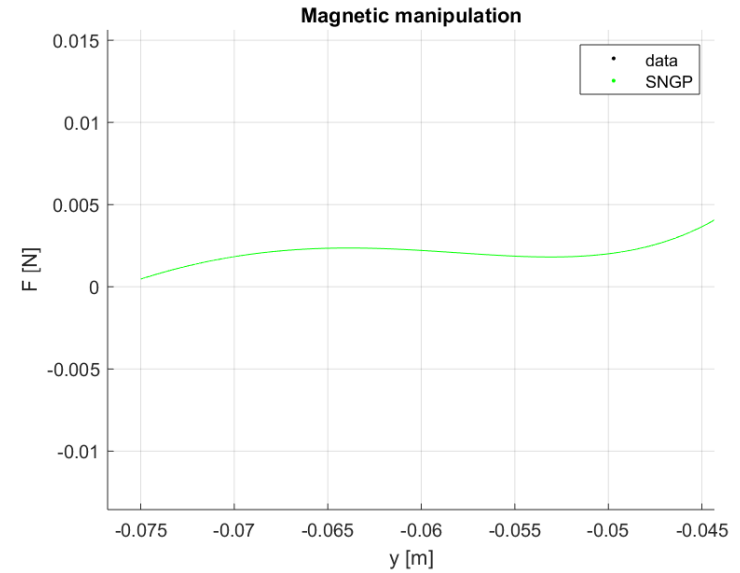
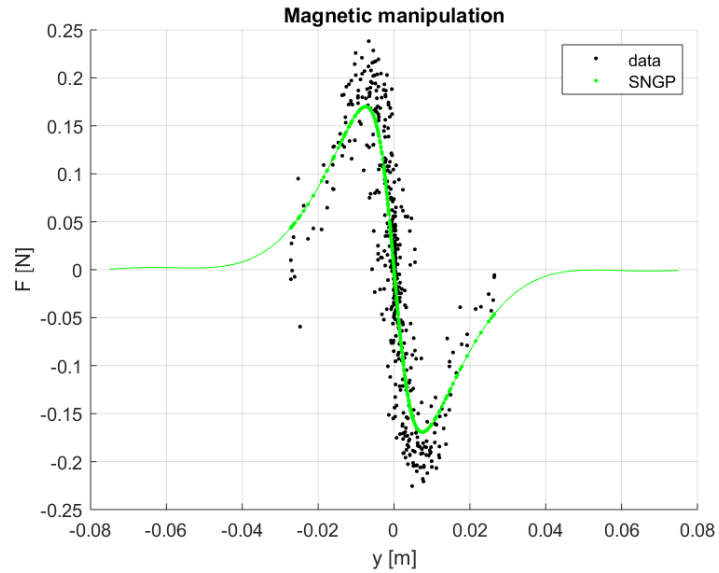
NSGA-II: Bi-objective Symbolic Regression

Models with small MSE on training data that fully comply with the constraints



NSGA-II: Bi-objective Symbolic Regression

Models with small MSE on training data that almost fully comply with the constraints



Strength Pareto Evolutionary Algorithm 2 (SPEA2)

SPEA2 maintains two sets of solutions

- **regular population** of newly generated solutions, and
- **archive**, which contains a representation of the nondominated front among all solutions considered so far.

The archive size is fixed, i.e., whenever the number of nondominated individuals is less than the predefined archive size, the archive is filled up by *good* dominated individuals.

A **truncation method** is invoked when the nondominated front exceeds the archive limit.

A member of the archive is only removed if

1. a solution has been found that dominates it or
2. the maximum archive size is exceeded and the portion of the front where the archive member is located is overcrowded.

Using the archive makes it possible not to lose certain portions of the current nondominated front due to random effects.

All individuals in the archive participate in selection.

SPEA2: Algorithm

Input: N is the population size, \bar{N} is the archive size.

1. **Initialization:** Generate an initial population P_0 and create the empty archive $\bar{P}_0 = \emptyset$. Set $t = 0$.
2. **Fitness assignment:** Calculate fitness of individuals in P_t and \bar{P}_t .
3. **Environmental selection:** Copy all nondominated individuals in P_t and \bar{P}_t to \bar{P}_{t+1} .
If size of \bar{P}_{t+1} exceeds \bar{N} then reduce \bar{P}_{t+1} using the truncation operator.
If size of \bar{P}_{t+1} is less than \bar{N} then fill \bar{P}_{t+1} with dominated solutions in P_t and \bar{P}_t .
4. **Termination:** If $t \geq T$ then return nondominated solutions in \bar{P}_{t+1} . Stop.
5. **Mating selection:** Perform binary tournament selection with replacement on \bar{P}_{t+1} in order to fill the mating pool.
6. **Variation:** Apply recombination and mutation operators to the mating pool and fill P_{t+1} with the generated solutions,
increment generation counter $t = t + 1$,
go to Step 2.



SPEA2: Fitness Assignment

Fitness assignment (fitness is to minimized) – for each individual both dominating and dominated solutions are taken into account.

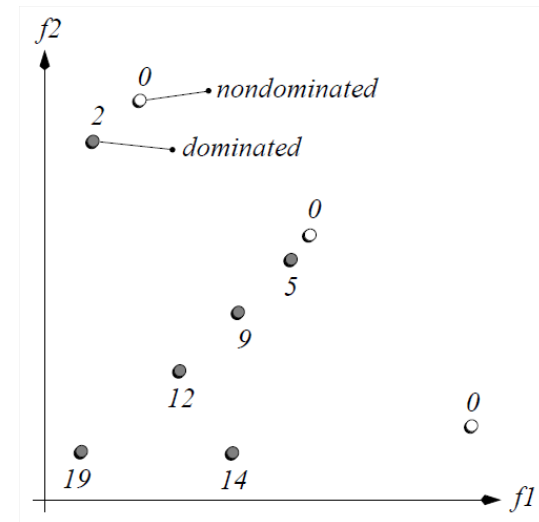
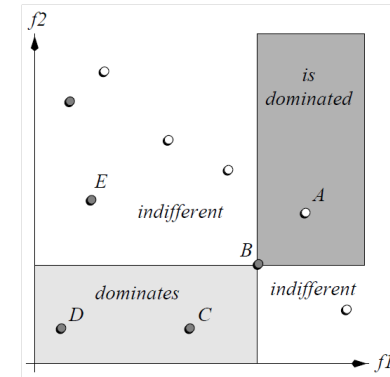
- Each individual i in the archive \bar{P}_t and the population P_t is assigned a **strength value** $S(i)$, representing the number of solutions it dominates.
- The raw fitness $R(i)$ of an individual i is calculated as

$$R(i) = \sum_{j \in P_t + \bar{P}_t, j \succ i} S(j)$$

that is $R(i)$ is determined by the strengths of its dominators in both archive and population.

$R(i) = 0$ corresponds to a nondominated solution.

Since the **raw fitness assignment** is based on the concept of Pareto dominance, it **may fail when most individuals do not dominate each other**.



SPEA2: Density Estimation

Density information is incorporated to discriminate between individuals having identical raw fitness values.

The density at any point is estimated as a (decreasing) function of the distance to the k -th nearest data point – calculated as the inverse of the distance to the k -th nearest neighbor.

- k equal to the square root of the sample size is used: $k = \sqrt{N + \bar{N}}$.
- Density $D(i)$ is calculated as

$$D(i) = \frac{1}{\sigma_i^k + 2}$$

where σ_i^k is the distance to the k -th nearest neighbor and it is made sure that $D(i) < 1$.

Final fitness is given as

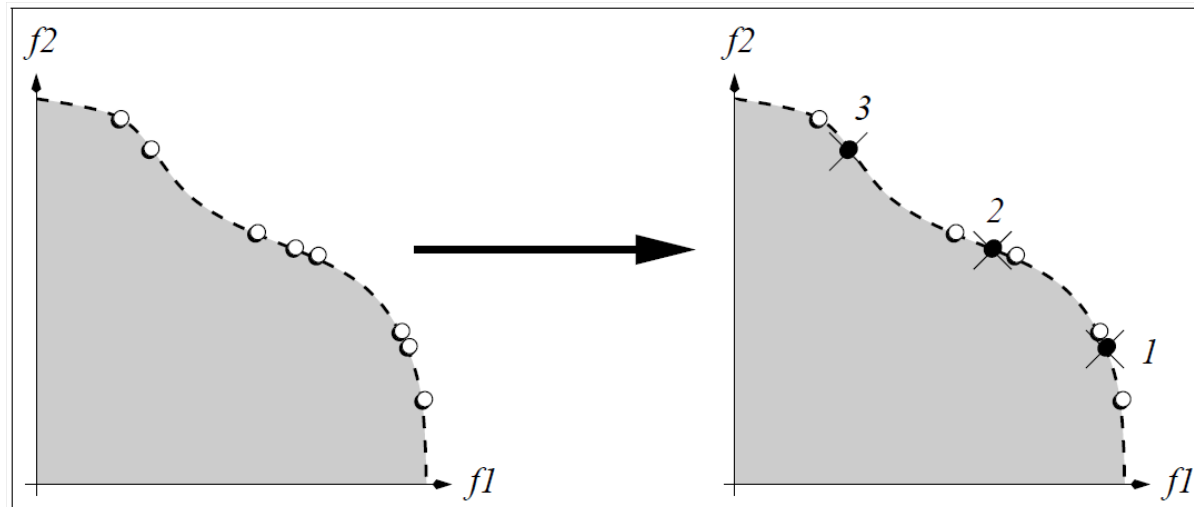
$$F(i) = R(i) + D(i)$$

SPEA2: Environmental Selection

If after copying all nondominated individuals from archive and population to the archive of the next generation

- the archive is too small (i.e. $|\bar{P}_{t+1}| < \bar{N}|$), the best $\bar{N} - |\bar{P}_{t+1}|$ dominated solutions (w.r.t. fitness) in the previous archive and population are copied to the new archive;
- the archive is too large (i.e. $|\bar{P}_{t+1}| > \bar{N}|$), individuals from \bar{P}_{t+1} are iteratively removed until $|\bar{P}_{t+1}| = \bar{N}$.

At each iteration, the individual which has the minimum distance to another individual is chosen (a tie is broken by considering the second smallest distances and so forth).



SPEA2: Conclusions

SPEA2

- uses the concept of **Pareto dominance** in order to assign scalar fitness values to individuals;
- uses a **fine-grained fitness** assignment strategy which **incorporates density information** in order to distinguish between solutions that are indifferent, i.e., do not dominate each other;
- uses environmental selection in order to keep the optimal diversity in the archive;
- seems to have advantages over NSGA-II in higher dimensional objective spaces.

MOEA Performance Measures

The result of a MOEA run is not a single scalar value, but a collection of vectors forming a non-dominated set.

- Comparing two MOEA algorithms requires comparing the non-dominated sets they produce. However, there is no straightforward way to compare different non-dominated sets.

Three goals that can be identified and measured:

1. The distance of the resulting non dominated set to the Pareto-optimal front should be minimized.
2. A good (in most cases uniform) distribution of the solutions found is desirable.
3. The extent of the obtained non dominated front should be maximized, i.e., for each objective, a wide range of values should be present.

S Metric cond.

Pros:

- Given two non-dominated sets, A and B , if each point in B is dominated by a point in A then A will always be evaluated as being better than B .
- Independent – the hypervolume calculated for the given set is not dependent on any other, or any reference set.
- Differentiates between different degrees of complete outperformance of two sets.
- Intuitive meaning/interpretation.

Cons:

- Requires defining some upper boundary of the region.
This choice does affect the ordering of non-dominated sets.
- It has a large computational overhead, $O(n^{k+1})$, where n is the number of nondominated solutions and k is the number of objectives, rendering it unusable for many objectives or large sets.
- It multiplies apples by oranges, that is, different objectives together.

Reading

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