



中山大學
SUN YAT-SEN UNIVERSITY

Lecture 8

Multi-objective Optimization

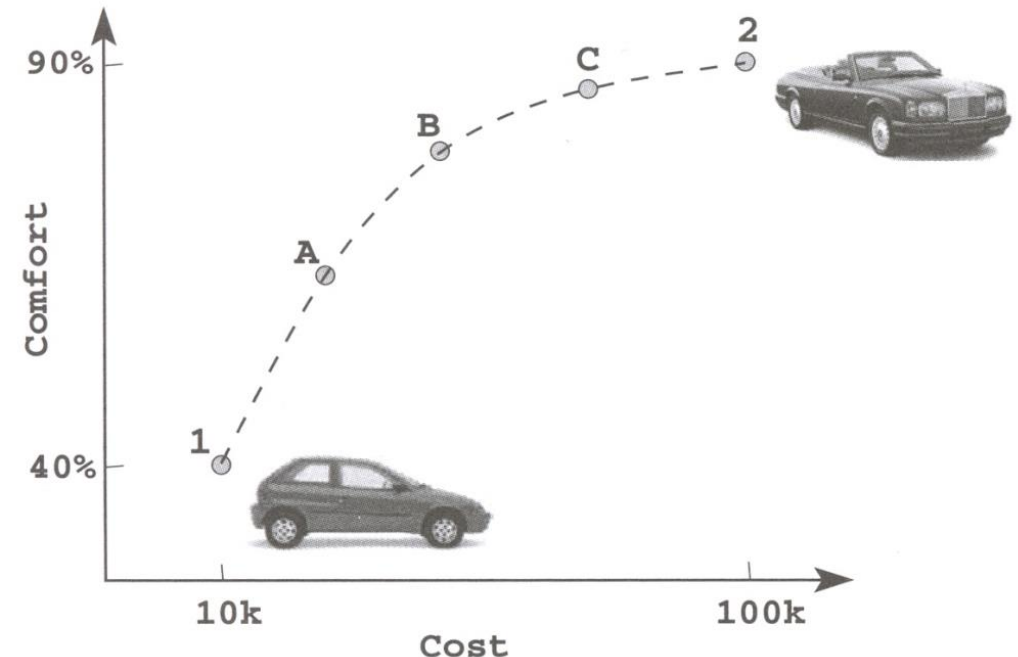
Algorithm

张子臻，中山大学计算机学院

zhangzizhen@gmail.com

Introduction

- Many real-world problems involve multiple objectives.
- These objectives are generally **conflicting** with each other.
 - A solution that is **extreme** with respect to one objective requires a **compromise** in other objectives.
 - A **sacrifice** in one objective is related to the **gain** in other objective(s).
- Illustrative example: Buying a car
 - Two extreme hypothetical cars 1 and 2.
 - Cars with a trade-off between cost and comfort: A, B, and C.
- Which solution out of all of the trade-off solutions is the best with respect to all objectives?
 - Without any further information those **trade-offs** are indistinguishable.
 - Thus, a number of optimal solutions is sought in multiobjective optimization.



Multi-objective Optimization

- A multi-objective optimization problem (MOOP) with M objectives is defined as follows: Given an n -dimensional decision variable vector $x = \{x_1, \dots, x_n\}$ in the solution space X , find a vector x that minimizes/maximizes a given set of M objective functions: $f(x) = \{f_1(x), \dots, f_M(x)\}$.
- The solution space X is generally restricted by a series of constraints.
- MOOP in the **general form**:

$$\begin{aligned}
 &\min f_m(x) && m = 1, 2, \dots, M \\
 &s.t \\
 &g_j(x) \geq 0 && j = 1, 2, \dots, J \\
 &h_k(x) = 0 && k = 1, 2, \dots, K \\
 &x_i^L \leq x_i \leq x_i^U && i = 1, 2, \dots, n
 \end{aligned}$$

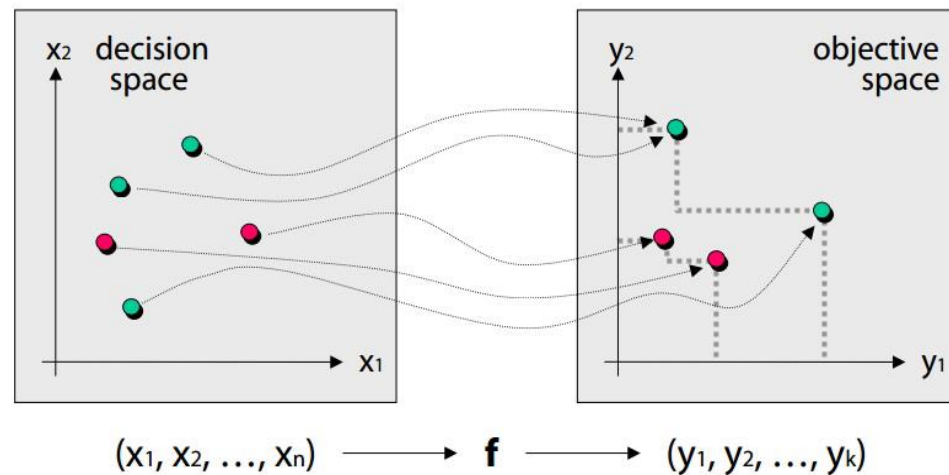
Definition

- (**Domination**) A feasible solution x is said to (weak) dominate another feasible solution y (or $x \preceq y$), if both the following condition hold:
 1. $f_i(x) \leq f_i(y), \forall i = 1, \dots, M.$
 2. $f_i(x) < f_i(y), \exists i = 1, \dots, M.$
- (**Non-dominated set**) Among a set of feasible solutions P , the non-dominated set of solutions P' are those solutions that are not dominated by any member of set P .
- (**Pareto optimal set**) The non-dominated set of the entire feasible search space S is called (globally) Pareto optimal set.
- (**Pareto front**) For a given Pareto optimal set, the corresponding objective function values in the objective space are called the Pareto front.

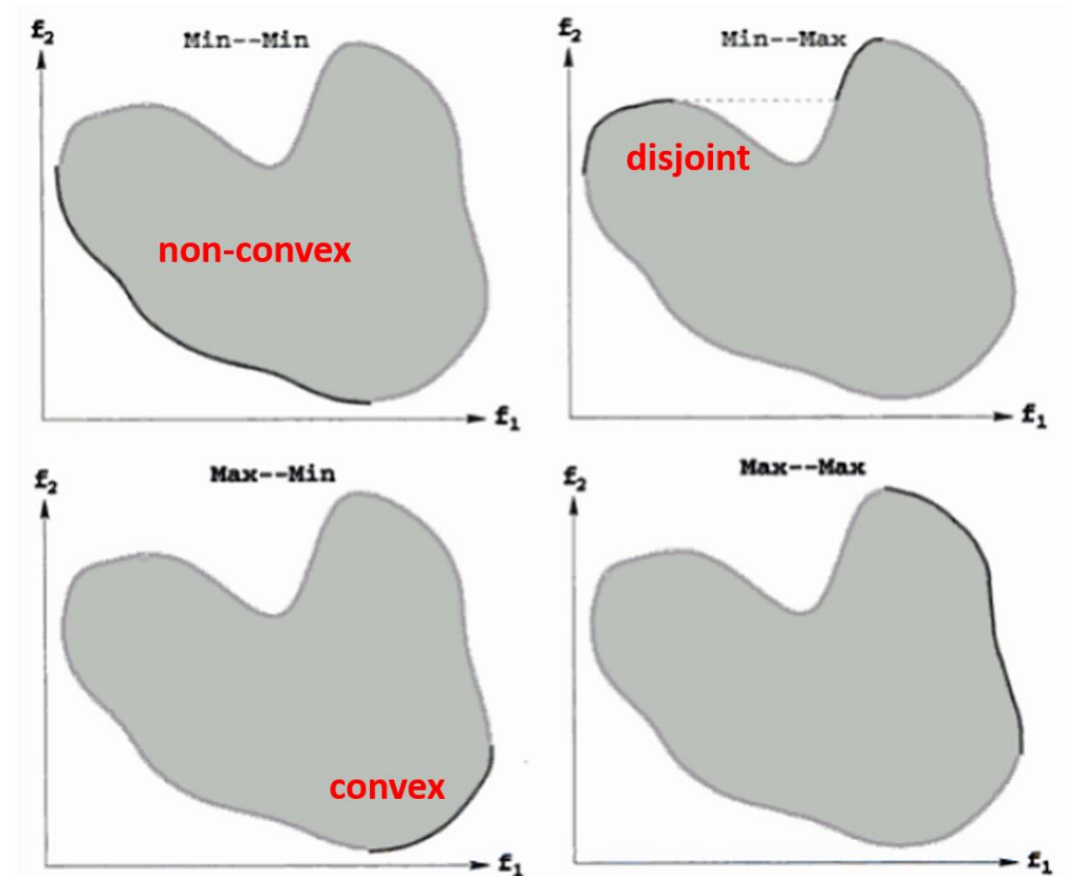
Pareto Optimality

- Pareto set & Pareto front

Pareto set ● Pareto front
 Pareto set approximation ● Pareto front approximation

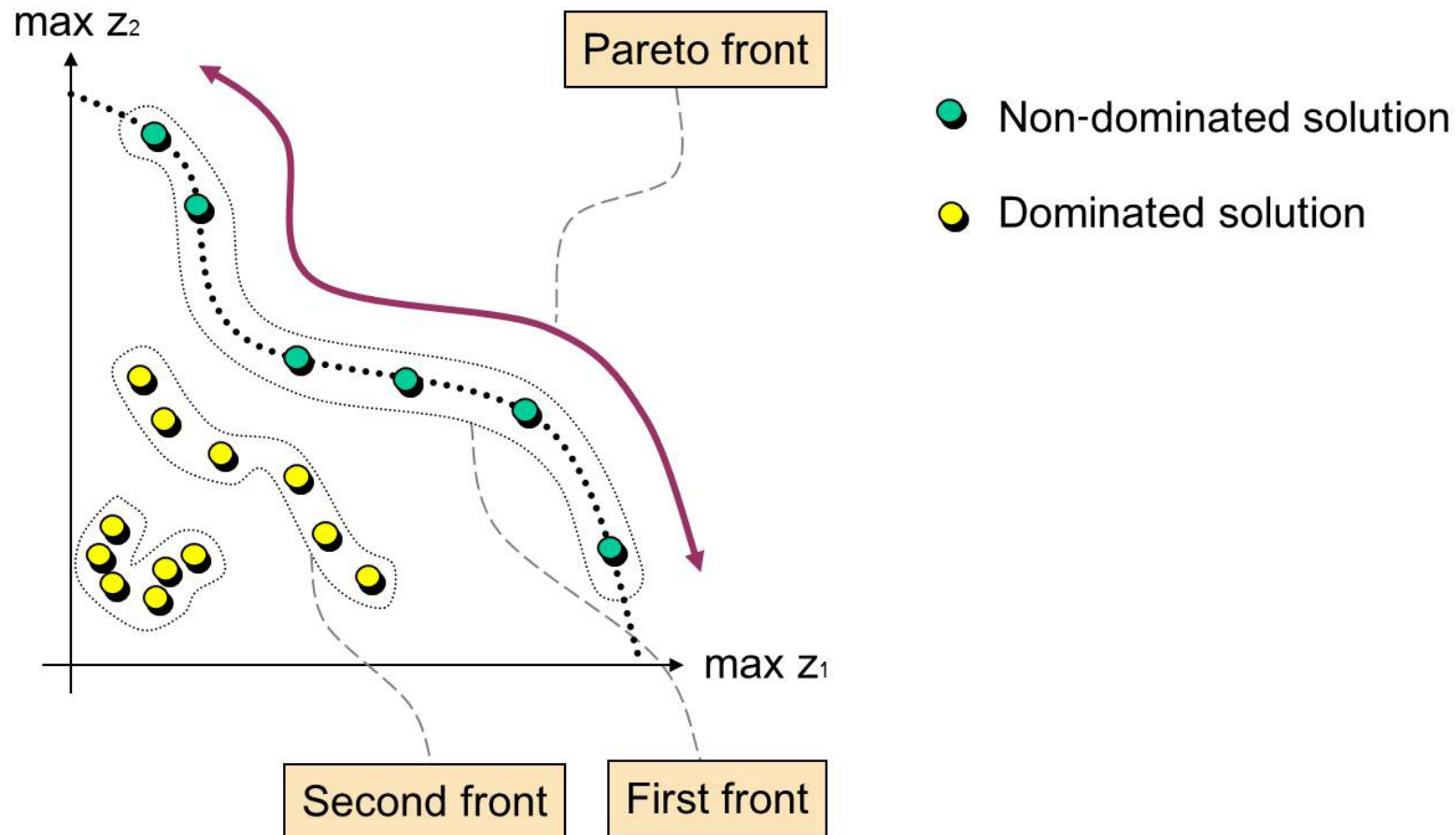


- Shapes of the Pareto front



Pareto Optimality

- Fronts can be classified into different levels.



Non-dominated Sorting

- The non-dominated sorting is to classified solutions into different levels.
- It is essentially a **topological sorting** process.

An $O(MN^2)$ non-dominated sorting algorithm

```

INPUT: Feasible solution set  $P$ ;
for each  $p \in P$  do
     $S_p = \emptyset$ ;
     $n_p = 0$ ;
    for each  $q \in P$  do
        if  $p \preceq q$  then
             $S_p = S_p \cup \{q\}$ ;
        else if  $q \preceq p$  then
             $n_p = n_p + 1$ ;
        end if
    end for
    if  $n_p = 0$  then
         $p_{rank} = 1$ ;
         $F_1 = F_1 \cup \{p\}$ ;
    end if
end for

 $i = 1$ ;
while  $F_i \neq \emptyset$  do
     $Q = \emptyset$ ;
    for each  $p \in F_i$  do
        for each  $q \in S_p$  do
             $n_q = n_q - 1$ ;
            if  $n_q = 0$  then
                 $q_{rank} = i + 1$ ;
                 $Q = Q \cup \{q\}$ ;
            end if
        end for
    end for
     $i = i + 1$ ;
     $F_i = Q$ ;
end while
  
```

Classical Method: Weighted Sum Method

- The Weighted sum method scalarizes a set of objectives into a **single objective** by **pre-multiplying** each objective with a **user-defined** weight.

$$\min F(x) = \sum_{m=1}^M w_m f_m(x)$$

$$\text{subject to } g_j(x) \geq 0 \quad j = 1, 2, \dots, J$$

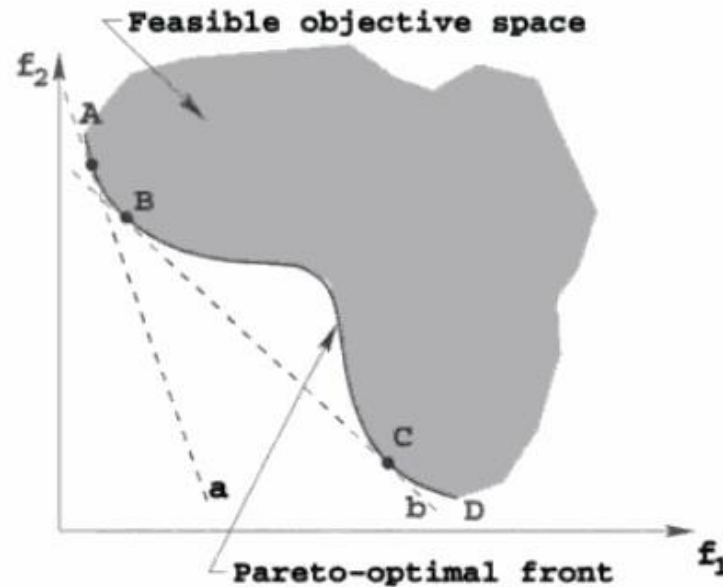
$$h_k(x) = 0 \quad k = 1, 2, \dots, K$$

$$x_i^L \leq x_i \leq x_i^U \quad i = 1, 2, \dots, n$$

- Q:** How to set up an appropriate weight vector w ?
- A:** By relative importance of objectives in the problem.
- Different objectives take different orders of magnitude. Sometimes we need a process to **normalize** each objective.

Weighted Sum Method

- Theorem: If the weight is positive for all objectives, the solution to the problem represented by weighted sum model is Pareto-optimal.
- If x is a Pareto-optimal solution of a convex multi-objective optimization problem, then there exists a non-zero positive weight vector w such that x is a solution to the problem given by weighted sum model.



Classical Method: Weighted Metric Method

$$\min l_p(x) = \left(\sum_{m=1}^M w_m |f_m(x) - z_m^*|^p \right)^{1/p}$$

$$\text{subject to } g_j(x) \geq 0$$

$$j = 1, 2, \dots, J$$

$$h_k(x) = 0$$

$$k = 1, 2, \dots, K$$

$$x_i^L \leq x_i \leq x_i^U$$

$$i = 1, 2, \dots, n$$

- When $p = 1$ is used, the resulting problem is equivalent to the **weighted sum approach**.
- When $p = 2$ is used, a **weighted Euclidean distance** of any point in the objective space from the *ideal* (or *utopian*) point z^* is minimized.
- When $p = \infty$, the problem is called **weighted Tchebyche problem**.

$$\min l_\infty(x) = \max_{m=1, \dots, M} w_m |f_m(x) - z_m^*|$$

$$\text{subject to } g_j(x) \geq 0$$

$$j = 1, 2, \dots, J$$

$$h_k(x) = 0$$

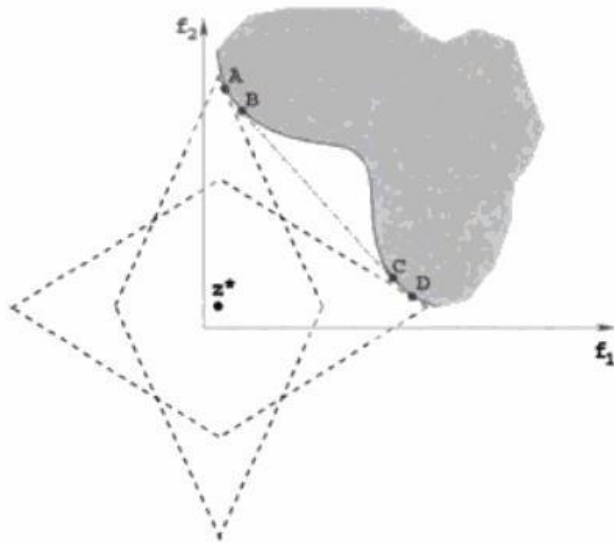
$$k = 1, 2, \dots, K$$

$$x_i^L \leq x_i \leq x_i^U$$

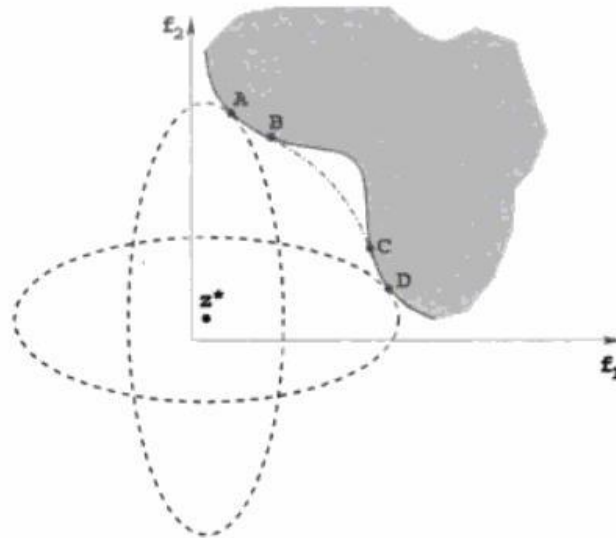
$$i = 1, 2, \dots, n$$

Weighted Metric Method

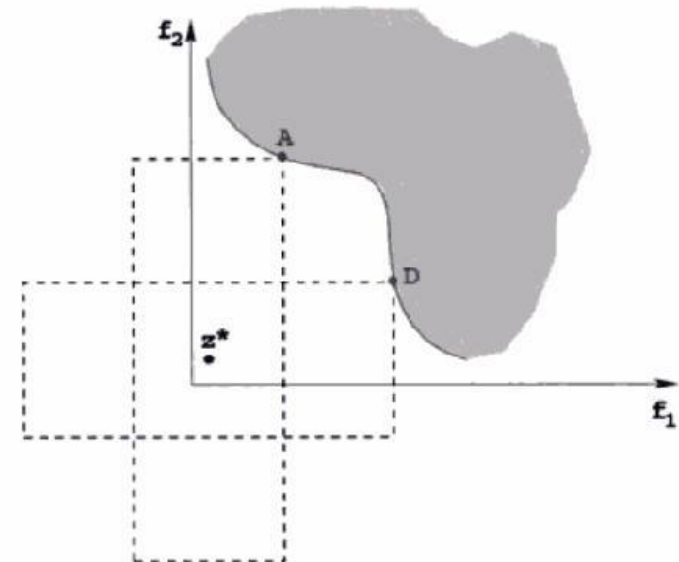
- Let x^* be a Pareto-optimal solution. Then there exists a positive weighting vector such that x is a solution of the weighted Tchebyche problem, where the reference point is the **ideal** (utopian) objective vector.



The weighted metric method with $p = 1$.



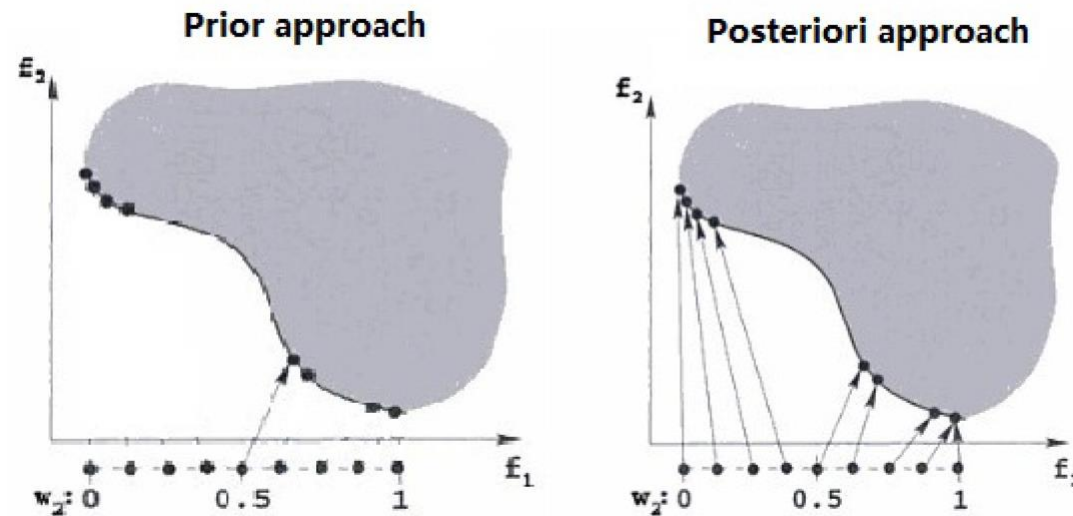
The weighted metric method with $p = 2$.



The weighted metric method with $p = \infty$.

Motivation for Finding Multiple Pareto-optimal Solutions

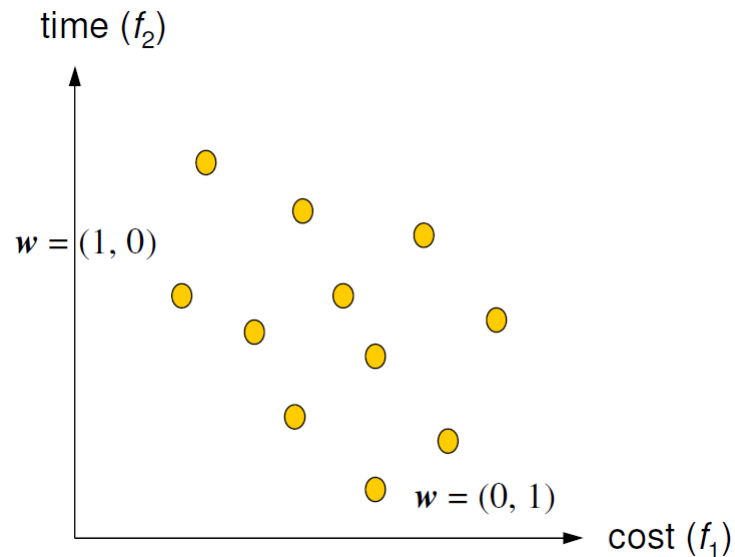
- **Q:** What should we do with multiple Pareto-optimal solution?
 - **A:** From a practical point of view, we only need one solution to implement.
 - **Q:** Which one of these multiple solutions are we interested in?
 - **A:** It depends on the trade-off relationship among objectives.
-
- A priori approach v.s. Posterior approach



A Priori Method

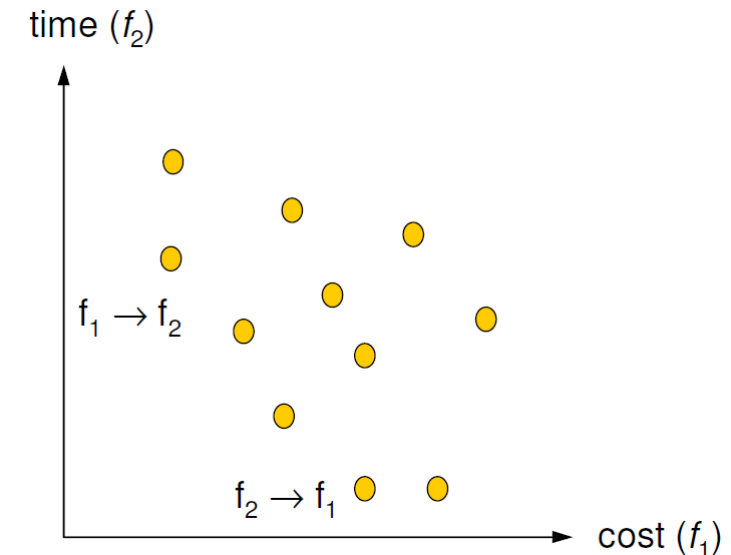
- A priori method - Aggregation

$$\begin{array}{ll} \text{minimize} & \sum_{i=1}^k w_i f_i(\mathbf{x}) \\ \text{subject to} & \mathbf{x} \in S, \end{array}$$



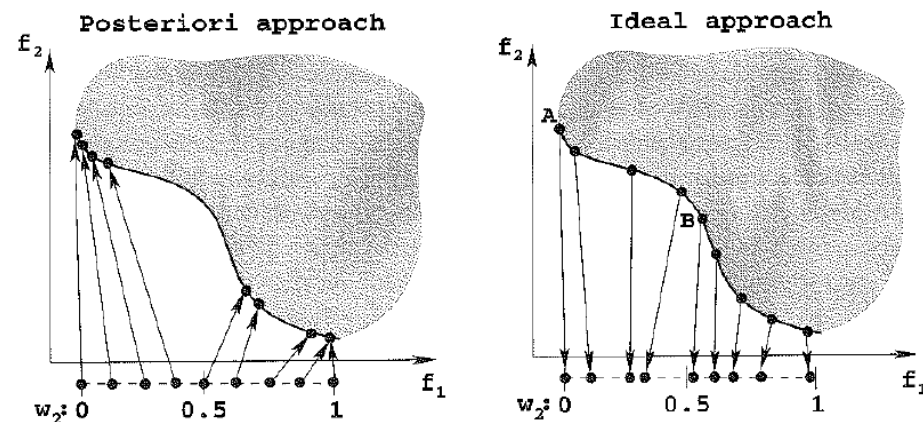
- A priori method - Lexicographical ordering

- Optimize f_1 primarily and f_2 secondarily
- Optimize f_2 primarily and f_1 secondarily



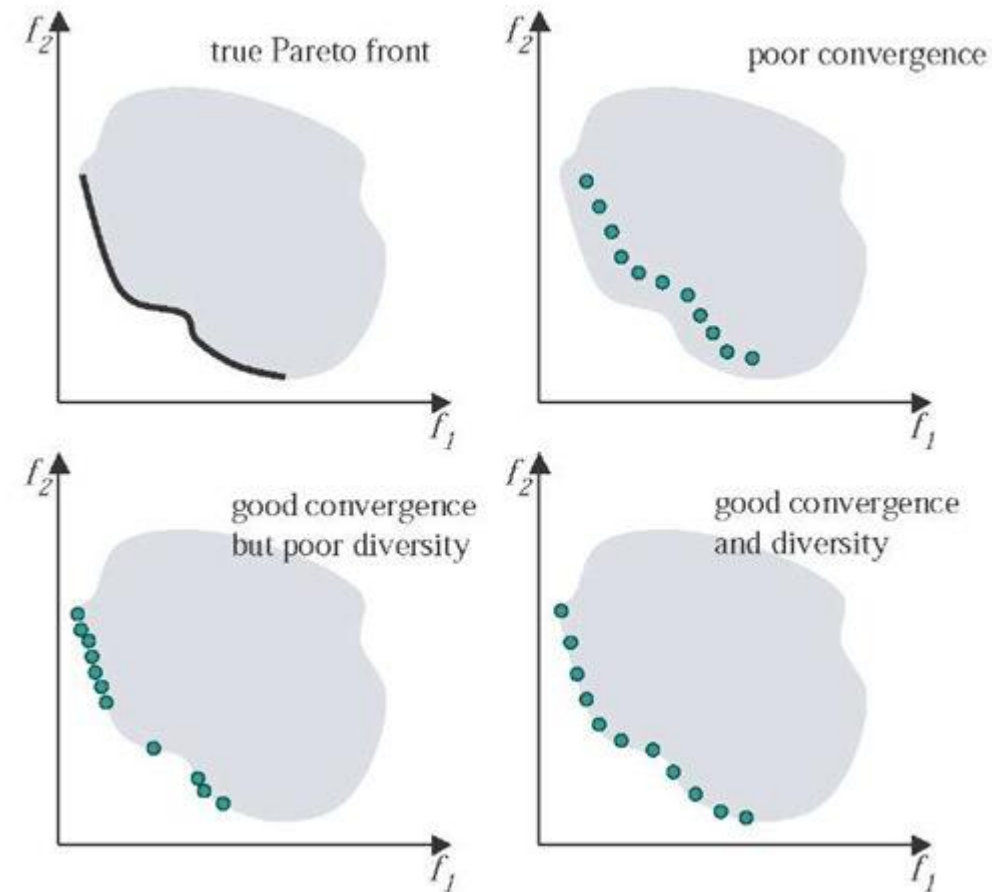
Posteriori Method

- Disadvantages of a priori methods
 - It is difficult to give the preference information in advance.
 - The decision maker needs to adjust the preference to obtain alternative solutions.
 - The condition becomes harder when there are multiple decision makers.
- Posteriori method
 - Find a set of Pareto optimal solutions first.
 - Choose one solution from the set by using other high-level information.
- Classical posteriori approach and the ideal approach



Goals of Multiobjective Optimization

- Goals of MOP are also multiobjective.
 - (Quality, Convergence, Proximity) To find a set of solutions **as close as possible** to the Pareto-optimal set.
 - (Diversity, Spread) To find a set of solutions **as diverse as possible**.

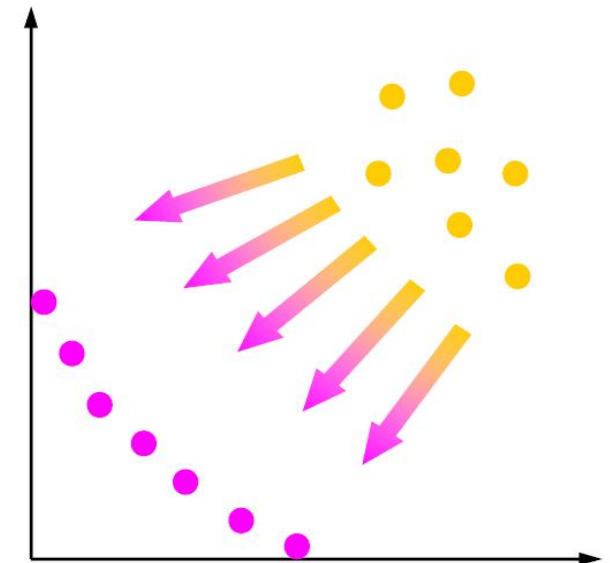


Multiobjective Optimization using Evolutionary Algorithm

- **Evolutionary algorithm** is a good option in achieving these goals.
- In general, an evolutionary algorithm is a **population-based** meta-heuristic optimization algorithm.

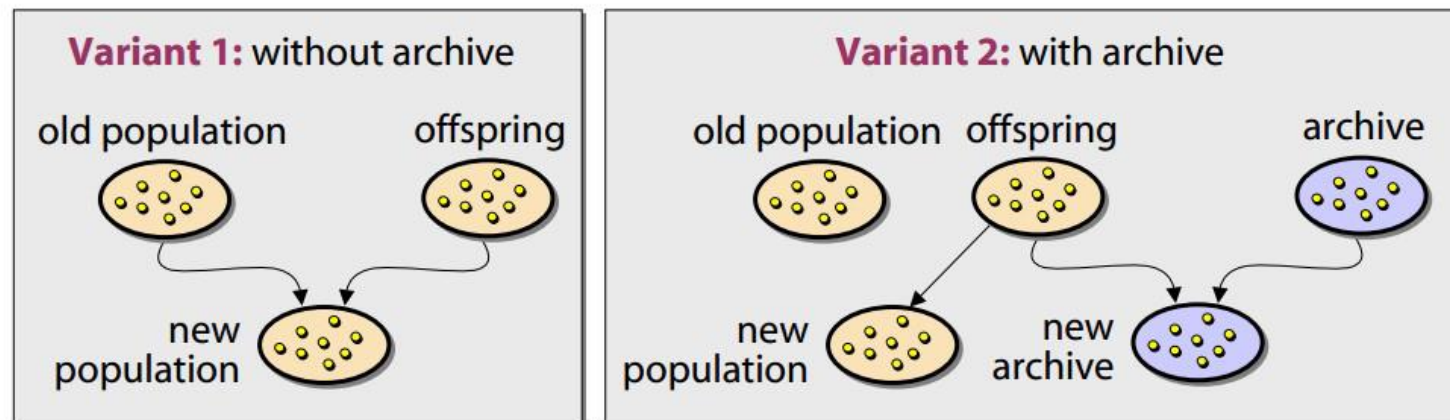
A Generic Evolutionary Algorithm Framework

```
Initialize population  $P_0$ ;  
 $t = 0$ ;  
while Termination criteria not reached do  
    Parent_selection();  
    Recombination();  
    Mutation();  
    Assign_fitness();  
    Selection();  
     $t = t + 1$ ;  
end while
```



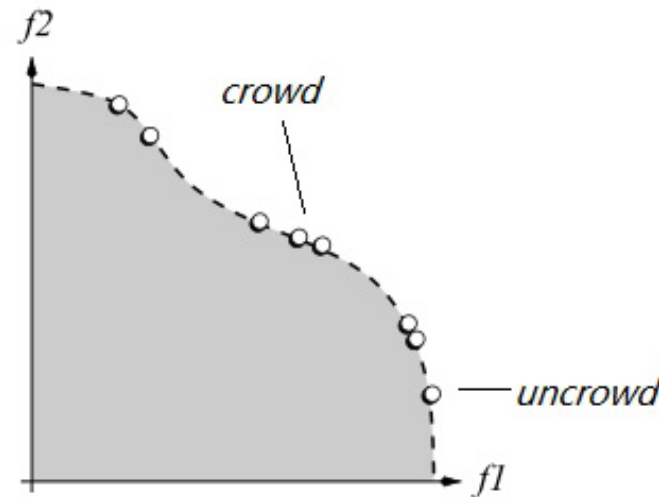
Challenges

- Q: How to guide search towards the Pareto-optimal front?
- A: Use Environmental selection. **Elitism** is important in evolutionary algorithms.
- Q: How to maintain the diversity of the population?
- A: Use **Density estimation** approaches.
- Q: How to preserve Pareto-optimal solutions?
- A: Use Population or Population + Archive strategy.



Density Estimation

- To maintain a diverse Pareto-optimal set, density estimation of individual in the population is introduced.



- Approaches:
 - Sharing function approach
 - Crowding distance approach
 - k-th nearest neighbor approach

Sharing Function

- Idea: diversity in the population is preserved by **degrading the fitness** of similar solutions.
- The algorithm of calculating the shared fitness value of i-th individual in the population of size N :

- Calculate **sharing function value** with all solutions in the population according to

$$Sh(d_{ij}) = \begin{cases} 1 - (\frac{d_{ij}}{\sigma_{share}})^\alpha, & \text{if } d_{ij} \leq \sigma_{share} \\ 0, & \text{otherwise.} \end{cases}$$

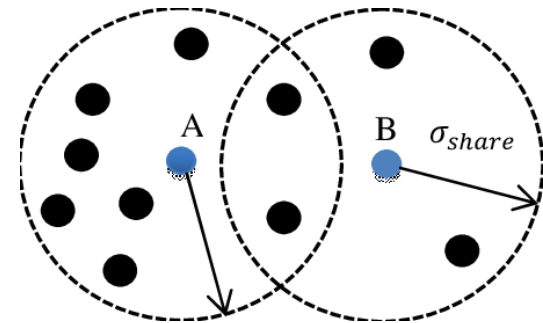
- Calculate **niche count** nc_i as follows

$$nc_i = \sum_{j=1}^N Sh(d_{ij})$$

- Calculate **shared fitness** as

$$f'_i = f_i / nc_i$$

- If $d = 0$ then $Sh(d) = 1$ meaning that two solutions are identical. If $d \geq \sigma_{share}$ then $Sh(d) = 0$ meaning that two solutions do not have any sharing effect on each other.



Density Estimation

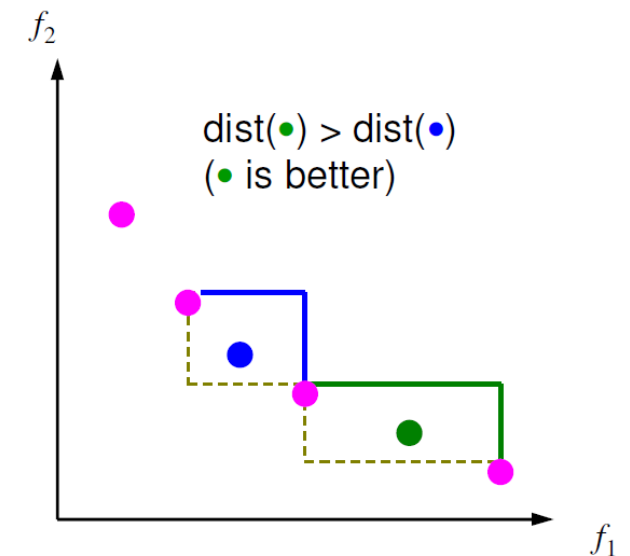
- Crowding distance approach
 - To get an estimate of the density of solutions surrounding a particular solution in the population, calculate the **average distance of two points** on either side of this point along each of the objectives.
 - This quantity i_{distance} serves as an estimate of the **perimeter of the cuboid** formed by using the nearest neighbors as the vertices (called **crowding distance**).

Crowding distance assignment

```

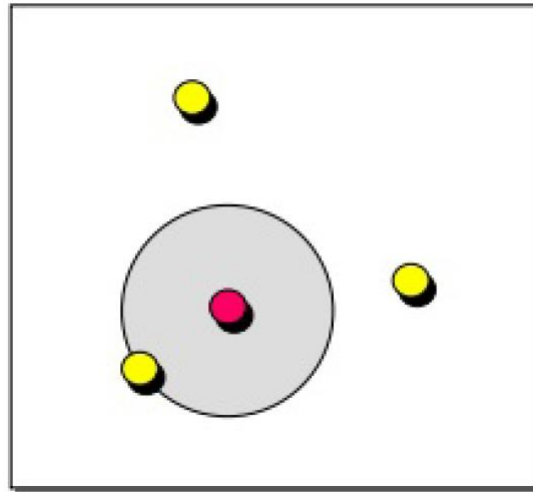
INPUT: Non-dominated set  $\mathcal{I}$ 
 $l = \mathcal{I}$ 
for each  $i$  do
  set  $\mathcal{I}[i]_{\text{distance}} = 0$ ;
end for
for each objective  $m$  do
   $\mathcal{I} = \text{sort}(\mathcal{I}, m)$ ;
   $\mathcal{I}[1]_{\text{distance}} = \mathcal{I}[l]_{\text{distance}} = \infty$ ;
  for  $i = 2$  to  $l - 1$  do
     $\mathcal{I}[i]_{\text{distance}} = \mathcal{I}[i]_{\text{distance}} + (\mathcal{I}[i + 1].m - \mathcal{I}[i - 1].m) / (f_m^{\max} - f_m^{\min})$ ;
  end for
end for

```



Density Estimation

- k-th nearest neighbor approach
 - For each individual i , the distances (in objective space) to all individuals j are calculated and stored in a list.
 - After sorting the list in increasing order, the k -th element gives the distance, denoted as k_i .



MOEA Approaches

- Weighted-Based Genetic Algorithm (WBGA)
- Non-dominated Sorting Genetic Algorithm II (NSGA-II)
- Strength Pareto Evolutionary Algorithm II (SPEA-II)
- Multiobjective Optimization Evolutionary Algorithm with Decomposition (MOEA/D)
- ...

Weighted-Based Genetic Algorithm (WBGA)

- The framework of WBGA is very similar to EA procedure.
- Consider a multi-objective maximization problem. The user first need to assign a set of **weight vector** w . **Normalize** each weight: $\bar{w}_i = \frac{w_i}{\sum_{j=1}^M w_j}$

x_w	1	2	3	4	5
Weight vector	(0.1,0.9)	(0.3,0.7)	(0.5,0.5)	(0.7,0.3)	(0.9,0.1)

- Each individual $x^{(i)}$ in the population is associated with a weight vector.

Individual	$x^{(1)}$	$x^{(2)}$	$x^{(3)}$	$x^{(4)}$	$x^{(5)}$	$x^{(6)}$	$x^{(7)}$	$x^{(8)}$
x_w	3	1	2	4	2	5	1	3

- Assign **fitness** to individuals: $F(x^{(i)}) = \sum_{j=1}^M w_j^{x_w^{(i)}} \frac{f_j(x^{(i)}) - f_j^{min}}{f_j^{max} - f_j^{min}}$
- Use the Sharing function to calculate the **niche count** nc_i for individual $x^{(i)}$
- Calculate the **shared fitness**: $F'_i = F_i / nc_i$

Non-dominated Sorting Genetic Algorithm II (NSGA-II)

- Fast non-dominated sorting approach
 - Computational complexity: $O(MN^2)$.
- Diversity preservation: crowded comparison approach
 - Every solution in the population has two attributes:
 1. non-domination rank (i^{rank})
 2. crowding distance ($i^{distance}$)
 - Crowded comparison operator: A partial order \prec_n is defined as:
$$i \prec_n j \text{ if } (i^{rank} < j^{rank}) \text{ or } ((i^{rank} = j^{rank}) \text{ and } (i^{distance} > j^{distance}))$$
- Elitist evolutionary model
 - Only the best solutions survive to subsequent generations.

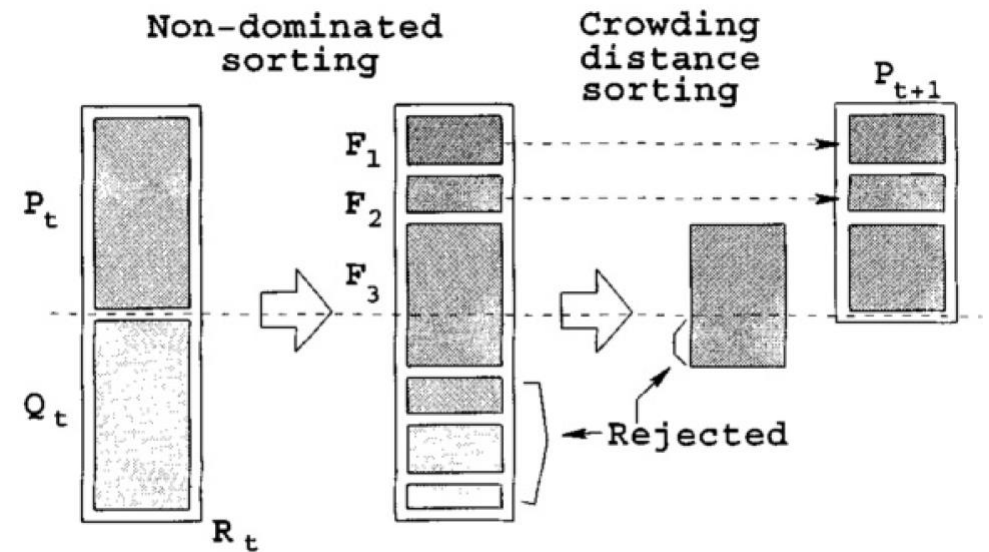
Non-dominated Sorting Genetic Algorithm II (NSGA-II)

- The overall complexity of the algorithm in each iteration is $O(MN^2)$.

NSGA-II in t -th iteration

```

 $Q_t = \text{make-new-pop}(P_t);$ 
 $R_t = P_t \cup Q_t;$ 
 $\mathcal{F} = \text{fast-non-dominated-sort}(R_t);$ 
 $P_{t+1} = \emptyset$  and  $i = 1;$ 
while  $|P_{t+1}| + |\mathcal{F}_i| \leq N$  do
    crowding-distance-assignment $(\mathcal{F}_i);$ 
     $P_{t+1} = P_{t+1} \cup \mathcal{F}_i;$ 
     $i = i + 1;$ 
end while
Sort $(\mathcal{F}_i, \prec_n);$ 
 $P_{t+1} = P_{t+1} \cup \mathcal{F}_i[1 : (N - |P_{t+1}|)];$ 
 $t = t + 1$ 
  
```



NSGA-II procedure

NSGA-II as a Constraint Handling Approach

- 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.
- Constrained-domination: A solution i is said to **constrained-dominate** a solution j , if any of the following conditions is true
 1. Solution i is feasible and solution j is not.
 2. Solutions i and j are both infeasible, but solution i has a smaller overall constraint violation.
 3. Solutions i and j are feasible, and solution i dominates solution j .
- **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.

MOEA/D

- Multiobjective Optimization Evolutionary Algorithm with Decomposition, “D” for problem **Decomposition**.
- The MOEA/D decomposes a multi-objective optimization problem into N scalar optimization subproblems.
- These subproblems are optimized **simultaneously** from the evolution of a population of solutions.
- In each generation, the population is composed of the best solutions found for each subproblem.
- Each subproblem is optimized considering only **neighboring** sub-problem information.
- It has low computational complexity compared to popular methods: NSGA-II, MOGLS.

Decomposition

- Weight vector-based aggregation function
 - Linear weighted sum
 - Tchebycheff
 - Boundary intersection
 - Penalty-based Boundary Intersection
- Every weight vector defines a subproblem, and every subproblem keeps the best solution.
 - A large number of weight vector
 - A very small sub-population (typically, the size is 1)

MOEA/D Framework

Input:

- MOP (1);
- a stopping criterion;
- N : the number of the subproblems considered in MOEA/D;
- a uniform spread of N weight vectors: $\lambda^1, \dots, \lambda^N$;
- T : the number of the weight vectors in the neighborhood of each weight vector.

Output: EP.

Step 1.1) Set $EP = \emptyset$.

Step 1.2) Compute the Euclidean distances between any two weight vectors and then work out the T closest weight vectors to each weight vector. For each $i = 1, \dots, N$, set $B(i) = \{i_1, \dots, i_T\}$, where $\lambda^{i_1}, \dots, \lambda^{i_T}$ are the T closest weight vectors to λ^i .

Step 1.3) Generate an initial population x^1, \dots, x^N randomly or by a problem-specific method. Set $FV^i = F(x^i)$.

Step 1.4) Initialize $z = (z_1, \dots, z_m)^T$ by a problem-specific method.

For $i = 1, \dots, N$, do

Step 2.1) Reproduction: Randomly select two indexes k, l from $B(i)$, and then generate a new solution y from x^k and x^l by using genetic operators.

Step 2.2) Improvement: Apply a problem-specific repair/improvement heuristic on y to produce y' .

Step 2.3) Update of z : For each $j = 1, \dots, m$, if $z_j < f_j(y')$, then set $z_j = f_j(y')$.

Step 2.4) Update of Neighboring Solutions: For each index $j \in B(i)$, if $g^{te}(y'|\lambda^j, z) \leq g^{te}(x^j|\lambda^j, z)$, then set $x^j = y'$ and $FV^j = F(y')$.

Step 2.5) Update of EP:

Remove from EP all the vectors dominated by $F(y')$.

Add $F(y')$ to EP if no vectors in EP dominate $F(y')$.

Aggregation Vectors

- Simplex-Lattice Design

$$\lambda_j^i \in \left\{ \frac{0}{H}, \frac{1}{H}, \dots, \frac{H}{H} \right\}, \forall i = 1, \dots, N, j = 1, \dots, M$$

where M is the number of objectives, H is a positive integer (user-defined).

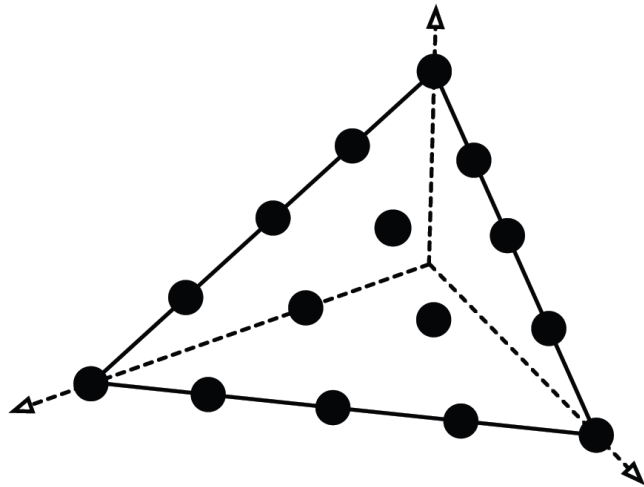
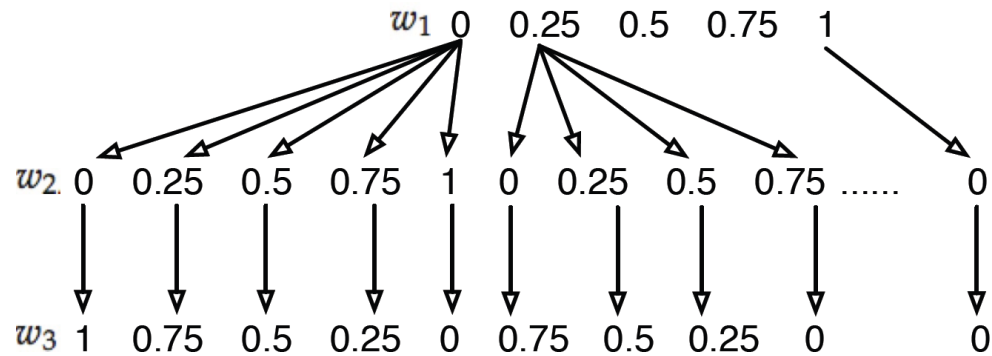
- The number of vectors (equal to the size of the population) is given by:

$$N = C_{H+M-1}^{M-1}$$

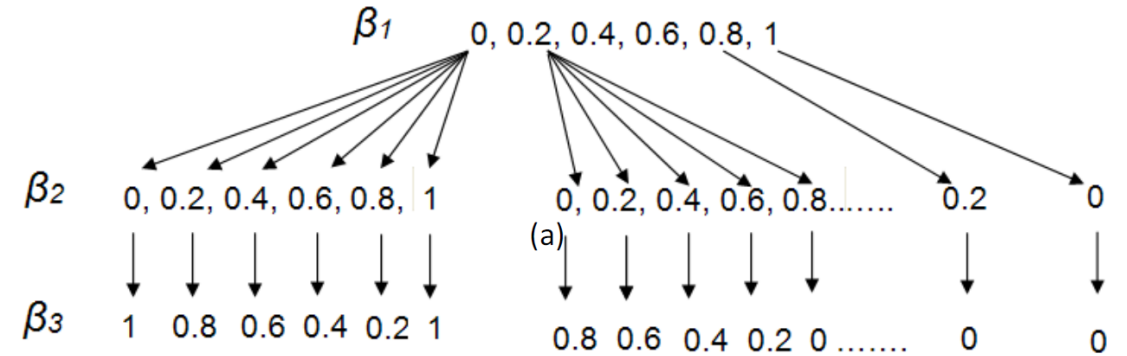
- For example, assume that M = 3 objectives and different H values, then
 - $H = 1, N = C_3^2 = 3, \lambda \in \{(1,0,0), (0,1,0), (0,0,1)\}$
 - $H = 2, N = C_4^2 = 6, \lambda \in \{(1,0,0), (0,1,0), (0,0,1), (0,1/2,1/2), (1/2,0,1/2), (1/2,1/2,0)\}$
 - $H = 3, N = C_5^2 = 10, \lambda \in \{(1,0,0), (0,1,0), (0,0,1), (0,1/3,2/3), (1/3,0,2/3), (1/3,2/3,0), (0,2/3,1/3), (2/3,0,1/3), (2/3,1/3,0), (1/3,1/3,1/3)\}$

Aggregation Vectors

- $M = 3, H = 4$



- $M = 3, H = 5$



Performance Measures of MOOP

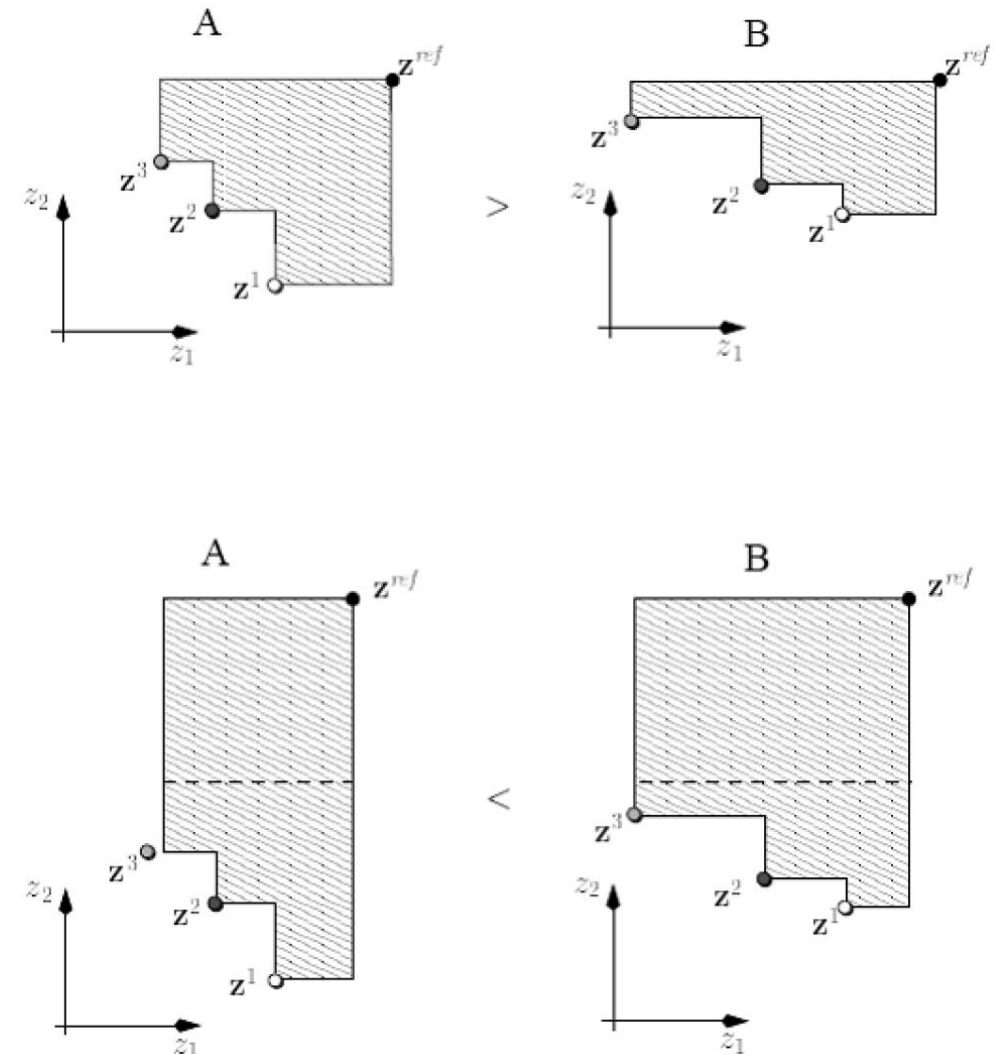
- The result of a MOEA run is not a single scalar value, but **a collection of vectors** forming a non-dominated set.
- Comparing two MOEA algorithms requires comparing the **non-dominated sets** they produce. However, there is no straightforward way to compare different non-dominated sets.
- Three goals 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.

Performance Metrics

- Number of fronts
- Generational distance (GD)
- Spacing (SP)
- Spread
- Hypervolume (HV) (*S-metric*)
- Coverage (*C-metric*)
- Inverted Generational Distance (IGD) (*D-metric*)
- Epsilon indicator

S-Metric

- It calculates the **hypervolume** of the multi-dimensional region enclosed by a set A and a reference point Z^{ref} .
- The hypervolume expresses the size of the region that is dominated by A .
- The bigger the value of this measure the better the quality of A is, and vice versa.



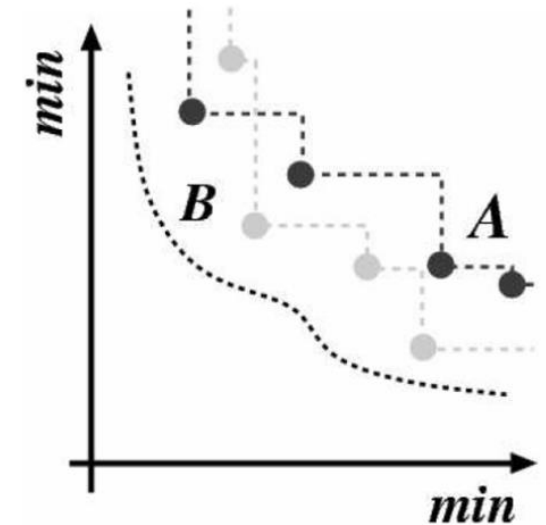
S-Metric

- 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.
- Cons:
 - Requires defining some upper boundary of the region.
 - It has a large computational overhead, especially for high-dimensional cases.

Reading: Nicola Beume, Carlos M. Fonseca, Manuel López-Ibáñez, Luís Paquete, and J. Vahrenhold. [On the complexity of computing the hypervolume indicator](#). *IEEE Transactions on Evolutionary Computation*, 13(5):1075–1082, 2009.

C-Metric

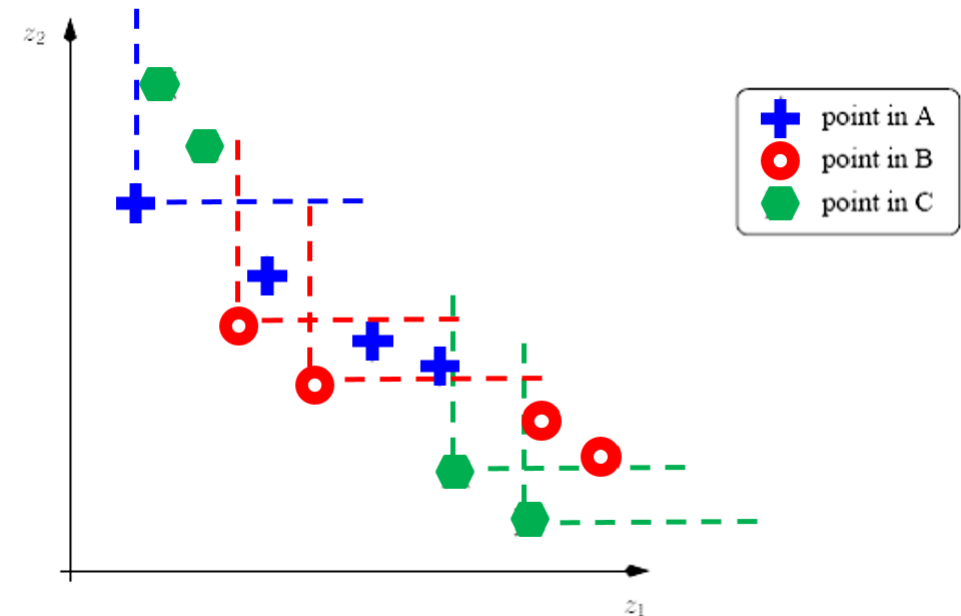
- Coverage of two sets $C(X,Y)$: given two sets of non-dominated solutions X and Y found by the compared algorithms, the measure $C(X,Y)$ returns a ratio of a number of solutions of Y that are dominated by or equal to any solution of X to the whole set Y .
 - It returns values from the interval $[0,1]$.
 - The value $C(X,Y) = 1$ means that all solutions in Y are covered by solutions of the set X .
 - And vice versa, the value $C(X,Y) = 0$ means that none of the solutions in Y are covered by the set X .
 - Always both orderings have to be considered, since $C(X,Y)$ is not necessarily equal to $1-C(Y,X)$.
 - Roughly speaking, $C(X,Y) > C(Y,X)$ indicates that X is better than Y .



$$C(A, B) = 0.25, C(B, A) = 0.75$$

C-Metric

- It has low computational overhead.
- If two sets are of different cardinality and/or the distributions of the sets are non-uniform, then it gives unreliable results.
- It is cycle-inducing: if three sets are compared using C-metric, they may not be ordered.
- Example:
 - $C(A,B) = 0$, $C(B,A) = 3/4$
 - $C(A,C) = 1/2$, $C(C,A) = 0$
 - $C(B,C) = 0$, $C(C,B) = 1/2$
 - B is considered better than A
 - A is considered better than C
 - C is considered better than B



D-Metric

- D-metric measures the **mean distance** over the points in a reference set, of the nearest point in an approximation set.

$$D(A, P^*) = \frac{\sum_{v \in P^*} d(v, A)}{|P^*|}$$

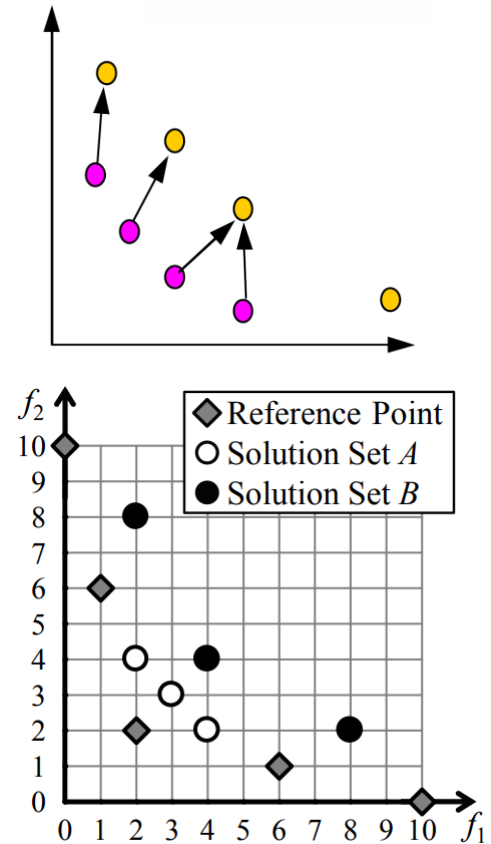
where A is the approximation set, P^* is a reference set (true PF), $d(v, A)$ denotes the **nearest distance** from v to solutions in A .

- Pros:

- It is cheap to compute.
- It can differentiate between different levels of complete outperformance given an appropriate choice of reference set.

- Cons:

- It is not Pareto compliant.
- The score is strongly dependent upon the distribution of points in the reference set.



Example with misleading IGD.

Applications

- Popular benchmark problems
 - Continuous functions
 - Knapsack
 - Traveling salesman problem
 - Permutation flow shop scheduling

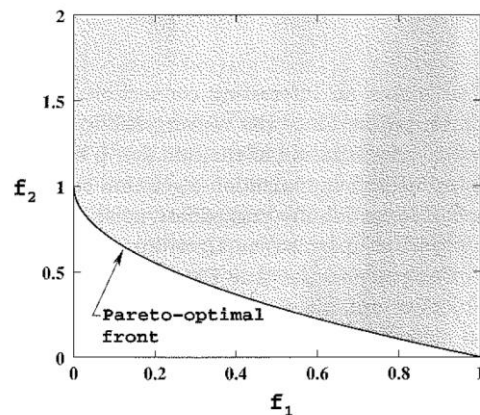


Figure 213 The search space near the Pareto-optimal region for ZDT1.

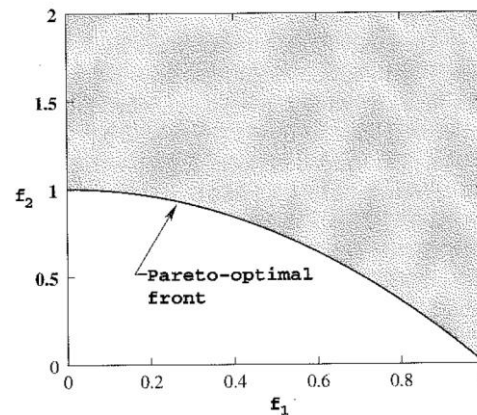


Figure 214 The search space near the Pareto-optimal region for ZDT2.

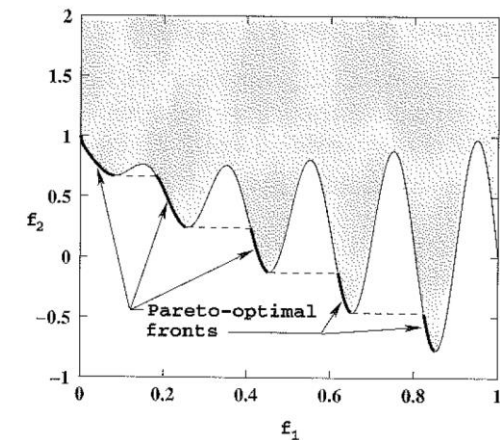


Figure 215 The search space near the Pareto-optimal region for ZDT3.

Thank you!

