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odified L-SHADE for Single Objective Real-Parameter Optimization



Contributed by
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1 Introduction

- The Problem introduction

2 SHADE

- Algorithms overview
- Adaptive Parameter
- Linear population size reduction

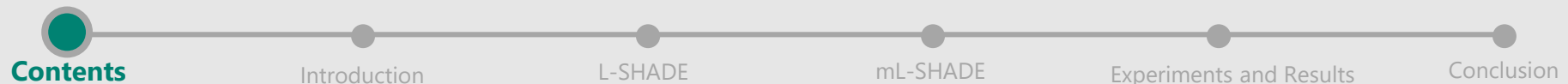
3 Proposed algorithm—mL-SHADE

- Proposed mL-SHADE mechanism

4 Experiments and Results

- Performance compression
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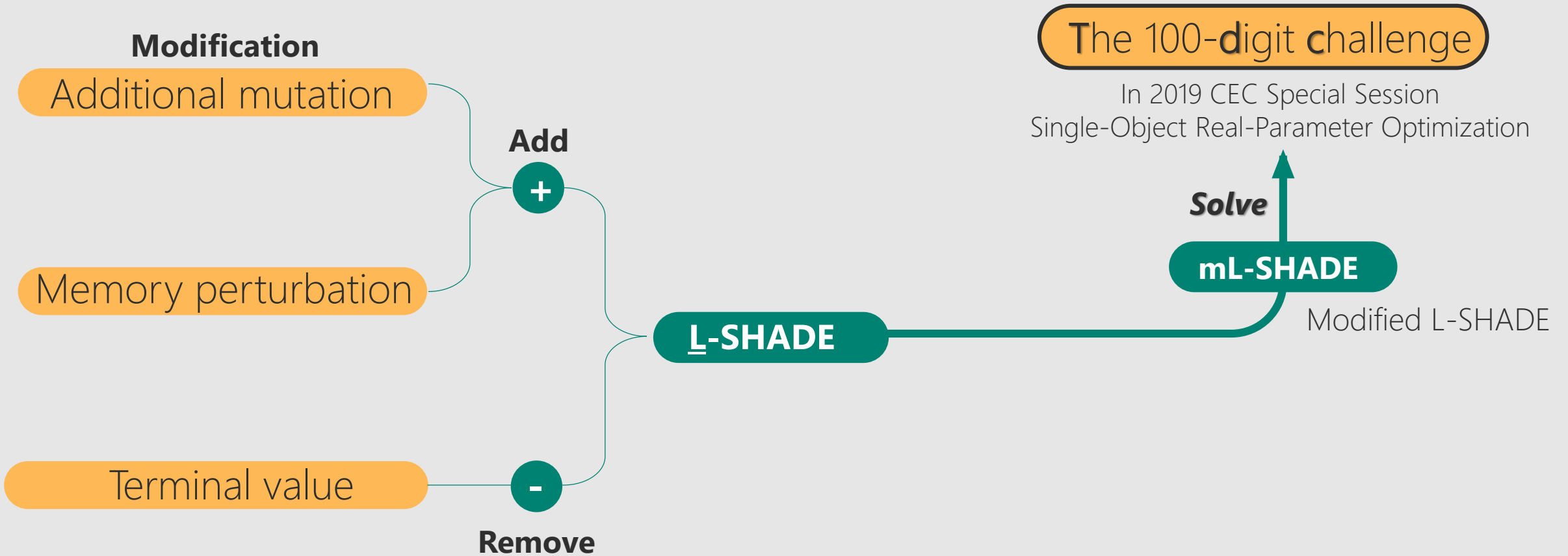


• CEC 2019 Competitions

- CEC-C02 Competition on "Evolutionary Multi-task Optimization"
- CEC-C03 Competition on "Online Data-Driven Multi-Objective Optimization Competition"
- CEC-C04 Competition on "Smart Grid and Sustainable Energy Systems"
- CEC-C05 Competition on "Evolutionary Computation in Uncertain Environments: A Smart Grid Application"

• CEC-C06 Competition on "100-Digit Challenge on Single Objective Numerical Optimization"

- CEC-C07 FML-based Machine Learning Competition for Human and Smart Machine Co-Learning on Game of Go
- CEC-C08 General Video Game AI Single-Player Learning Competition
- CEC-C09 Strategy Card Game AI Competition
- CEC-C10 Nonlinear Equation Systems Competition

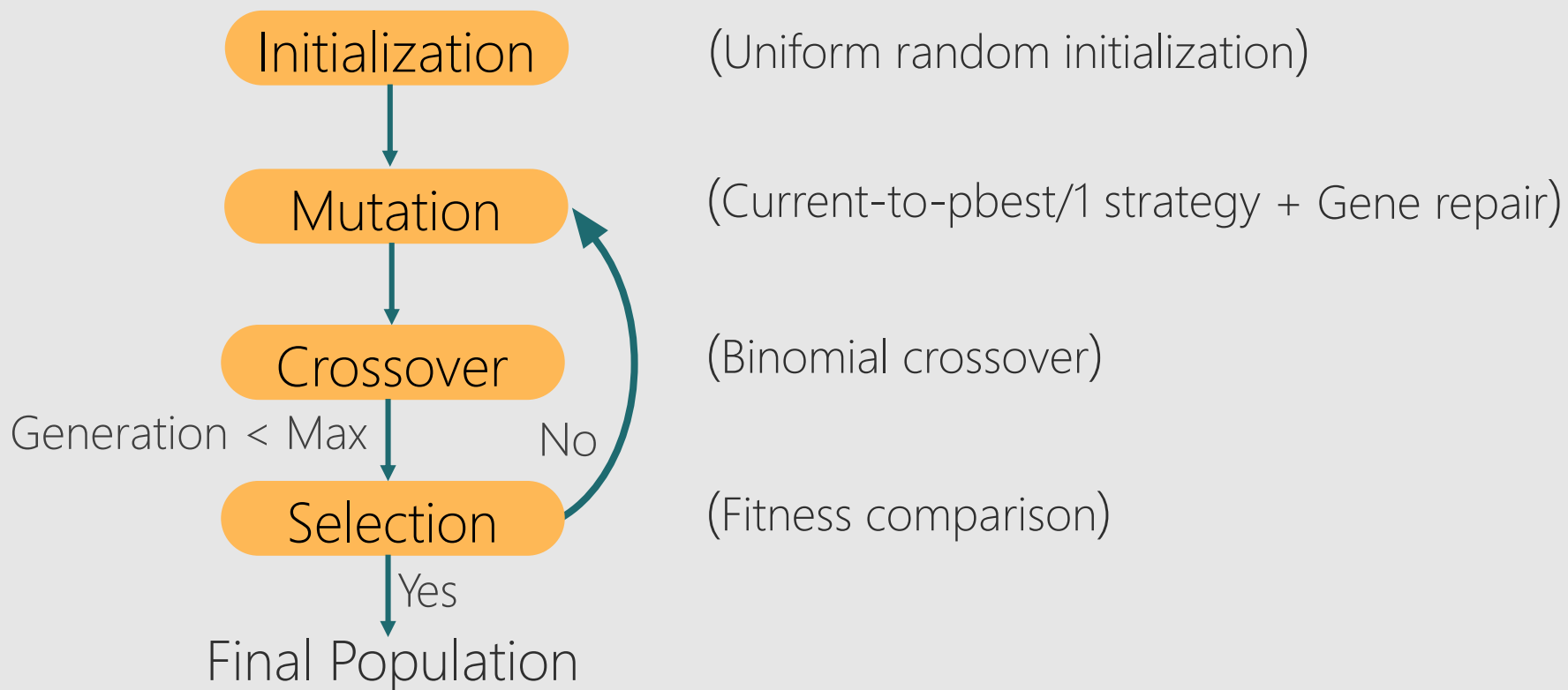


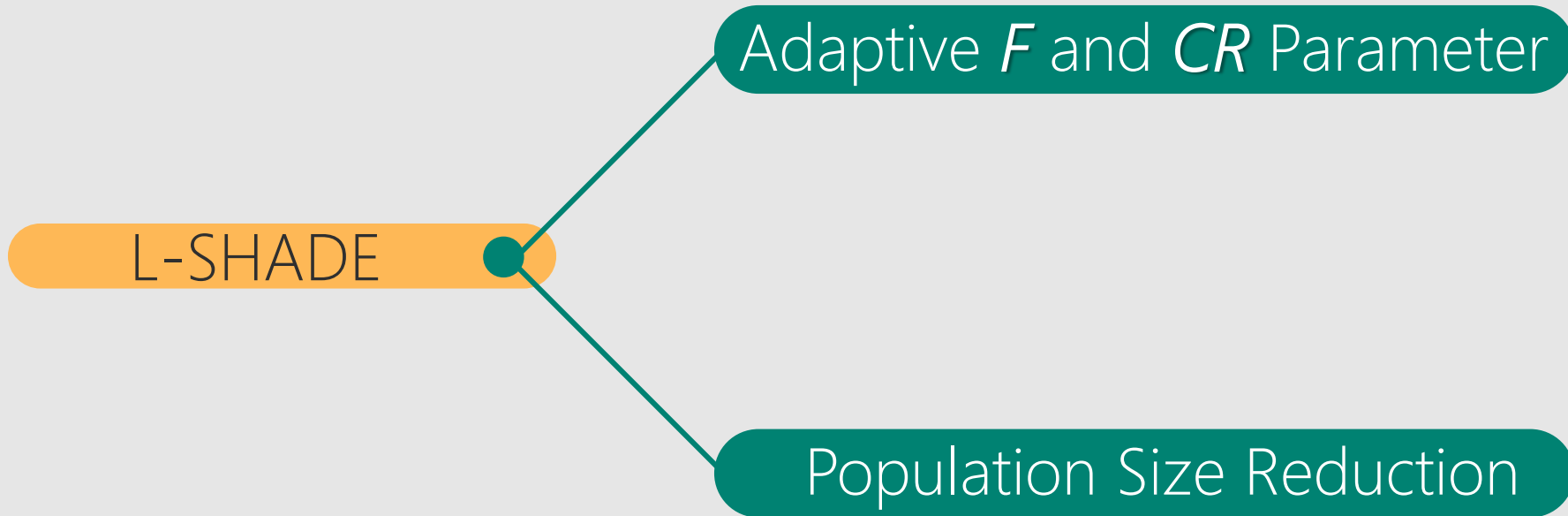
[L-SHADE] R. Tanabe, and A. S. Fukunaga, "Improving the Search Performance of SHADE Using Linear Population Size Reduction," in IEEE CEC, pp. 1658–1665, 2014.



- Algorithms overview

L-SHADE / mL-SHADE







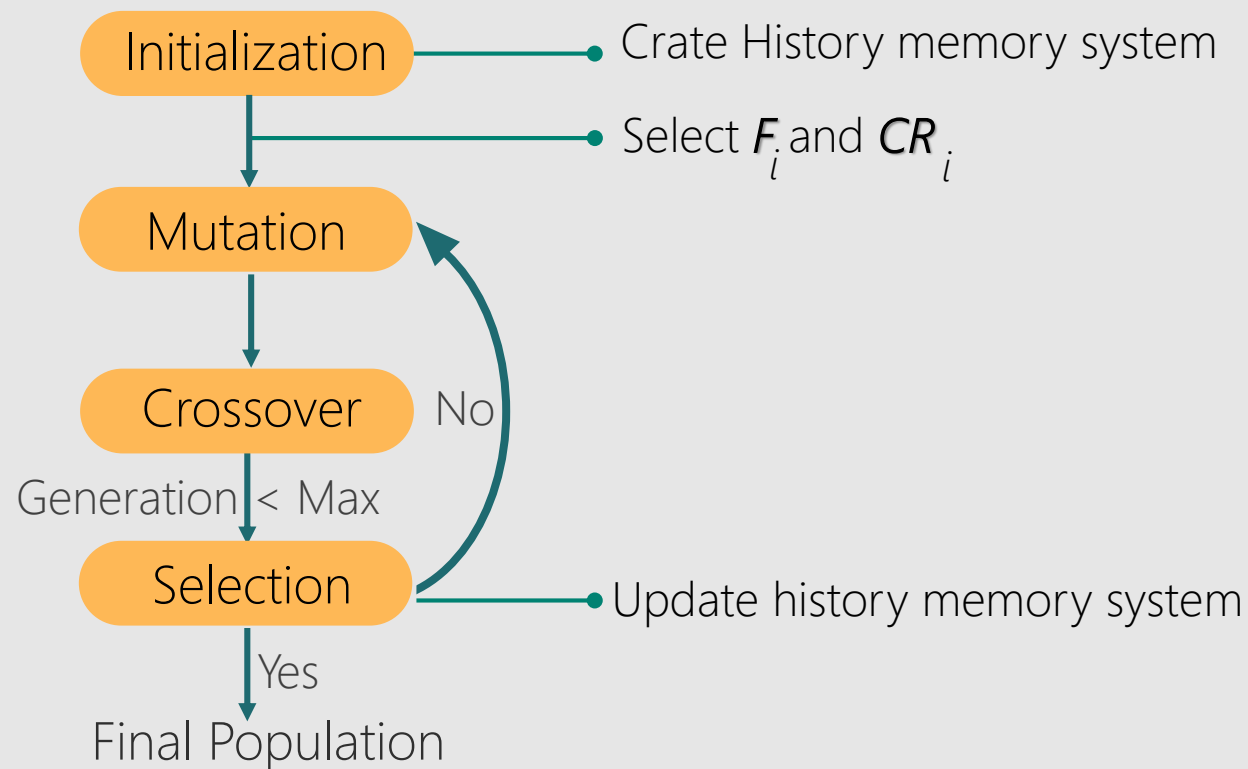
Adaptive Parameter

History memory system

| | | | | | |
|----------|------------|------------|-------|--------------|------------|
| M_F | $M_{F,1}$ | $M_{F,2}$ | | $M_{F,H-1}$ | $M_{F,H}$ |
| M_{CR} | $M_{CR,1}$ | $M_{CR,2}$ | | $M_{CR,H-1}$ | $M_{CR,H}$ |

- M is mean value of the successful F and CR which can generate the better solution in each iteration
- Each mean value will be utilized to generate F and CR next iteration

L-SHADE





- Adaptive Parameter

H

| | | | | | |
|----------|-----|-----|-------|-----|-----|
| M_F | 0.5 | 0.5 | | 0.5 | 0.5 |
| M_{CR} | 0.5 | 0.5 | | 0.5 | 0.5 |

In the initialization stage, all element of history table will be set to be 0.5, and each of them will be updated when the set of better solution is found

Initialization

Mutation

Crossover

Selection



Adaptive Parameter

| | | | | | |
|----------|------------|------------|-------|--------------|------------|
| | H | | | | |
| M_F | $M_{F,1}$ | $M_{F,2}$ | | $M_{F,H-1}$ | $M_{F,H}$ |
| M_{CR} | $M_{CR,1}$ | $M_{CR,2}$ | | $M_{CR,H-1}$ | $M_{CR,H}$ |

For each target vector x_i will generated F_i and CR_i as follow.

r_i index is selected randomly from $[1,H]$

$$F_i = \text{randc}_i(M_{F,r_i}, 0.1)$$

$\text{randc}_i()$ is a Cauchy distribution

$$CR_i = \begin{cases} \text{randn}_i(M_{CR,r_i}, 0.1) & \text{if } M_{CR,r_i} \neq 0(\perp) \\ 0 & \text{otherwise} \end{cases}$$

$\text{randn}_i()$ is a normal distribution

** 0 (\perp terminal value)

Initialization

Mutation

Crossover

Selection



Adaptive Parameter

At the selection, if the trial vector's (\vec{u}) fitness is better than or equal to target vector's (\vec{x}), their fitness value, F_i , and CR_i will be stored in S table.

$$\Delta f_k = |f(u_{i,G}) - f(x_{i,G})|$$

Fitness improvement

1

$$w_k = \frac{\Delta f_k}{\sum_{l=1}^{|S|} \Delta f_l}$$

Improvement-based weight

2

3

$$mean_{wL}(F) = \frac{\sum_{k=1}^{|S|} w_k \cdot F_k^2}{\sum_{k=1}^{|S|} w_k \cdot F_k}$$

weighted Lehmer mean

$$mean_{wL}(CR) = \frac{\sum_{k=1}^{|S|} w_k \cdot CR_k^2}{\sum_{k=1}^{|S|} w_k \cdot CR_k}$$

CR

Initialization

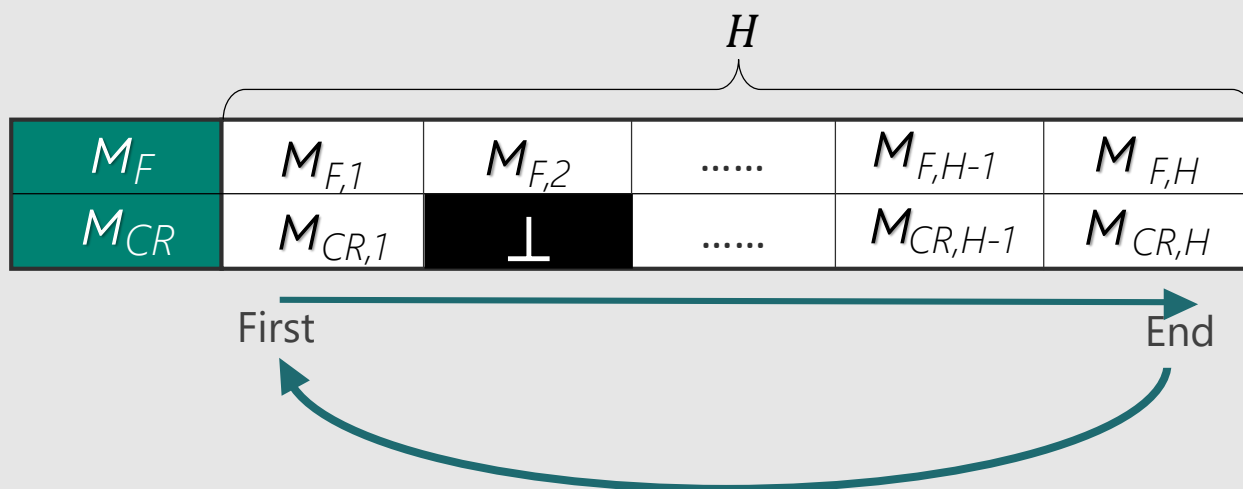
Mutation

Crossover

Selection



Adaptive Parameter



Initialization

Mutation

Crossover

Selection

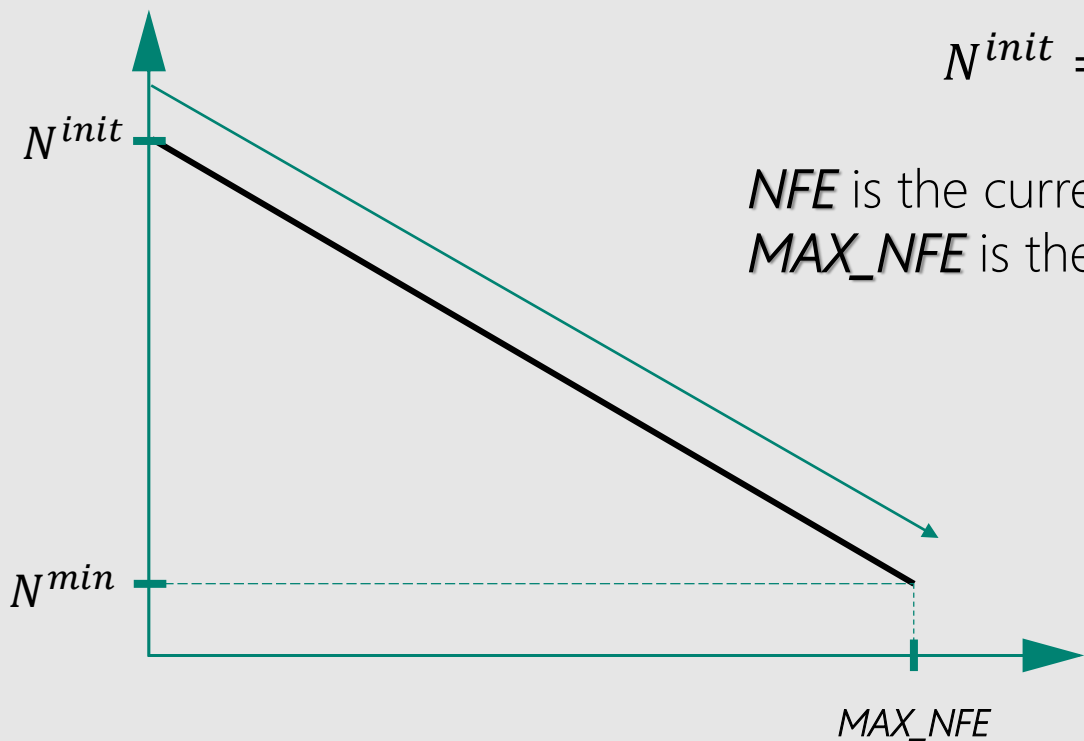
1. If S table is not empty, mean value of $M_{F,k}$ and $M_{CR,k}$ will be updated by new mean F and CR
2. If mean value of CR is 0, then $M_{CR,k}$ will be set as the terminal value \perp (0) and that element will never be changed to be the other number again.



- Linear population size reduction

$$NP_{G+1} = \text{round}\left(\left(\frac{N^{min} - N^{init}}{MAX_NFE}\right) \times NFE + N^{init}\right)$$
$$N^{init} = 18 \times D, N^{min} = 4$$

NFE is the current number of fitness evaluations
MAX_NFE is the maximum number of fitness evaluations



Initialization

Mutation

Crossover

Selection



L-SHADE



➔ mL-SHADE

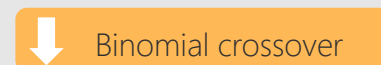
- Terminal value
- + Memory perturbation
- + Additional mutation operator



- Remove Terminal value

As L-SHADE will update the $M_{CR,k}$ element inside history memory table to be \perp every time, when it found the mean of CR equal 0 and never change to be the other value again. It also forces the target vector to CR as 0, it select the \perp from history table.

This can end the exploration and start to exploitation.



we found that in some cases, all M_{CR} element are set to terminal value when the evolution phase is *very early*.

***This may affect the performance of the algorithm, so we remove the terminal value in mL-SHADE algorithm.*

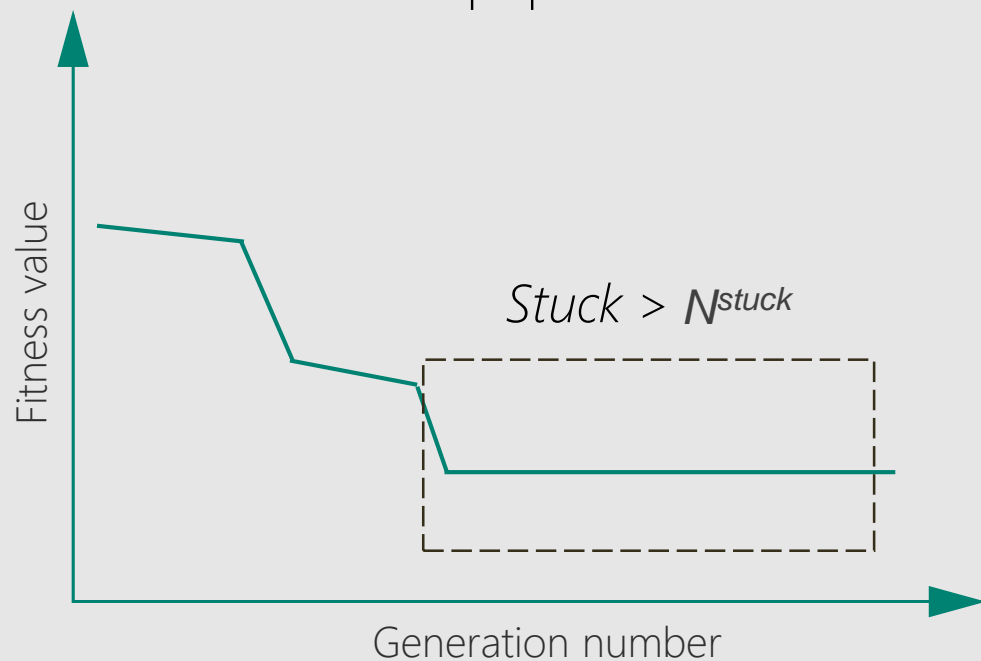




- Memory perturbation

We found that the memory may **not be updated** for a long time, which means that the fitness value has not improved.

One of the reasons why fitness value stops improving is that the control parameters are not suitable for the current population.



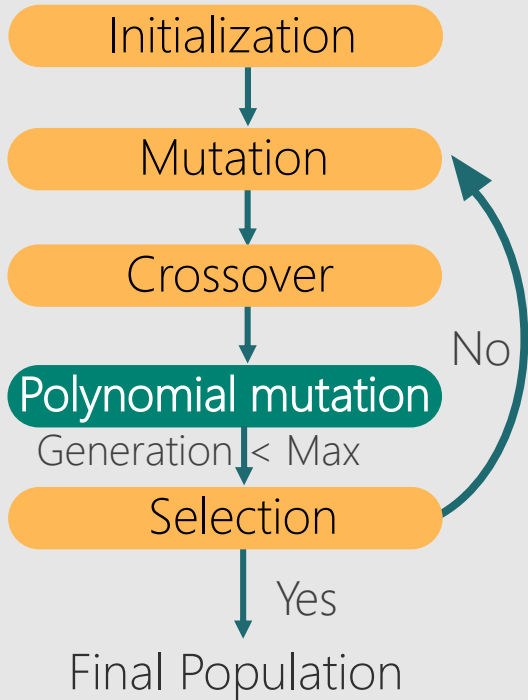
$$M_{CR,k} = 1.0 - M_{CR,k}$$

$$M_{F,k} = 1.0 - M_{F,k}$$

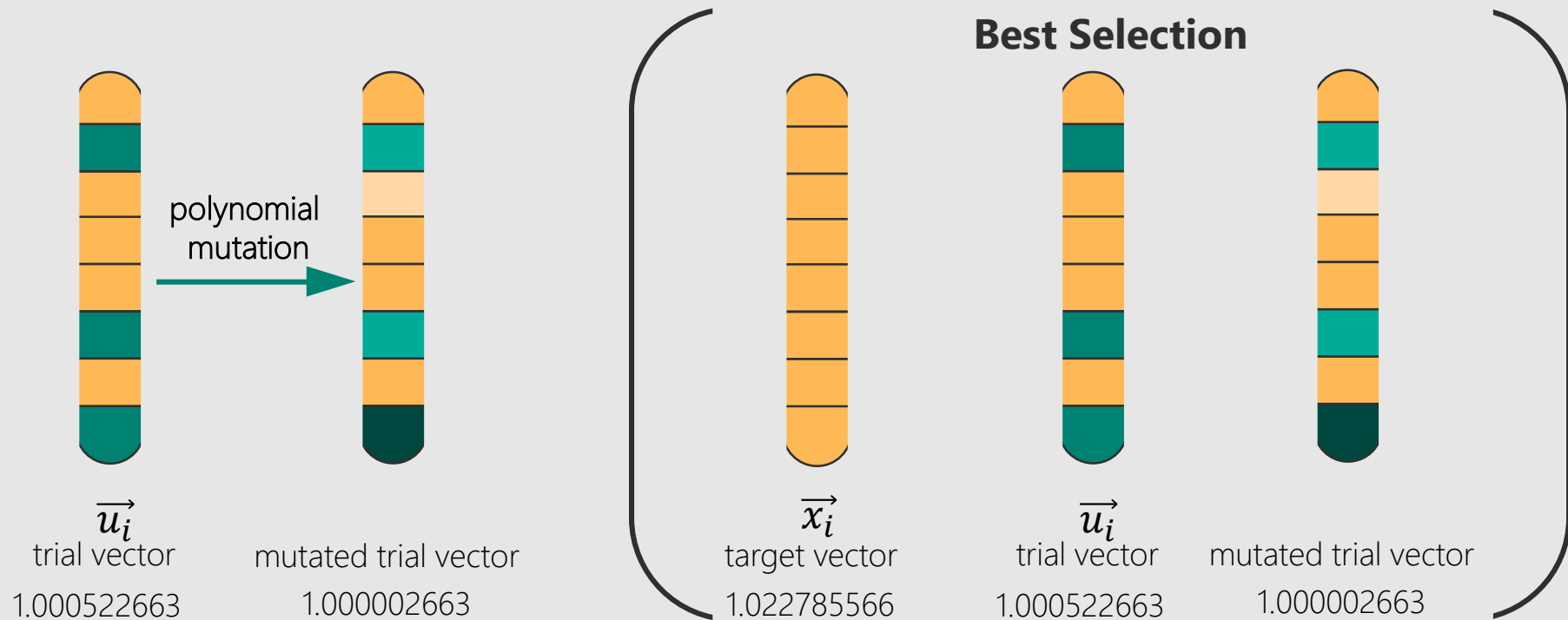


- Additional **m**utation operation (polynomial mutation)

mL-SHADE



After the trial vector is generated, **polynomial mutation (PM)** is applied to generate a mutated trial vector, and choose the better one to be the final trial vector.





- 100-Digit Challenge on Single Objective Numerical Optimization (CEC C06) was utilized to test the performance of our algorithm
- We compared mL-SHADE with the other seven algorithms including L-SHADE
- The source code of those algorithms can be downloaded from organizer's website.



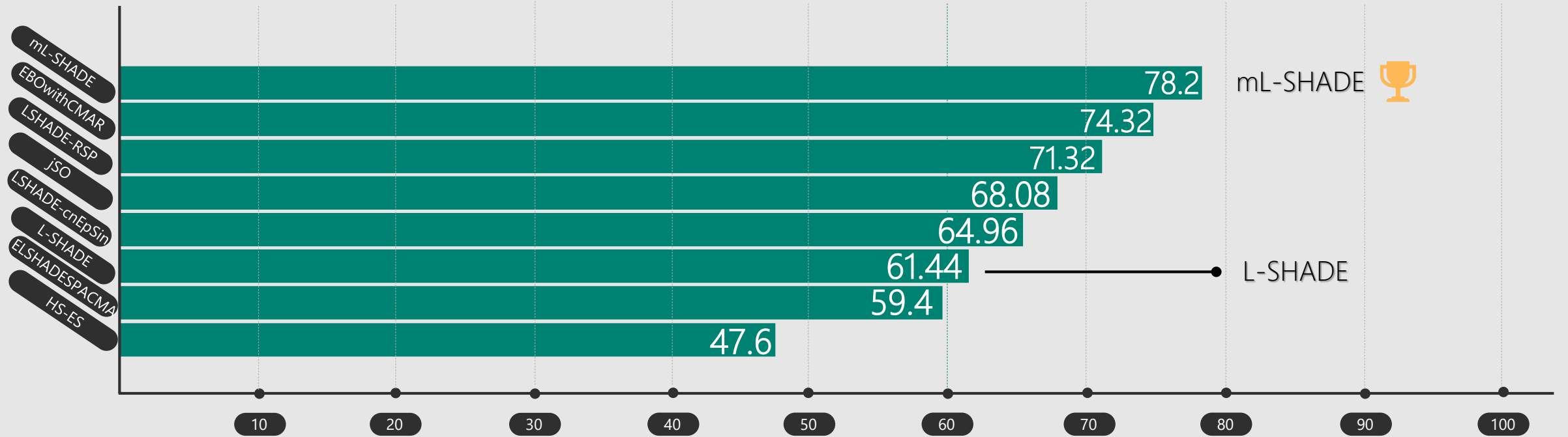
• Parameter setting

| Parameter | Meaning | mL-SHADE | L-SHADE |
|-------------|---|----------------|-----------------|
| N^{init} | size of the initial population | $18 \cdot D$ | $18 \cdot D$ |
| N_{min} | minimal population size | 4 | 4 |
| H | size of the history memory | 6 | 6 |
| r^{arc} | archive size $ A = \text{round}(r^{arc} \cdot N^{init})$ | 1.0 | 2.6 |
| p | required in the cur-to- p best/1 mutation | 0.11 | 0.11 |
| m_r | probability of polynomial mutation | 0.05 | N/A |
| p_m, η | parameters of polynomial mutation | $1/D, 10$ | N/A |
| $MaxNFE$ | maximum number of fitness evaluations | $2 \cdot 10^6$ | $10000 \cdot D$ |

| No. | N^{stuck} | No. | N^{stuck} |
|-----|------------------|-----|------------------|
| 1 | 400 | 6 | 400 |
| 2 | 400 | 7 | 400 |
| 3 | 6 (same as H) | 8 | 400 |
| 4 | 400 | 9 | 6 (same as H) |
| 5 | 400 | 10 | 400 |



Results





Results

Ranking:

1. mL-SHADE 
2. EBOwithCMAR
3. L-SHADE-RSP
4. jSO
5. L-SHADE-cnEpSin
6. L-SHADE
7. ELSHADESPACMA
8. HS-ES

| No. | mL-SHADE | L-SHADE | ELSHADESPACMA | jSO |
|-------|------------|----------------|---------------|-------|
| 1 | 10 | 10 | 10 | 10 |
| 2 | 10 | 10 | 10 | 10 |
| 3 | 10 | 7.16 | 5.44 | 10 |
| 4 | 10 | 0.24 | 0.84 | 3.88 |
| 5 | 10 | 10 | 10 | 10 |
| 6 | 10 | 10 | 10 | 10 |
| 7 | 5.04 | 1 | 1.08 | 1 |
| 8 | 1 | 1 | 1.16 | 1.08 |
| 9 | 2.16 | 2.04 | 3 | 2.12 |
| 10 | 10 | 10 | 7.88 | 10 |
| Score | 78.2 | 61.44 | 59.4 | 68.08 |
| No. | LSHADE-RSP | LSHADE-cnEpSin | EBOwithCMAR | HS-ES |
| 1 | 10 | 10 | 10 | 7.6 |
| 2 | 10 | 10 | 10 | 0 |
| 3 | 10 | 6.12 | 10 | 1.72 |
| 4 | 6.76 | 4.6 | 10 | 4.96 |
| 5 | 10 | 10 | 10 | 10 |
| 6 | 10 | 10 | 10 | 10 |
| 7 | 1.36 | 0.84 | 0.56 | 0.16 |
| 8 | 1 | 1.04 | 1.68 | 0.16 |
| 9 | 2.2 | 2.36 | 2.08 | 3 |
| 10 | 10 | 10 | 10 | 10 |
| Score | 71.32 | 64.96 | 74.32 | 47.6 |



• Results

- We made a minor changes after submitting the paper, which let us get a higher score.

The repair method of CR value is modified.

- Rule 1: If CR is bigger than 1, CR is set as 1.
- Rule 2: If CR is negative, it will be set as its absolute value, then apply the Rule 1.

| No. | mL-SHADE |
|-------|----------|
| 1 | 10 |
| 2 | 10 |
| 3 | 10 |
| 4 | 10 |
| 5 | 10 |
| 6 | 10 |
| 7 | 8.12 |
| 8 | 1.08 |
| 9 | 2.12 |
| 10 | 10 |
| Score | 81.32 |



- In our experiments, we found three problems (7-9) that are difficult to all tested algorithms. **No single algorithm** can get the highest score in more than one of them.
- For the future work, we will continue our research to study how different algorithms fit different functions and then to propose a **better integration**.
- Another direction is to develop an adaptive method to adjust the value of the important parameter N^{stuck} in our memory perturbation mechanism.

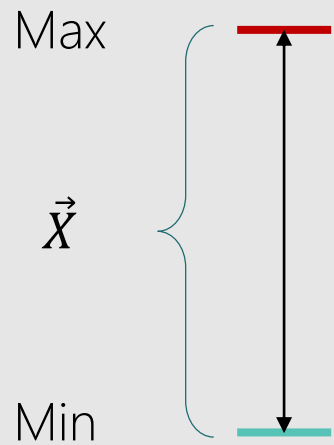
Thanks for your attention



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• Uniform random initialization



- Basically, the parameter of each initial vector is be selected randomly by using the below equation.
- The variable $x_{j,min}$ and $x_{j,max}$ are the minimum and maximum value of each parameter. $rand_{ij}$ is the random real value between 0 to 1.

$$x_{j,i,G} = x_{j,min} + rand_{i,j}[0,1] * (x_{j,max} - x_{j,min})$$





- Current-to-pbest/1 strategy

$$v_{i,g} = x_{i,g} + F_i \cdot (x_{pbest,g} - x_{i,g}) + F_i \cdot (x_{r1,g} - x_{r2,g})$$

Where $v_{i,g}$ is the mutated vector of $x_{i,g}$

$x_{pbest,g}$ is randomly selected from top 100p% solutions in the current population.

$x_{r1,g}$ is randomly selected from current population.

$x_{r2,g}$ is randomly selected from the combination of current population and the set of reprints which are replaced by trial vector.



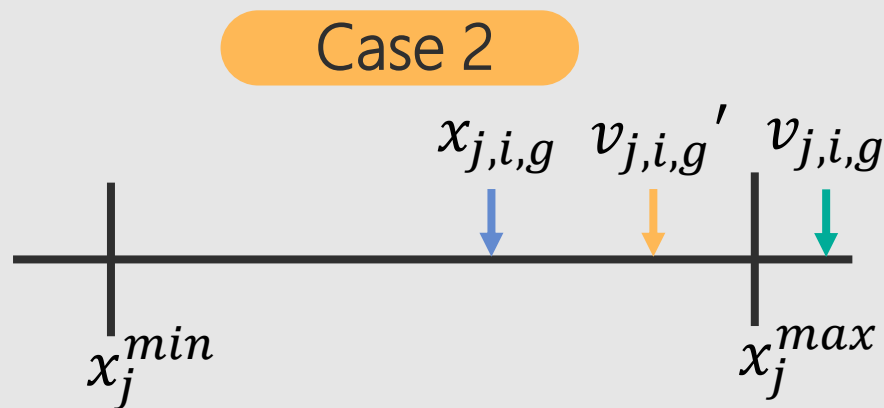
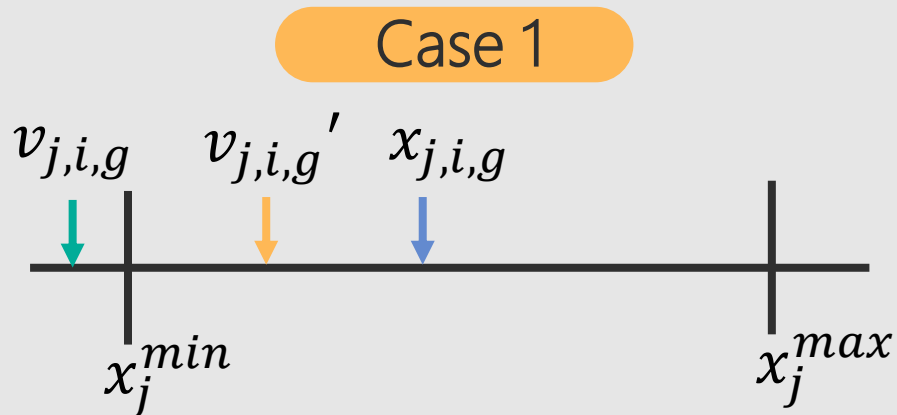


• Gene repair

$$v_{j,i,g}' = \begin{cases} (x_j^{\min} + x_{j,i,g})/2 & \text{if } v_{j,i,g} < x_j^{\min} \\ (x_j^{\max} + x_{j,i,g})/2 & \text{if } v_{j,i,g} > x_j^{\max} \end{cases}$$

Case 1

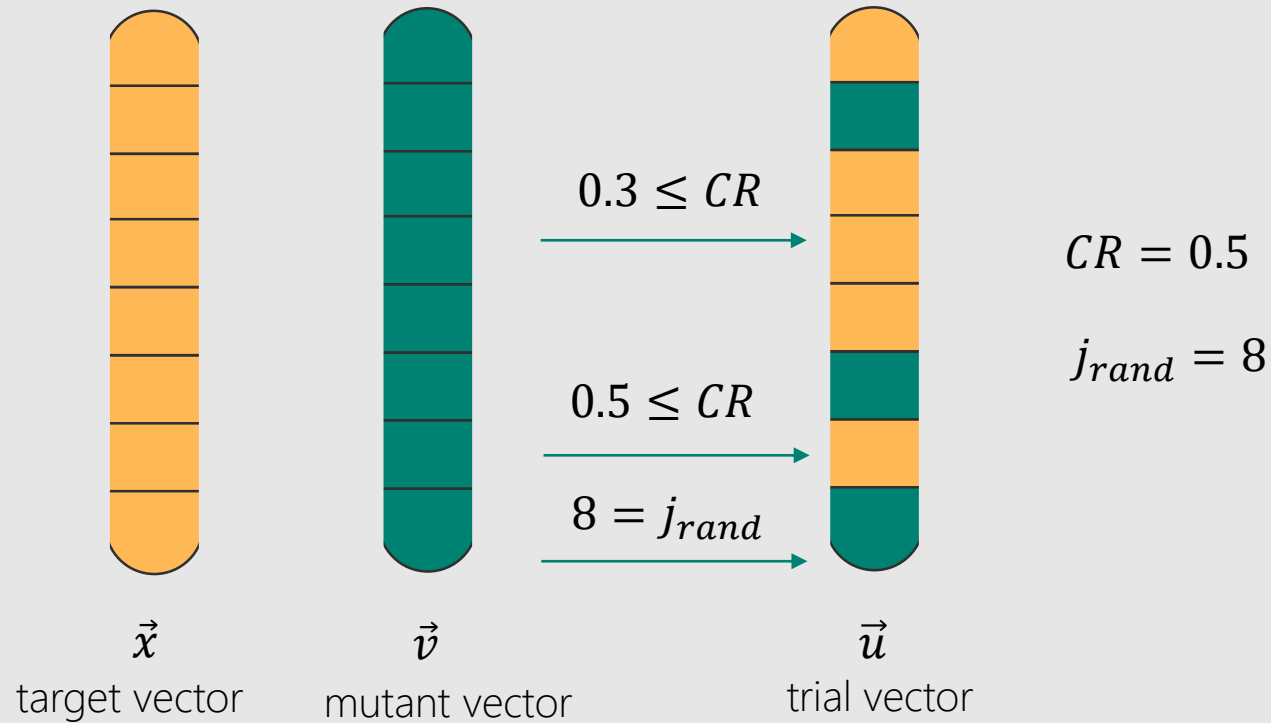
Case 2





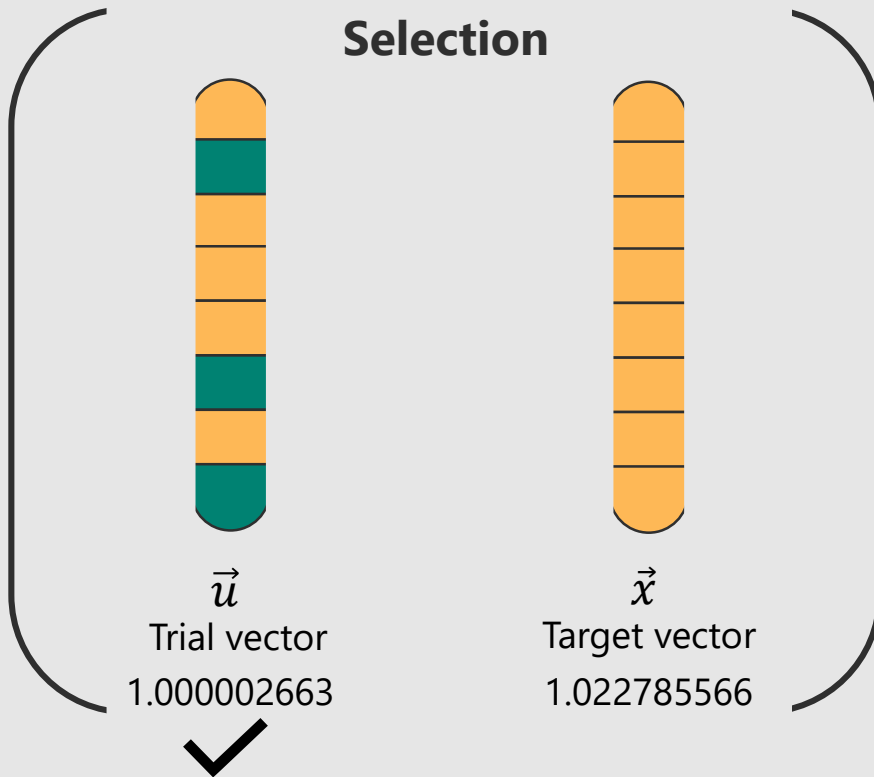
Binomial crossover

$$u_{j,i,g} = \begin{cases} v_{j,i,g} & \text{if } rand[0,1) \leq CR_i \text{ or } j = j_{rand} \\ x_{j,i,g} & \text{otherwise} \end{cases}$$





• selection



The trial vector $\vec{u}_{i,g}$ will be selected to be the candidate solution in the iteration if its fitness is not better than the target vector $\vec{x}_{i,g}$.

$$x_{i,g+1} = \begin{cases} u_{i,g} & \text{if } f(u_{i,g}) \leq f(x_{i,g}) \\ x_{i,g} & \text{otherwise} \end{cases}$$

