

Evolution Strategies $(\mu/\varrho + \lambda)$ -ES for test functions optimization

1 Evolution Strategies in version (1+1)-ES

```
1: population  $\leftarrow$  initialize ( $\mu$ )
2:  $g \leftarrow 0$ 
3: while terminal_condition not TRUE do
4:   offspring  $\leftarrow$  prepare ( $\lambda$ )
5:   for  $o = 1$  to  $\lambda$  do
6:     parents  $\leftarrow$  marriage (population,  $\varrho$ )
7:      $s \leftarrow s\_recombination$  (parents)
8:      $x \leftarrow x\_recombination$  (parents)
9:      $s \leftarrow s\_mutation$  ( $s$ )
10:     $x \leftarrow x\_mutation$  ( $x$ ,  $s$ )
11:     $f \leftarrow fitness\_function$  ( $x$ )
12:    offspring $o$   $\leftarrow$  struct ( $x$ ,  $s$ ,  $f$ )
13:  end for
14:  population  $\leftarrow$  selection (offspring +  $\mu$ )
15:   $g \leftarrow g + 1$ 
16: end while
17: return population
```

2 Algorithm description

- *initialize*(μ) returns the μ individuals, which could be a structure/object:
 - x — vector with d numbers which represents optimization function arguments - point in the domain space. \mathbb{R}^n). Initialized randomly from uniform distribution and the range = *domain*.
 - s — vector with d numbers which represents standard deviation to control the mutation strength in every dimension. Initialized randomly from uniform distribution and the range = $[0.1 : 10]$.
 - function — fitness function (calculated from the formula e.g. Schwefel function)

domain is a given domain for a specific test function (benchmark) that will be solved by ES.

- Initial parameters
 - μ — number of individuals in the population: recommended values from 1 to 100
 - ϱ — number of individuals in the pool of parents for reproduction: recommended values from 1 to μ
 - λ — the number of offsprings: recommended values from 1 to 100 .
 - max_num_gen - maxim number of generations e.g. 10000
- A *teminal_condition* control maximum number of steps or finding the solution close to the know global optimum e.g. $fitness_function(best) - f_global < 0,0001$.

- Selection of the pool of parents

```

function parents = marriage (population,  $\varrho$ )
1: parents  $\leftarrow$  prepare ( $\varrho$ )
2: selected_individuals  $\leftarrow$  randperm (length (population))
3: for  $i = 1$  to  $\varrho$  do
4:   parents $i$   $\leftarrow$  populationselected_individuals $i$ 
5: end for
6: return parents

```

- Pseudo-code of the recombination algorithm for endogenous parameters

```

function s = s_recombination (parents)
1:  $n \leftarrow$  length (parents.s)
2: s  $\leftarrow$  prepare ( $n$ )
3: for  $i = 1$  to  $n$  do
4:   s $i$   $\leftarrow$  mean_s (parents,  $i$ )
5: end for
6: return s

```

where: $mean_s (parents, i)$ is the average value of all the parents.

- Pseudo-code of recombination

```

function x = x_recombination (parents)
1:  $n \leftarrow$  length (parents)
2: x  $\leftarrow$  prepare ( $n$ )
3: for  $i = 1$  to  $n$  do
4:   x $i$   $\leftarrow$  mean_x (parents,  $i$ )
5: end for
6: return x

```

where, $mean_x (parents, i)$ is the average value of all the parents.

- Endogenous parameters mutation

```

function out_s = s_mutation (s)
1:  $n \leftarrow$  length (s)
2: out_s  $\leftarrow$  prepare ( $n$ )

```

```

3:  $c \leftarrow 1$ 
4:  $\tau_0 \leftarrow e^{\frac{c}{\sqrt{2n}} \cdot \aleph(0,1)}$ 
5:  $\tau \leftarrow \frac{c}{\sqrt{2\sqrt{n}}}$ 
6: for  $i = 1$  to  $n$  do
7:    $out\_s_i \leftarrow \tau_0 \cdot s_i \cdot e^{\tau \cdot \aleph(0,1)}$ 
8: end for
9: return  $out\_s$ 

```

where:

- $\aleph(0, 1)$ is a random number from a normal distribution with the mean equal to 0 and standard deviation of 1
- variable c is positive number could be set to 1.

- Problem parameters mutation

function $out_x = x_mutation(x, s)$

```

1:  $n \leftarrow length(y)$ 
2:  $out\_x \leftarrow prepare(n)$ 
3: for  $i = 1$  to  $n$  do
4:    $out\_x_i \leftarrow x_i + s_i \cdot \aleph(0, 1)$ 
5: end for
6: return  $out\_x$ 

```

where, $\aleph(0, 1)$ is a random number from a normal distribution with the mean equal to 0 and standard deviation of 1

- Selection called plus (+)

function $new_population = selection(offspring, population, \mu)$

```

1:  $new\_population \leftarrow sort([offspring, population], ascd)$ 
2:  $new\_population \leftarrow new\_population(1 : \mu)$ 
3: return  $new\_population$ 

```

3 Benchmarks

To test the algorithm the following function should be used:

1. RASTRIGIN FUNCTION

- <http://www.sfu.ca/~ssurjano/rastr.html>,
- $domain = [-5.12, 5.12]$
- Global minimum $f(0, 0, \dots, 0) = 0$

2. GRIEWANK FUNCTION

- <http://www.sfu.ca/~ssurjano/griewank.html>,
- $domain = [-600, 600]$
- Global minimum $f(0, 0, \dots, 0) = 0$

3. SPHERE FUNCTION

- <http://www.sfu.ca/~ssurjano/spheref.html>,
- $domain = [-5.12, 5.12]$
- Global minimum $f(0, 0, \dots, 0) = 0$

4. ZAKHAROV FUNCTION

- <http://www.sfu.ca/~ssurjano/zakharov.html>,
- $domain = [-5, 10]$
- Global minimum $f(0, 0, \dots, 0) = 0$

5. EASOM FUNCTION

- <http://www.sfu.ca/~ssurjano/easom.html>,
- $domain = [-100, 100]$
- Global minimum $f(\pi, \pi) = -1$

6. STYBLINSKI-TANG FUNCTION

- <http://www.sfu.ca/~ssurjano/stybtang.html>,
- $domain = [-5, 5]$
- Global minimum $f(-2.903534, \dots, -2.903534) = -39.16599d$, where d is dimension