# CSE/ECE 848 Introduction to Evolutionary Computation

Module 3 - Lecture 10 - Part 2
Evolutionary Strategies Mutation and Recombination

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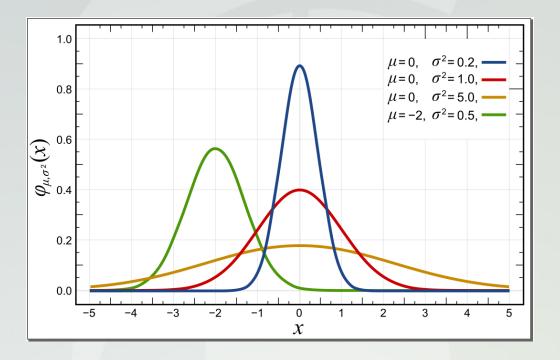
## Mutation

- As mentioned previously, mutation is the driving force
- x is the individual parameter value

$$x_i' = x_i + \sigma * N(0,1)$$

 where σ is the stepsize, and N(0,1) represents a single standard Gaussian random variable







- Early ES were concerned with the concept of stepsize σ
- "Stepsize" meant the strength of mutations
- However, the rate of change of each element of vector x should change over time as the element "converges" to better and better answers
- The issue of stepsize is the same as for any gradient hillclimber: If the step size is too small, little progress is made in exploring the space. If the stepsize is too big, the answer is missed because of overstepping in the space



## **Heuristic Mutation Strategy**

Rechenberg used two common functions to estimate optimal step sizes:

Linear corridor

$$f_1(x) = F(x_1) = c_0 + c_1 x_1$$
  
 $\forall_i \in \{2, ..., n\} : -b/2 \le x_i \le b/2$ 

The sphere

$$f_2(x) = c_0 + c_1 * \sum_{i=1}^{n} (x_i - x_i^*)^2$$



## The 1/5 Rule

 He solved the optimal expected convergence rates stepsizes for those two (which were respectively)

$$p_{opt} \approx 0.184 \ p_{opt} \approx 0.270$$

 and decided that the best rate of successful mutations to failed mutations should be about 0.20, or one fifth.



### This led to the basic mutation rule:

- If more than 1/5th of the mutations cause an improvement (in the objective function) then multiply σ by a factor 1+s
- If less than 1/5th of the mutations cause an improvement, then multiply σ by (1+s)^(-1/4)

Algorithm: (1+1)-ES with 1/5 success-rule					
1. Initialize $\boldsymbol{X}_0,\sigma_0$					
2. repeat					
3. $\widetilde{\boldsymbol{X}}_n = \boldsymbol{X}_n + \sigma_n \mathcal{N}(\boldsymbol{0}, \boldsymbol{I})$	Sample one offspring				
4. if $f(\widetilde{\boldsymbol{X}}_n) \leq f(\boldsymbol{X}_n)$ then	If $f(\text{offsp.}) \leq f(\text{parent})$				
5. $X_{n+1} = \widetilde{X}_n$	New parent $=$ offsp.				
$6. \qquad \sigma_{n+1} = 1.5  \sigma_n$	Step-size is increased				
7. else	If offspring strictly worse				
8. $\boldsymbol{X}_{n+1} = \boldsymbol{X}_n$	$New\ parent = old\ parent$				
9. $\sigma_{n+1} = 1.5^{-1/4}  \sigma_n$	Step-size is decreased				
10. until stopping criteria is met					

After one successful and 4 unsuccessful mutations, this results in:

$$E(\sigma_{n+1}|\sigma_n) = \left((1.5)^{-1/4}\right)^{4/5} (1.5)^{1/5} \sigma_n$$

$$E(\sigma_{n+1}|\sigma_n) = \sigma_n$$

## First ES

- The first ES was based on a (1+1)-ES strategy, where the child competes with the parent, and mutation is driven by the 1/5 rule.
- Note, population size is 1, and σ is global
- This is a kind of stochastic gradient method, best characterized as a local hill climber

## 1/5th rule and what it's good for

The 1/5 rule was derived for unimodal linear functions

Thus not particularly useful for practical problems but indicative of a requirement:

One should adapt as the problem difficulty changes!

#### Test functions for single-objective optimization [edit]

Test	
Functions	>

http://en.wikipedia.org/wiki/Test\_functions\_for\_optimization

Name	Plot	Formula	Global minimum	Search domain
Rastrigin function		$f(\mathbf{x}) = An + \sum_{i=1}^{n} \left[ x_i^2 - A\cos(2\pi x_i)  ight]$ where: $A = 10$	$f(0,\ldots,0)=0$	$-5.12 \le x_i \le 5.12$
Ackley's function	War and	$\begin{split} f(x,y) &= -20 \exp \left[ -0.2 \sqrt{0.5 \left( x^2 + y^2 \right)} \right] \\ &- \exp [0.5 \left( \cos 2\pi x + \cos 2\pi y \right)] + e + 20 \end{split}$	f(0,0)=0	$-5 \leq x,y \leq 5$
Sphere function	And the second second	$f(\boldsymbol{x}) = \sum_{i=1}^n x_i^2$	$f(x_1,\dots,x_n)=f(0,\dots,0)=0$	$-\infty \leq x_i \leq \infty, 1 \leq i \leq n$
Rosenbrock function		$f(m{x}) = \sum_{i=1}^{n-1} \left[ 100 ig( x_{i+1} - x_i^2 ig)^2 + (x_i - 1)^2  ight]$	$\text{Min} = \begin{cases} n = 2 & \to & f(1, 1) = 0, \\ n = 3 & \to & f(1, 1, 1) = 0, \\ n > 3 & \to & f(1, \dots, 1) = 0 \end{cases}$	$-\infty \leq x_i \leq \infty, 1 \leq i \leq n$
Beale's function		$f(x,y) = \left(1.5 - x + xy\right)^2 + \left(2.25 - x + xy^2\right)^2 + \left(2.625 - x + xy^3\right)^2$	f(3, 0.5) = 0	$-4.5 \leq x,y \leq 4.5$
Goldstein-Price function		$f(x,y) = \left[1 + (x+y+1)^2 \left(19 - 14x + 3x^2 - 14y + 6xy + 3y^2\right)\right]$ $\left[30 + (2x - 3y)^2 \left(18 - 32x + 12x^2 + 48y - 36xy + 27y^2\right)\right]$	f(0,-1)=3	$-2 \leq x,y \leq 2$
Booth's function		$f(x,y) = (x+2y-7)^2 + (2x+y-5)^2$	f(1,3)=0	$-10 \leq x,y \leq 10$
Bukin function N.6		$f(x,y) = 100\sqrt{\left y - 0.01x^2\right } + 0.01\left x + 10\right .$	f(-10,1) = 0	$-15 \leq x \leq -5, -3 \leq y \leq 3$
Matyas function		$f(x,y) = 0.26 \left( x^2 + y^2  ight) - 0.48 xy$	f(0,0)=0	$-10 \leq x,y \leq 10$

## Recombination in (µ+1)-ES

- ( $\mu$ +1)-ES allows the possibility of creating new individuals based on a combination of features of the parents, where  $\mu$  > 1
- Choose p parent vectors (1 <= p <= n), and mix characters from these p parent vectors to create a child
- Thus p=2 is similar to GA crossover



- Global intermediary recombination: Position i is average over all p parents
- Local intermediary recombination: Select two out of p parents for each child position i and take a weighted average
- Discrete recombination: copy a value from a randomly chosen parent for each child position
- Other scenarios are possible, too
- While recombination is used in ES, it is not the primary driving force, which is the previously described mutation