

# **Genetic Algorithms and Their Applications**

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# What are EAs?

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- Evolutionary Computing (EC) refers to the study of the foundations and applications of certain heuristic techniques based on the principles of natural evolution.
- The aim of designing evolutionary algorithms (EAs) is to mimic some of the processes taking place in natural evolution in algorithmic way.
- Four major categories of EAs (depending more on historical development rather than major functional differences).

$$EC = GA \cup GP \cup ES \cup EP$$

GA = Genetic Algorithms, Holland

GP=Genetic Programming, Koza

ES = Evolution Strategies, , Rechenberg, Schwefel

EP = Evolutionary Programming, Fogel

# Basic metaphor

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Although, biological evolution are not yet completely understood; strong experimental evidence is there to support the following points.

- Evolution is a process operating on *chromosomes* rather than on organisms.
- Natural selection is the mechanism that selects organisms which are *well-adapted* to the environment to reproduce more often than those which are not.
- Evolutionary process takes place during the reproduction stage that includes mutation (causing offspring to be different from parents) and recombination (combines chromosome segments of the parents to produce offspring).

# Skeleton of an EA

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Generate  $[P(0)]$  (initial population)

$t \leftarrow 0$

WHILE NOT Termination-Criteria

DO

    Evaluate  $[P(t)]$  (population at time  $t$ )

$P'(t) \leftarrow \text{Select } [P(t)]$

$P''(t) \leftarrow \text{Apply Reproduction-Operators on } [P'(t)]$

$[P(t+1)] \leftarrow \text{Replace by } [P(t), P''(t)]$

$t \leftarrow t + 1$

END

RETURN Best-Solution

# When should an EA be used?

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- The search space is large
- The search space is known not to be perfectly smooth
- The search space is not unimodal / not well understood
- Fitness function is noisy
- Search time should be minimum

# How GA Evolved?

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- Genetic Algorithms have been developed by John Holland in 1960 and his students at the University of Michigan.
- He got inspiration from the Charles Darwin's (1859) "On the Origin of Species by Means of Natural Selection or the Preservation of Favored Races in the struggle for Life".

## Their goal was :

- To abstract and rigorously explain the adaptive process of natural systems.
- To design artificial systems software that retains the important mechanics of natural systems science.

# What are genetic algorithms?

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- ❖ GAs are adaptive computational procedures modeled on the mechanics of natural genetic systems. They act as biological metaphor and try to emulate some of the processes observed in natural evolution.
- ❖ Natural evolution operates on encoding of biological entities in the form of a collection of genes called a chromosome. Similarly, GAs operate on string representation of possible solutions (individuals/ chromosomes) containing the features.
- ❖ Selection obeys Darwinian survival of the fittest (determined by the objective function) strategy. Nature acts as environment, objective function plays the same role.
- ❖ Variation is introduced mainly through genetic operations like recombination (crossover) and mutation.

# Similarities of natural evolution and GA terminologies

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<b>Natural evolution</b>	<b>GA</b>
Chromosome	String
Gene	Feature
Allele	Feature value
Genotype	String structure
Phenotype	Decoded structure



# Components of a GA

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- Population of individuals
- Encoding/decoding (of individuals) mechanism
- Objective function & associated fitness evaluation criterion
- Selection procedure
- Genetic operators (recombination/crossover, mutation)
- Probabilities to perform genetic operations
- Replacement technique
- Termination conditions

# Population

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- A set of individuals (chromosomes) representing the parameter set

$x_1, x_2, \dots, x_p$

$x_1 \rightarrow 0\ 0\ 0\ 0$

$x_2 \rightarrow 0\ 1\ 0\ 0$

... ..

$x_p \rightarrow 1\ 1\ 0\ 0$

chromosome: 0 0 0 0 0 10 0 ... 1 1 0 0

- Each member refers to a coded *possible* solution
- Fixed/variable size
- Generally, initial population is chosen randomly

# Encoding/decoding mechanism

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## Coding

- ❖ Converts parameter values into chromosome representation
- ❖ For the *continuous* valued parameters decimal to the binary conversion used.

For example       $13 == 01101$  (for 5 bit representation )

- ❖ For a parameter having *categorical* values a particular bit position in the chromosome representation is set to 1 if it comes from that category.

For example the parameter *marital status* can have values from {married, unmarried, divorced, widow}. So, unmarried == 0100  
widow==0001

- ❖ These strings (representing the parameters of a problem) are concatenated to form a chromosome.

# Decoding

❖ Decoding is the reverse of encoding.

❖ For continuous valued parameter the binary representation is converted to continuous value by the following formula

$$\text{Lower bound} + \frac{\sum_{i=0}^{\text{\#bits}-1} \text{bit}_i * 2^i}{2^{(\text{\#bits})}-1} * (\text{Upper bound} - \text{Lower bound})$$

$$01101 == 40 + (13/31) * (60 - 40) = 48.387$$

❖ For categorical valued parameters the value is found by consulting the range of the parameter.

$$0001 == \text{widow}$$

$$0100 == \text{unmarried}$$

# Evaluation and selection

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- A measure of chromosome's performance. More suitable strings should get high fitness values.
- Selection gives more chance to better fitted individuals (Mimics natural selection procedure)
- Popular selection techniques
  - ❖ Roulette wheel selection
  - ❖ Linear normalization selection
  - ❖ Tournament selection
  - ❖ Stochastic Universal Sampling

# Roulette wheel selection

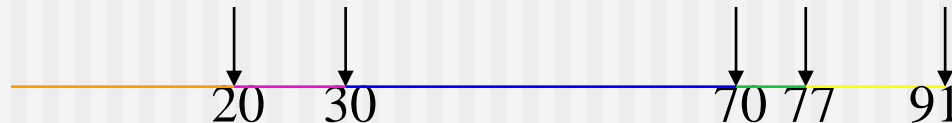
- Sum the fitness of all the chromosomes of the population. Call it *total-fitness*.
- Generate a random number  $n$  in  $[0, \textit{total-fitness}]$
- Return the first chromosome whose fitness when added to the fitness of the preceding population member is greater or equal to  $n$ .

## Example:

Let there be five chromosomes with fitness 20, 10, 40, 7, 14  
Then *total-fitness*=91.

Say, the random number drawn ( $n$ ) is 45.

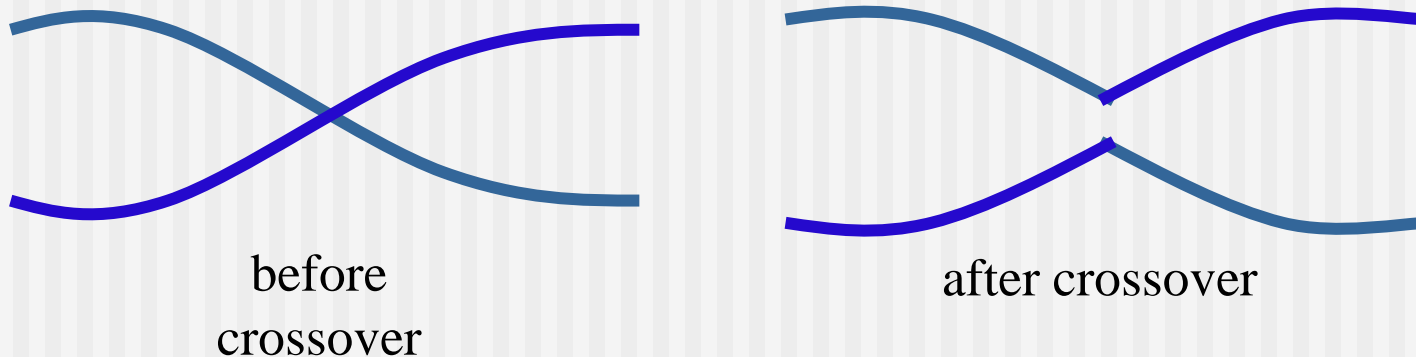
Select the 3<sup>rd</sup> chromosome (since  $20+10+40 > 45$ ).



# Recombination/crossover

- ❖ Exchange of information; exploitation
- ❖ Choose mating pairs (from the selected chromosomes).
- ❖ Check (using  $p_c$ ) whether this pair should go for recombination or not. If yes, interchange chromosome segments using cross-sites.  
→ one point, two point, multi point, uniform,...

## One point crossover

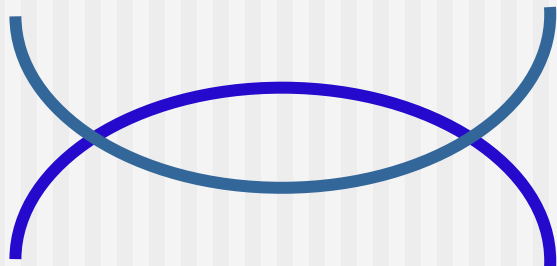


➤ Parent1: x y x y x y x y child1: x y x y x b a b

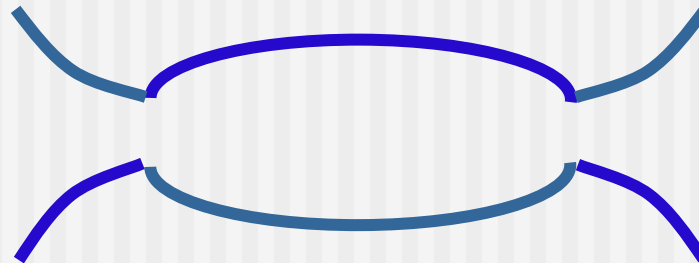
Parent2: a b a b a b a b child2: a b a b a y x y

# Recombination (contd.)

## Two point crossover



before  
crossover



after  
crossover

➤ parent1: xy xy xy xy

parent2: ab ab ab ab

child1: xy ab a y xy

child2: ab xy x b ab



# Mutation

- ❖ Introduces diversity, helps to regain lost genetic material
- ❖ Exploration
- ❖ Bit mutation.
- ❖ Check (using  $p_m$ ) whether this bit should be mutated or not.  
If yes, flip the bit.
- ❖ 00**1**000 → 00**0**000

## Probabilities to perform genetic operations

- ❖ May be fixed or made variable.
- ❖  $p_c$  : 0.6 to 0.9       $p_m$  : 0.001 to 0.01

# Replacement techniques

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❖ **Generational** - replaces all the individuals at a time

- Create  $N$  children through reproduction

- Replace the old population with these new individuals

❖ **Steady state** - replaces a few individuals at a time

- Create  $m$  ( $< N$ ) children through reproduction

- Delete  $m$  members of the population to make room for them

- Insert the children into the population

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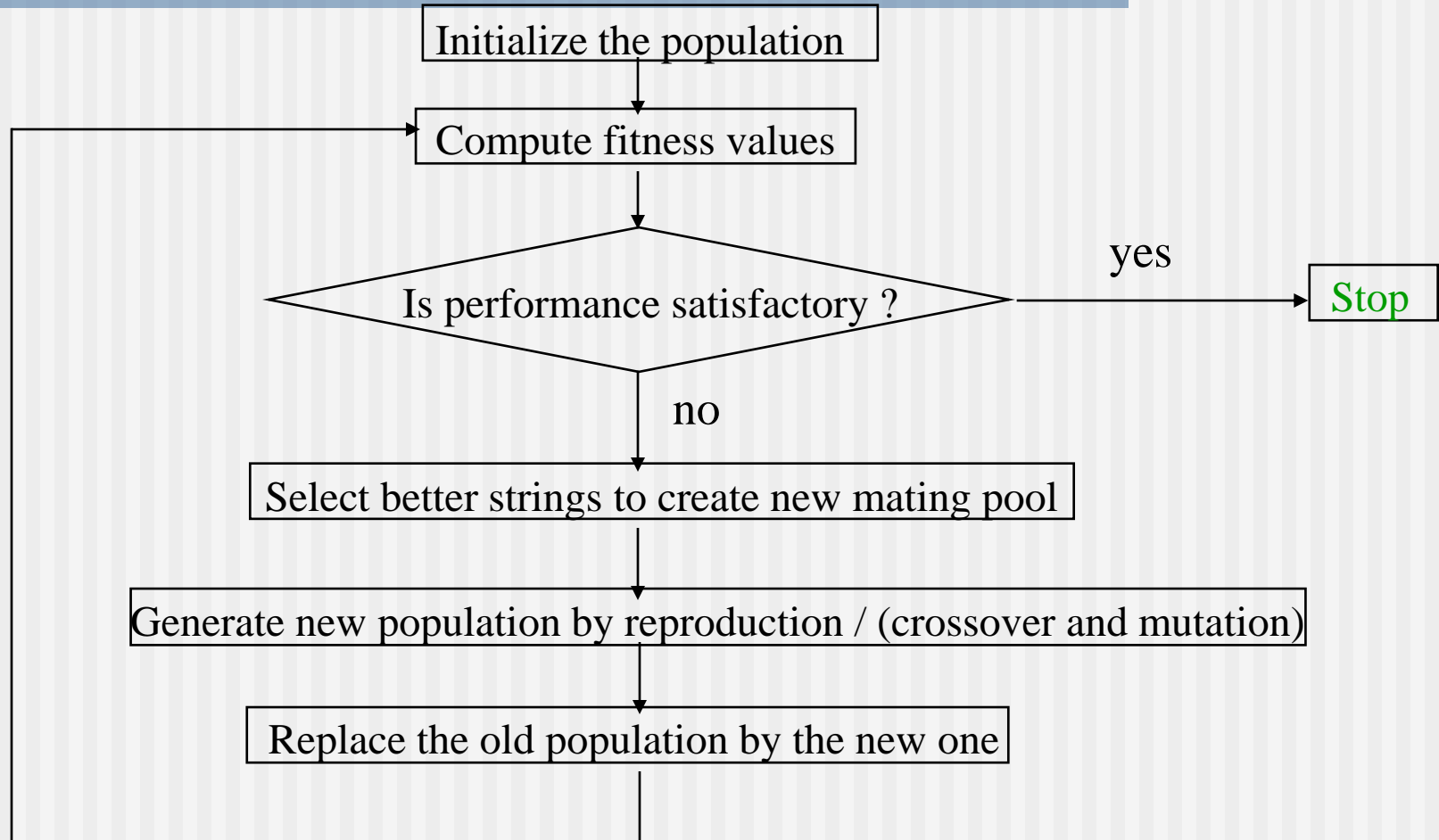
## Terminating criterion

- ❖ Execute for a fixed number of generations/iterations.
- ❖ Until a string with a certain fitness value is located.
- ❖ Until the population attains a certain degree of homogeneity (most of the individuals become similar).

## Elitism *(optional)*

Keeps track of /store the best solution obtained so far.

# Flow diagram of a GA



# Example 1

Maximize  $f(x) = x^2$

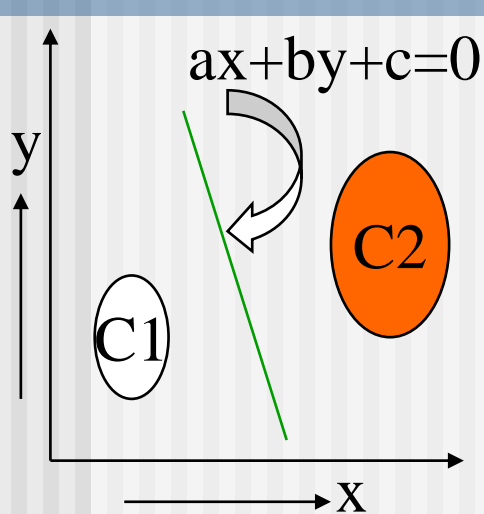
String#	Initial Pop	x-value	$f(x)=x^2$	Expected count ( $f_i/f_{av}$ )	Actual count from r-wheel
1	01101	13	169	0.58	1
2	11000	24	576	1.97	2
3	01000	8	64	0.22	0
4	10011	19	361	1.23	1

sum = 1170; average = 293; max = 576

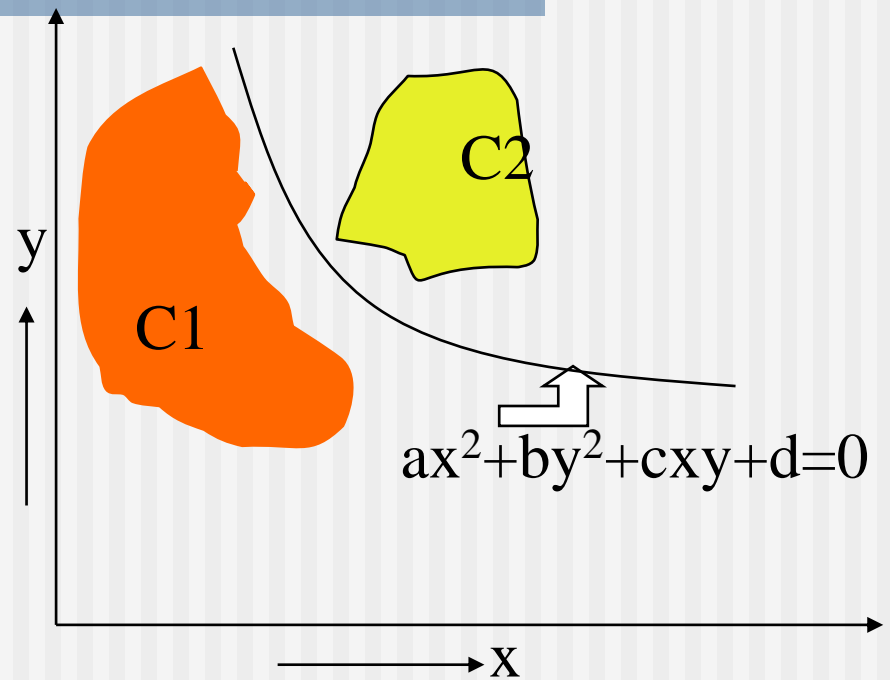
Mating pool (after reproduction)	Mate	cross-site (random select)	New Pop	x-value	$f(x)=x^2$
01101	2	4	01100	12	144
11000	1	4	11001	25	625
11000	4	2	11011	27	729
10011	3	2	10000	16	256

sum = 1754; average = 439; max = 729

## Example 2 (classification)



Parameters :  $a, b$  &  $c$



Parameters :  $a, b, c$  &  $d$

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class	P length	S length				
c1						p1
c2						p2
c1						p3

# Distinguishing characteristics of GAs

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- ❖ Multiple point searching (population based)
  - implicit parallelism
  - sometimes helps to prevent getting stuck to local optima
- ❖ Works on coded parameter set
  - resolution of the solutions can be controlled
- ❖ Search space may be discontinuous
- ❖ Uses probabilistic state transition rules
- ❖ Does not require any auxiliary information



# Deviation from conventional GAs

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- ❖ Distributed GA
- ❖ Parallel GA
- ❖ Structured GA
- ❖ Hybridization with neural networks and fuzzy logic
  - Fuzzy-GA
  - Neuro-fuzzy GA
  - Neuro-GA
- ❖ Hybridization with simulated annealing

# Applications areas by domain

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- ❖ Numerical, combinatorial and constrained optimization
- ❖ Scheduling, TSP, Graph Coloring
- ❖ Industrial design by parameterization
- ❖ Network design by construction, routing
- ❖ Automatic programming - evolves computer programs for specific tasks (Genetic Programming)
- ❖ Pattern recognition - classification, clustering, prediction
- ❖ Image processing --- segmentation, enhancement
- ❖ Data mining --- rule mining, clustering
- ❖ Bioinformatics – docking, prediction of structure of protein
- ❖ Economics - financial prediction
- ❖ Molecular biology – molecular conformation

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Thank you