

# Evolutionary Algorithms

Inteligencia Artificial en los Sistemas de Control Autónomo

## Objectives

- Describe the most relevant EAs

## Bibliography

- Eiben, A.E. and Smith, J.E. Introduction to Evolutionary Computing. Springer 2003.

# Table of Contents

## 1. Genetic Algorithms

- Introduction
- Representation
- Mutation
- Recombination
- Selection

## 2. Genetic Programming

- Introduction
- Representation
- Mutation
- Recombination
- Initialization
- Bloat in Genetic Programming

## 3. Evolution Strategies

- Introduction
- Representation
- Mutation
- Recombination
- Parent and survivor selection

# Genetic Algorithms

## Introduction (I)

Introduced by Holland in the 70's

- John H. Holland ``Adaptation in Natural and Artificial Systems'', MIT Press
- GA is the most popular EA
- Usually EAs confused with GA

Canonical GA (which is not canonical)

- Fixed length strings
- Binary codification
- Holland's Theorem

Representation	Bit strings
Recombination	r-point
Mutation	Bit flip
Parent select	Fitness prop
Survivor select	Generational

# Genetic Algorithms

## Introduction (II)

GAs are a family of algorithms, with common features

- Representation in strings, named **chromosomes**
- Mutation and recombination
- Usually fixed length

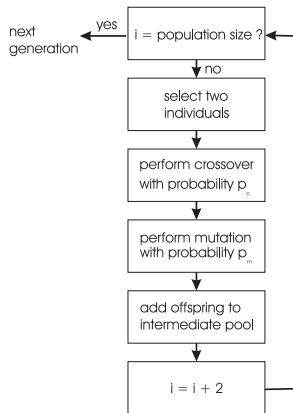
GAs are like a toolbox with customizable components

- Representations, genetic operators, selections mechanism, ...
- These components are interdependent

Rule of thumb: Small genotype changes  $\Rightarrow$  Small phenotype changes

# Genetic Algorithms

## Introduction (III)



# Genetic Algorithms

## Representation: Binary



One of the oldest and widely used codifications

- Consequence of Holland's Theorem
- Strong historical influence

Often used to codify non-binary information (not recommended)

- Pure binary codification
- Gray coding
- Custom codification



Hint: Use binary codification to represent binary information

# Genetic Algorithms

## Representation: Integer

4	3	2	1	0	4	2	3	3
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Chromosome as a sequence of integers

- More natural codification for many problems
- Optimization of integer values
- Integer representation ( $\{1, 2, 3, 4\} = \{\text{North, East, South, West}\}$ )



# Genetic Algorithms

## Representation: Floating-point

1.1	0.2	3.0	33.2	0.0	-3.2	130.1	88.3	-7.1
-----	-----	-----	------	-----	------	-------	------	------

Chromosome as a sequence of floating-point values

- Common in optimization problems
- Solutions with continuous nature

# Genetic Algorithms

## Representation: Permutation

4	3	2	5	6	1
---	---	---	---	---	---

Some problems involve order

- Sequence of integers
- No repeated numbers
- Range of valid numbers
- Special genetic operators

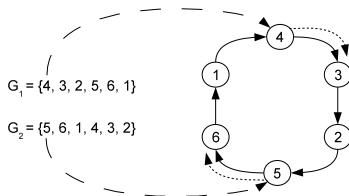
Information can be contained in

- The locus (position)

$$[3, 1, 2, 4] \Rightarrow [C, A, B, D]$$

- The allele (value)

$$[3, 1, 2, 4] \Rightarrow [B, C, A, D]$$



Integer codification to solve TSP

# Genetic Algorithms

## Mutation

**Mutation:** Genetic operator that uses one parent

- Introduces randomness into the genotype
- Depends on representation

Main objectives

- Avoid local minima (premature convergence)
- Enhances exploration

Often dependent on the **mutation rate**

- Significant influence in the algorithm behaviour
- Higher mutation rate, higher exploration

# Genetic Algorithms

## Mutation for binary representations

Flip bit with probability  $p_m$



Optimal  $p_m$  depends on the problem and goals

- Need of high fitness population
- Need of high fitness individual
- Need of genetic diversity
- Modality of the problem
- Algorithm dynamics

Rule of thumb:  $p_m = \frac{1}{\text{length}}$

# Genetic Algorithms

## Mutation for integer representations

Two main mutations applied to each gene

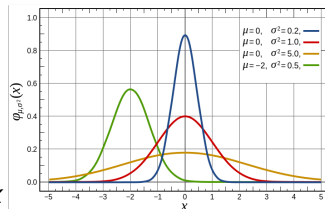
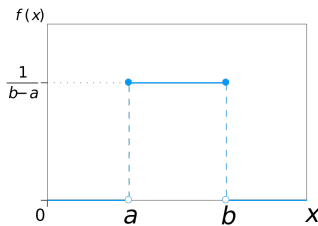
- **Random resetting:** Choose new random value with  $p_m$
- **Creep mutation:** Add small (positive or negative) random value with  $p_m$

# Genetic Algorithms

## Mutation for floating-point representations

Set new value with value drawn from a distribution

- **Uniform mutation** Choose new random value from  $[L, U]$  with  $p_m$
- **Non-uniform mutation** Usually adding a value drawn from a zero-mean gaussian distribution



# Genetic Algorithms

## Mutation for permutation representations

Genes are no longer independent

- No gene mutation,  $p_m$  affects the whole chromosome

### Swap mutation



### Insert mutation



### Scramble mutation



### Inversion mutation



# Genetic Algorithms

## Recombination

Recombination creates one individual from two or more parents

- Also known as crossover (specially for two parents)
- Basic feature in GA
- Parents selection mechanism needed

Usually applied to all new individuals

- Not used when elitism is applied
- Sometimes applied with  $p_c \in [0, 5, 1]$

Objectives of recombination

- Combine parents' behavior  $\Rightarrow$  No new genetic material
- Constructive role
- Enhances exploitation

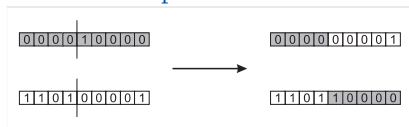


# Genetic Algorithms

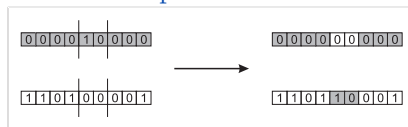
## Recombination: Binary and integer representations

Three crossover mechanisms for binary and integer encodings

One-point crossover



Two-points crossover



Uniform crossover



# Genetic Algorithms

## Recombination: Floating point representations (I)

### Discrete recombination

- Analogous to binary recombination
- No new genetic material

### Arithmetic recombination

- Combines the parents' genes
- Weighted sums of genes:  $z_i = \alpha x_i + (1 - \alpha) y_i$
- Usually,  $\alpha = 0,5$  (average values)
- Different arithmetic recombinations

# Genetic Algorithms

## Recombination: Floating point representations (II)

### Whole arithmetic recombination (All genes are included)



### Simple arithmetic recombination (Similar to one-point crossover)



### Single arithmetic recombination (Similar to uniform crossover)

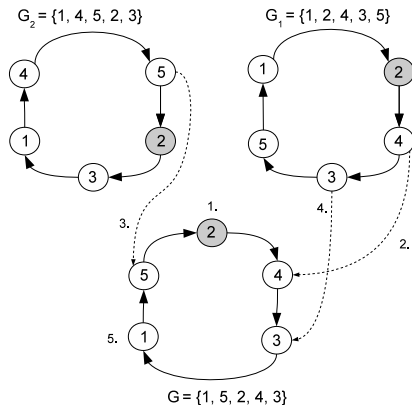


# Genetic Algorithms

## Recombination: Permutation representations

### Specialized recombinations

- Partially Mapped Crossover
- Edge Crossover
- Order Crossover
- Cycle Crossover



# Genetic Algorithms

## Selection

Two purposes for selection

- Parent selection: Individuals to generate offspring
- Survivor selection: Individuals to replace

Usually same methods applied to both

# Genetic Algorithms

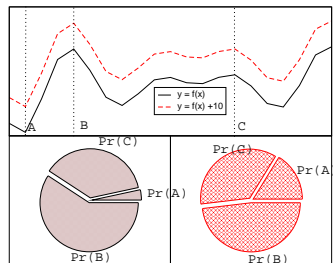
## Selection: Fitness Proportional Selection

Selection probability proportional to fitness

- Premature convergence
- Lack of selective pressure for close fitness values
- Selective pressure not customizable
- Susceptibility to function transposition

Historically relevant

$$p_s = \frac{f_i}{\sum_{j=1}^n f_j}$$



# Genetic Algorithms

## Selection: Ranking Selection

Selection probability proportional to rank

- Individuals are sorted by fitness
- Arbitrary rank to probability mapping
- Avoid problems with super individuals
- Selective pressure independent of fitness
- Selective pressure not customizable

### Linear mapping

$$P_{lin,rank}(i) = \frac{(2-s)}{\mu} + \frac{2i(s-1)}{\mu(\mu-1)}$$

$$1,0 < s < 2,0$$

### Exponential mapping

$$P_{exp,rank}(i) = \frac{1-e^{-i}}{c}$$

$c$  = normalization factor

## Genetic Algorithms

## Selection: Tournament Selection

## Algorithm of tournament size $k$

1. Select randomly  $k$  chromosomes
2. Compute their fitness
3. Select the fittest one
4. Go to 1

## Customizable selective pressure

- Depends on  $k$  and  $\mu$

## De facto standard

- Good for parallel computation
- Efficient implementation

Usually  $k = 2$  in GA, in GP  $k = 7$



# Genetic Algorithms

## Selection: Survival selection

Two strategies

- Generational (all the population is replaced)
- Steady-state (partial replacement)

Survival selection algorithms

- Fitness-Based Replacement (inverse of the previous ones)
- Age-Based Replacement
- Elitism

# Genetic Programming

## Introduction (I)

GP is a family of algorithms

- Evolve programs
- Self-programming computers
- GP, Linear GP, Cartesian GP, EDA, ...

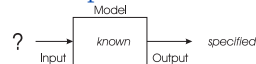
GP introduced by Koza in the 90's

Koza, J.R. ``Genetic Programming: On the Programming of Computers by Means of Natural Selection'', MIT Press. 1992

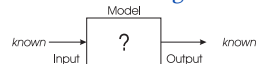
GA and ES focused on optimization

- GP focused on Machine Learning

### Optimization



### Modelling



### Simulation



# Genetic Programming

## Introduction (II)

Example: Credit scoring problem within a bank. Develop a model describing good customers

Id	Children	Salary	Status	Credit
Id-1	2	45.000	Married	0
Id-2	0	30.000	Single	1
Id-3	1	40.000	Married	1
Id-4	2	60.000	Divorced	1
...				
Id-X	2	50.000	Married	1

Possible model:

IF (children=2) AND (Salary>80.000) THEN good ELSE bad

# Genetic Programming

## Introduction (III)

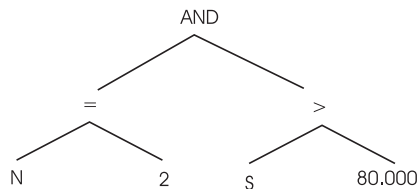
General form

IF (Formula)  
THEN good  
ELSE bad

In EC terms

Phenotype: Formula

Fitness: Classification accuracy



(children=2) AND (Salary>80.000)

# Genetic Programming

## Representation (I)

GP representation differs in two aspects

- Nonlinear structure
- Variable size

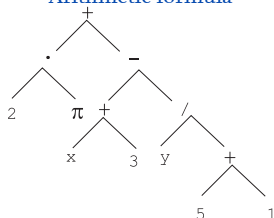
New representation and genetic operators

- Same selection (done in phenotypic space)

# Genetic Programming

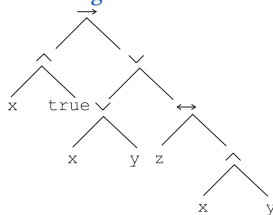
## Representation (II)

Arithmetic formula



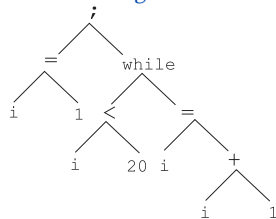
$$\left(2\pi + ((x+3) - \frac{y}{5+1})\right)$$

Logical formula



$$(x \wedge \text{true}) \rightarrow ((x \vee y) \vee (z \leftrightarrow (x \vee y)))$$

Program



```
i=1;
while (i<20) {
  i = i+1;
}
```

# Genetic Programming

## Representation (III)

Two types of nodes

- **Function set** Internal nodes. It has an associated number of attributes
- **Terminal set** Leaves of the tree

Danger: Inviabile trees

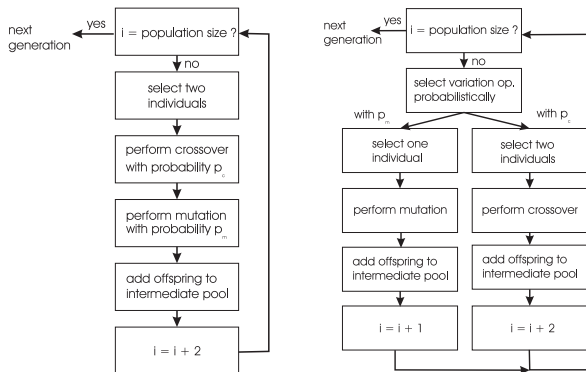
- Grammar-aware GP variants
- Strongly Typed Genetic Programming (STGP), Grammatical Evolution (GE), ...

(Complex representation example)

# Genetic Programming

## Mutation (I)

### Application of genetic operators in GP contrast to GA





# Genetic Programming

## Mutation (II)

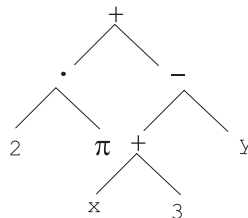
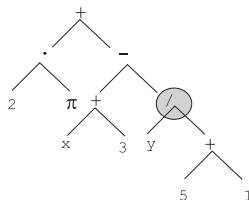
### Subtree mutation

1. Select a random node
2. Delete subtree
3. Add new random subtree

### Parameters

- Probability of choosing a terminal node

Highly correlated with **code bloat**



# Genetic Programming

## Mutation (III)

### Alternative mutation operators

- Size-fair subtree mutation
- Node replacement mutation (point mutation)
- Hoist mutation
- Shrink mutation

# Genetic Programming

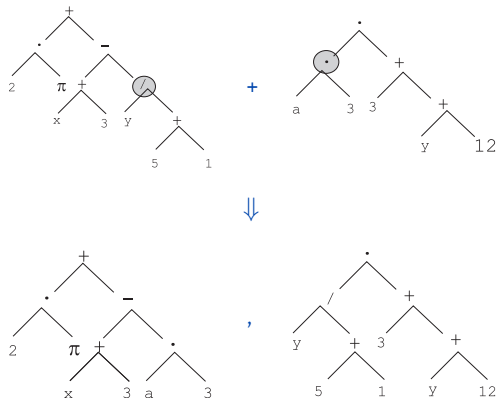
## Recombination (I)

### Subtree crossover

1. Take a random node from both parents
2. Swap subtrees

### Parameters

- Probability of choosing a terminal node



# Genetic Programming

## Recombination (II)

### Alternative recombination operators

- Homologous crossover
- Uniform crossover
- Size-fair crossover
- Node replacement mutation (point mutation)
- Hoist mutation
- Shrink mutation

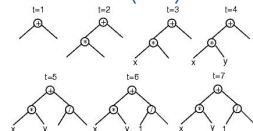
# Genetic Programming

## Initialization

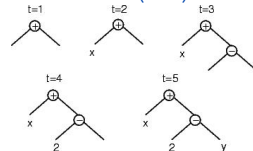
### Three initialization methods

- **Full.** Introduces non-terminals nodes until max depth
- **Grow.** Introduces terminal or non-terminal with equal probability
- **Ramped half-n-half.** Applies full or grow with equal probability

### Full ( $D=2$ )



### Grow ( $D=2$ )



# Genetic Programming

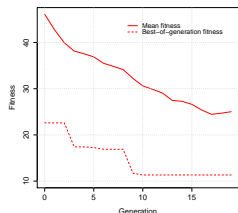
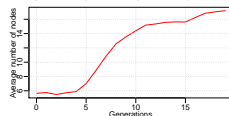
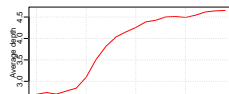
## Bloat in Genetic Programming

**Code bloat:** Uncontrolled grow of tree sizes

- Intrinsic to variable-length representations
- Undesirable effects
- Perhaps, the worse problem in GP

Countermeasures

- Depth limitation in genetic operators
- Parsimony pressure
- Tree pruning
- Multiobjective techniques



# Genetic Programming

## Example of reporting

Cuadro 1: Main parameters used to obtain the approximations for secrets ID in the Genetic Tango attack against David-Prasad authentication protocol.

Parameter	ID
Population	500
Generations	10
Terminal Set	A, B, D, E, F, $P_{ID1}$ , $P_{ID2}$
Function set	And, or, xor
Fitness	Hamming distance to secret
Fitness tags	5
Fitness sessions	100
Min. depth	1
Max. depth	3
Selection	Lexicographic tournament
Tournament size	4
Crossover	0.9
Reproduction	0.1
Elitism size	1
Terminals	0.1
Non terminals	0.9
Initialization	Rampld H-H

# Evolution Strategies

## Introduction (I)

Introduced by Rechenberg and Schwefel in the 60's

- Motivated by wing shape optimization
- Real-function optimization

ES properties

- Emphasis on mutation
- Mutation is gaussian noise
- Self-adaptation

Representation	Real-valued vectors
Recombination	Discrete
Mutation	Gaussian perturbation
Parent selection	Uniform
Survivor selection	$(\mu, \lambda)$ or $(\mu + \lambda)$
Speciality	Self-adaptation



# Evolution Strategies

## Introduction (II)

### Example of basic ES

- Representation: Vector of real values
- Recombination: Not used
- Mutation: Gaussian noise with **step-size**  $\sigma$

### Adaptative $\sigma$ (1/5 rule)

- Theoretical foundations
- Based on the ratio of success mutations ( $p_s$ )
- After  $k$  iterations a new  $\sigma$  is computed

$$\sigma = \begin{cases} \sigma/c & \text{if } p_s > 1/5, \\ \sigma \cdot c & \text{if } p_s < 1/5, \\ \sigma & \text{if } p_s = 1/5 \end{cases}$$

where  $0,817 \leq c \leq 1$  is a parameter

# Evolution Strategies

## Representation

Nowdays ES is usually self-adapted

- Step size ( $\sigma$ ) is included in the genotype
- Evolution includes variables and parameters

One or more  $\sigma$  values

- One  $\sigma$ :  $\langle \underbrace{x_1, x_2, \dots, x_n}_{\bar{x}}, \sigma \rangle$
- Several:  $\sigma : \langle \underbrace{x_1, x_2, \dots, x_n}_{\bar{x}}, \underbrace{\sigma_1, \sigma_2, \dots, \sigma_{n_\sigma}}_{\bar{\sigma}} \rangle$

# Evolution Strategies

## Mutation

Genetic operators to modify  $\sigma$

- Mutation with one step size:

$$\begin{aligned}x'_i &= x_i + N_i(0, \sigma') \\ \sigma' &= \sigma \cdot e^{\cdot N(0, \tau)}, \tau \propto 1/\sqrt{n}\end{aligned}$$

$\tau$  is analogous to learning rate in ANN

- Mutation with n step sizes:

$$\begin{aligned}x'_i &= x_i + N_i(0, \sigma_i) \\ \sigma' &= \sigma \cdot e^{\cdot N(0, \tau') + N_i(0, \tau)}\end{aligned}$$

with  $\tau' \propto 1/\sqrt{2n}$  and  $\tau \propto 1/\sqrt{2\sqrt{n}}$

# Evolution Strategies

## Recombination

Secondary operator in ES

- **Discrete recombination.** Like uniform crossover in GA
- **Intermediate recombination.** Like arithmetic crossover in GA

ES tends to use **global recombination**

- More than two parents

# Evolution Strategies

## Parent and survivor selection

The whole population is seen as parent

- Select individual with uniform probability
- No selective pressure in parent selection

After creating the offspring, the  $\lambda$  fittests individuals are selected

- Deterministic procedure

Two selection mechanisms depending on who can be selected

- $(\mu, \lambda)$  **selection**. Only the offspring.
- $(\mu + \lambda)$  **selection**. Parents and offspring

$(\mu, \lambda)$  selection is more popular