# **Evolutionary Algorithms**

Inteligencia Artificial en los Sistemas de Control Autónomo Máster Universitario en Ingeniería Industrial

Departamento de Automática





### Objectives

• Describe the most relevant EAs

### Bibliography

 $\bullet~$  Eiben, A.E. and Smith, J.E. Introduction to Evolutionary Computing. Springer 2003.

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### 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	1-point
Mutation	Bit flip
Parent select	Fitness prop
Survivor select	Generational



### Introduction (II)

GAs are a family of algorithms, with common features

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

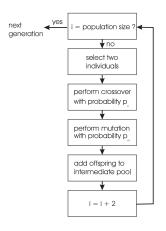
GAs are like a toolbox with customizable components

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

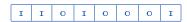
Rule of thumb: Small genotype changes ⇒ Small phenotype changes



### Introduction (III)



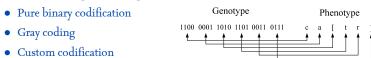
### Representation: Binary



One of the oldest and widely used codifications

- Consequence of Holland's Theorem
- Strong historical influence

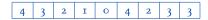
Often used to codify non-binary information (not recommended)



Hint: Use binary codification to represent binary information



### Representation: Integer



#### Chomosome as a sequence of integers

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



Representation: Floating-point



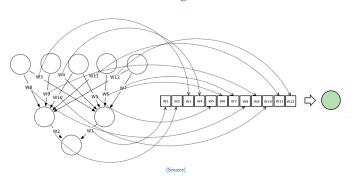
Chomosome as a sequence of floating-point values

- Common in optimization problems
- Solutions with continous nature

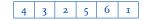


### Representation: Floating point (II)

### ANN encoding with a GA



### Representation: Permutation



Some problems involve order

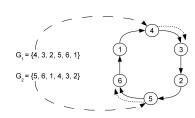
- Sequence of integers
- No repeated numbers
- Range of valid numbers
- Special genetic operators

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

#### 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



### 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:  $p_m = \frac{1}{l_{enoth}}$ 



### Mutation for integer representations

Two main mutations applied to each gene

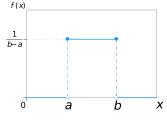
- Random resetting: Choose new random value with p<sub>m</sub>
- Creep mutation: Add small (positive or genative) random value with p<sub>m</sub>

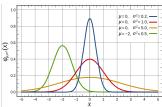


### Mutation for floating-point representations

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







### Mutation for permutation representations

### Genes are no longer independent

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

Swap mutation	Insert mutation	
123456789	123456789	
Scramble mutation	Inversion mutation	
123456789	123456789	

#### 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  $\mathfrak{p}_{\mathfrak{c}} \in [0,5,1]$

#### Objectives of recombination

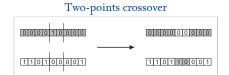
- Combine parents' behavior ⇒ No new genetic material
- Constructive role
- Enhances explotation

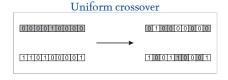


### Recombination: Binary and integer representations

#### Three crossover mechanisms for binary and integer encodings

### 





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



Recombination: Floating point representations (II)

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



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

0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 0.10.20.30.40.50.60.50.50.6

0.30.20.30.20.30.20.50.50.6

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

0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.5 0.9 0.3 0.2 0.3 0.2 0.3 0.2 0.3 0.2 0.3 0.3 0.2 0.3 0.2 0.3 0.2 0.3 0.5 0.3

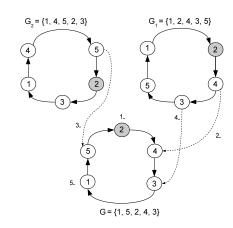


0.3 0.2 0.3 0.2 0.3 0.2 0.3 0.2 0.3

## Recombination: Permutation representations

### Specialized recombinations

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



#### Selection

#### Two purposes for selection

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

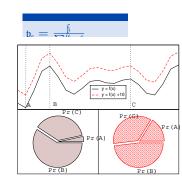
Usually same methods applied to both

### 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





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

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

### Exponential mapping

$$P_{exp_r\alpha nk}(\mathfrak{i})=\frac{1-e^{-\mathfrak{i}}}{c}$$

c = normalization factor



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

ullet Depends on  ${
m k}$  and  $\mu$ 

De facto standard

- Good for parallel computation
- Efficient implementation

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



Selection: Survival selection

#### Two strategies

- Generational (all the population is remplaced)
- Steady-stade (partial remplacement)

#### Survival selection algorithms

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



### 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

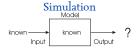
Koza, J.R. "Genetic Programming: On the Programming of Computers by Means of Natural Selection", MIT Press. 1992

#### GA and ES focused on optimization

GP focused on Machine Learning







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	I
Id-3	I	40.000	Married	I
Id-4	2	60.000	Divorced	I
Id-X	2	50.000	Married	I

Possible model:

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

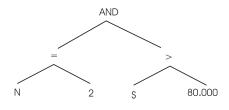


Introduction (III)

#### General form

IF (Formula)
THEN good
ELSE bad

In EC terms Phenotype: Formula Fitness: Classification accuracy



(children=2) AND (Salary>80.000)

### Representation (I)

#### GP representation differs in two aspects

- Nonlinear structure
- Variable size

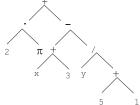
New representation and genetic operators

• Same selection (done in phenotipic space)



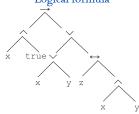
### Representation (II)



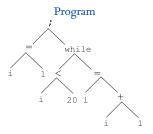


$$\left(2\pi\!+\!\left((\mathbf{x}\!+\!3)\!-\!\frac{\mathbf{y}}{5+1}\right)\right)$$

### Logical formula



$$\begin{array}{l} (x \wedge \mathsf{true}) \to \\ ((x \vee \gamma) \vee (z \leftrightarrow (x \vee \gamma))) \end{array}$$



Representation (III)

#### Two types of nodes

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

#### Danger: Inviable trees

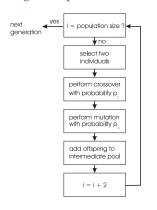
- Grammar-aware GP variants
- Strongly Typed Genetic Programming (STGP), Grammatical Evolution (GE), ...

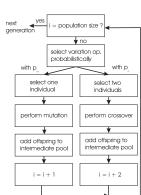
(Complex representation example)



### Mutation (I)

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

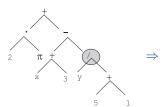




### Mutation (II)

#### Subtree mutation

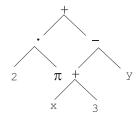
- 1. Select a random node
- 2. Delete subtree
- 3. Add new random subtree



#### Parameters

 Probability of choosing a terminal node

Highly correlated with code bloat



Mutation (III)

#### Alternative mutation operators

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



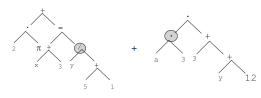
### Recombination (I)

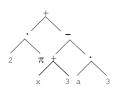
#### Subtree crossover

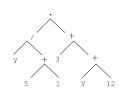
- Take a random node from both parents
- 2. Swap subtrees

#### Parameters

• Probability of choosing a terminal node







## Recombination (II)

### Alternative recombination operators

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



#### 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

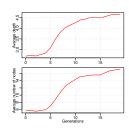
## Bloat in Genetic Programming

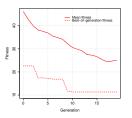
## Code bloat: Uncontrolled grow of tree sizes

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

#### Countermeasures

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







# 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	I
Max. depth	3
Selection	Lexicographic tournament
Tournament size	4
Crossover	0.9
Reproduction	O.I
Elitism size	I
Terminals	0.1
Non terminals	0.9
Initialization	Rampled H-H

# 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 (ps)
- After k iterations a new  $\sigma$  is computed

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

where  $0.817 \le c \le 1$  is a parameter



## 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  $\sigma$ :  $\langle \underbrace{x_1, x_2, ..., x_n}_{\overline{x}}, \sigma \rangle$
- $\bullet \ \, \text{Several:} \, \sigma: \big\langle \underbrace{x_1, x_2, ..., x_n}_{\Xi}, \underbrace{\sigma_1, \sigma_2, ..., \sigma_{n_\sigma}}_{\bar{\sigma}} \big\rangle$

#### Mutation

#### Genetic operators to modify $\sigma$

Mutation with one step size:

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

au is analogous to learning rate in ANN

Mutation with n step sizes:

$$\mathbf{x}_{i}' = \mathbf{x}_{i} + \mathbf{N}_{i}(0, \sigma_{i})$$

$$\sigma' = \sigma \cdot e^{\cdot \mathbf{N}(0, \tau') + \mathbf{N}_{i}(0, \tau)}$$

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



## 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



#### 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 offpring.
- $(\mu + \lambda)$  selection. Parents and offpring

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



# Search phases

Initial phase: Random distribution, high genetic diversity Advanced phase: Begins to converge Convergence: Around one or few points, low genetic diversity

Premature convergence if population not located in global maxima





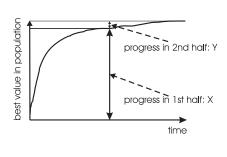


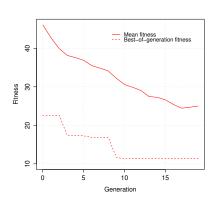
(Animation)



# Working with an Evolutionary Algorithm

## Fitness dynamics



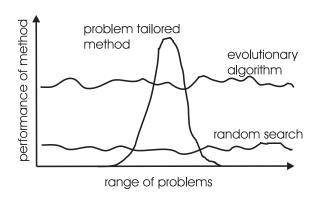


Few long runs or many short runs?



# Working with an Evolutionary Algorithm

## When EAs are useful





# Working with an Evolutionary Algorithm

#### Advanced EAs

- Multiobjective Evolutionary Algorithms (MOEAs)
- Optimization with constrains
- Coevolution
- Dinamic optimization
- Islands models
- Memetic algorithms
- Hyperheuristics

