Evolutionary Algorithms

Inteligencia Artificial en los Sistemas de Control Autónomo Máster en Ciencia y Tecnología desde el Espacio

Departamento de Automática





Objectives

• Describe the most relevant EAs

Bibliography

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

Table of Contents

- 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



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

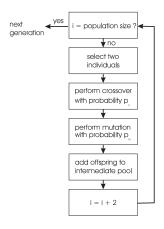
GAs are like a toolbox with customizable components

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

Rule of thumb: Small genotype changes ⇒ Small phenotype changes



Introduction (III)



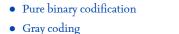
Representation: Binary



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)

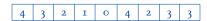


Custom codification



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
- $\bullet \ \ Integer \ representation (\{1,2,3,4\} = \{ \mathsf{North}, \mathsf{East}, \mathsf{South}, \mathsf{West} \}) \\$

Representation: Floating-point



Chomosome as a sequence of floating-point values

- Common in optimization problems
- Solutions with continous nature



Representation: Permutation

4	2	2	_	6	т
4	3		5	U	1

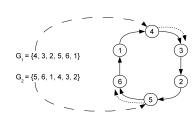
Some problems involve order

- 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

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: $p_m = \frac{1}{length}$



Mutation for integer representations

Two main mutations applied to each gene

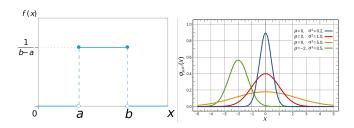
- \bullet Random resetting: Choose new random value with p_m
- Creep mutation: Add small (positive or genative) random value with p_m



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

 \bullet 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 $p_c \in [0,5,1]$

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

One-point crossover



Two-points crossover



Uniform crossover





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)



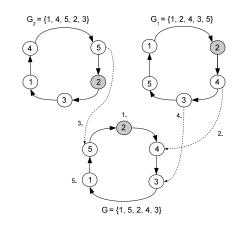
Single arithmetic recombination (Similar to uniform crossover)



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

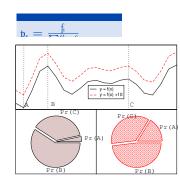


Selection: Fitness Proportional Selection

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

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

Exponential mapping

$$P_{exp_rank}(i) = \frac{1-e^{-i}}{c}$$

c = normalization factor



Selection: Tournament Selection

Algorithm of tournament size k

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

Customizable selective pressure

• Depends on k and μ

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, ...

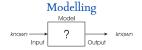
GP introduced by Koza in the 90's

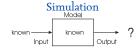
Koza, J.R. "Genetic Programming: On the Programming of Computers by Means of Natural Selection", MIT Press. 1992

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	О
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

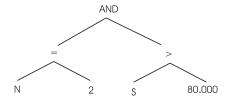


Introduction (III)

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

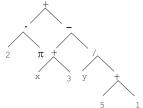
New representation and genetic operators

• Same selection (done in phenotipic space)



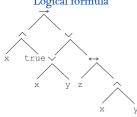
Representation (II)



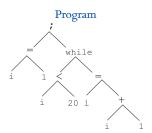


$$\left(2\pi + \left((x+3) - \frac{\gamma}{5+1}\right)\right)$$

Logical formula



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



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

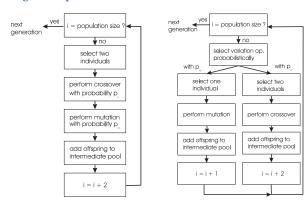
- Grammar-aware GP variants
- $\bullet~$ Strongly Typed Genetic Programming (STGP), Grammatical Evolution (GE), ...

(Complex representation example)



Mutation (I)

Application of genetic operators in GP contrast to GA





Mutation (II)

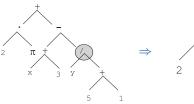
Subtree mutation

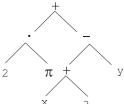
- T. Select a random node
- 2. Delete subtree
- 3. Add new random subtree

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



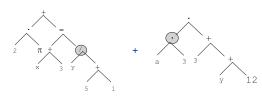
Recombination (I)

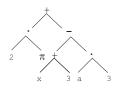
Subtree crossover

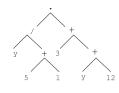
- Take a random node from both parents
- 2. Swap subtrees

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

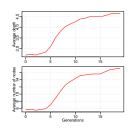
Bloat in Genetic Programming

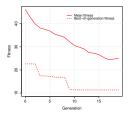
Code bloat: Uncontrolled grow of tree sizes

- 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	IO
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	O.I
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	(μ,λ) or $(\mu+\lambda)$
Speciality	Self-adaptation



Introduction (II)

Example of basic ES

- Representation: Vector of real values
- Recombination: Not used
- Mutation: Gaussian noise with step-size σ

Adaptative σ (1/5 rule)

- Theoretical foundations
- Based on the ratio of success mutations (p_s)
- After k iterations a new σ 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 (σ) is included in the genotype
- Evolution includes variables and parameters

One or more σ values

• One
$$\sigma$$
: $\langle \underbrace{x_1, x_2, ..., x_n}_{\bar{x}}, \sigma \rangle$

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



Mutation

Genetic operators to modify σ

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 λ fittests individuals are selected

• Deterministic procedure

Two selection mechanisms depending on who can be selected

- (μ, λ) **selection**. Only the offpring.
- ullet $(\mu + \lambda)$ selection. Parents and offpring

 (μ, λ) selection is more popular

