Evolutionary Programming and Genetic Programming

Motto:

"How can computers learn to solve problems without being explicitly programmed? In other words, how can computers be made to do what is needed to be done, without being told exactly how to do it?"

— Attributed to Arthur Samuel (1959)

The origins:

- L. Fogel (1960) development of methods, inspired by the natural evolution, which generate, in an automatic way, systems with some intelligent behavior;
- D. Fogel (1990) in the last two decades the evolutionary programming became more oriented toward solving problems (optimization and design) than simulating generic intelligent behaviours

Particularities

- Various encoding variants (e.g. real vectors, state diagrams, neural networks structures)
- Based only on mutation, no recombination
- Current variants: self-adaptive

First (traditional) direction :

- Evolve systems (e.g. finite state machine) with prediction ability
- The fitness of such a structure is measured by analyzing the behavior of the system = prediction ability
- The fitness is a quality measure related to the behaviour of the system

Finite State Machines (FSM):

FSM = (S, I, O, T,s0) S – set of states I – input alphabet O – output alphabet T:SxI->SxO - transition rules s0 – initial state

A simple test problem:

design a FSM to check if a binary string has an even or an odd number of elements equal to 1 (parity problem)

- S={even,odd}
- I={0,1}
- O={0,1}

FSM output:

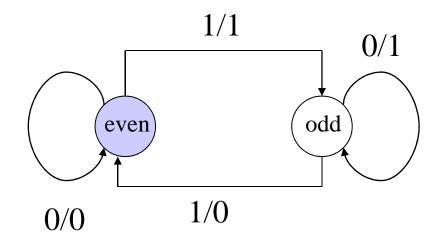
final state = 0 (the sequence has an even number of components equal to 1)

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final state = 1 (the sequence has an odd number of components equal to 1)
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State diagram = labeled directed graph

EP Design:

- choose: S, I, O



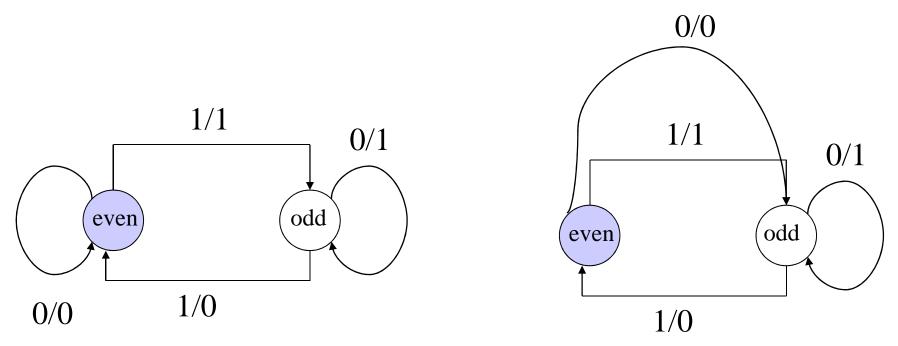
Population initialization: generate random FSMs

- Generate labels for nodes
- Generate arcs
- Generate labels

Mutation:

- Mutation of the output symbol
- Redirect an arc (mutate the target node)
- Add/eliminate nodes
- Change the initial state

Mutation example: change the target node of an arc



Encoding particularities: the population contains labeled graphs Types of representation: explicit / implicit

Variants of explicit representation:

- Fixed number of nodes + fixed number of edges => only the edge weights are adaptable
 - Array of fixed length containing the weight values = linearized version of the weight matrix
- Fixed number of nodes + variable number of edges
 - Array of weight values (there is used one special value corresponding to removed edges)
- Variable number of nodes => both the presence indicators of nodes as well as the edge weights can be changed
 - Binary array containing the presence indicators of the nodes
 - List of the edge weights

Mutation:

- Change of the edge weights (by replacing with randomly selected values – for discrete weights, by random additive perturbation – for real weights)
- Complementing some randomly selected values from the vector containing the presence indicators for nodes => add/remove nodes

Remark: each mutation type is applied with a given probability, e.g.:

- p1: edge weight changing
- p2: edge addition
- p3: edge removal
- p4: node addition
- p5: node removal

(the probabilities sum to 1: p1+p2+p3+p4+p5=1)

Crossover:

- For fixed structure graphs => linear structure => standard crossover operators can be applied
- For graphs with arbitrary structure:
 - Interchange two subgraphs (exchange the set of edges which correspond to a subset of nodes)

Rmk: implicit representations are characterized by evolving synthetic descriptions of the structures.

Example:

- representation: (number of nodes, connectivity degree, distribution probability for weights)
- evaluation: a particular instance is generation based on the description

Evaluation of a configuration:

- simulation for a test set
- the fitness is considered to be proportional with the success rate

Current status in the field

• it has been redirected to the evolutionary design of computational structures (e.g. neural networks)

Second (current) direction: it is related to optimization methods similar to evolution strategies

- there is only a mutation operator (no recombination)
 - the mutation is based on random perturbation of the current configuration (x'=x+N(0,s))
 - s is inversely correlated with the fitness value (high fitness leads to small s, low fitness leads to large values for s)
- starting from a population with m elements, by mutation are constructed m children and the survivors are selected from the 2m elements by tournament or by truncation.
- There are self-adaptive variants, called MetaEP; these variants are similar to self-adaptive Evolution Strategies

MetaEP

$$(x_1, ..., x_n, s_1, ..., s_n) \rightarrow (x'_1, ..., x'_n, s'_1, ..., s'_n)$$

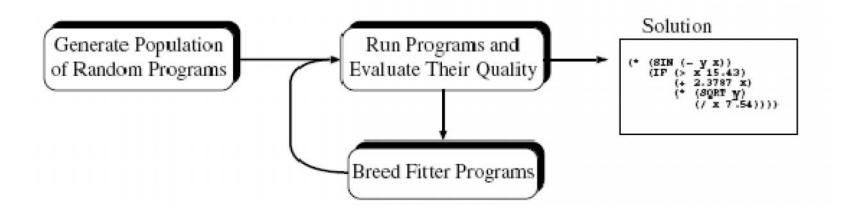
 $s'_i = s_i (1 + \Gamma N(0.1)), \ \Gamma \cong 0.2$
 $x'_i = x_i + s'_i N(0.1)$

Remark: currently the normal mutation used to self-adapt the control parameters has been replaced with a log-normal distribution (as in the case of SE)

Principal contributor: J. Koza (1990)

Official web site: www.genetic-programming.org

- GP is an automated method for creating a working computer program from a high-level statement of a problem.
- GP starts from a high-level statement of "what needs to be done" and automatically creates a computer program to solve the problem.



The result is a program or an "executable" expression

Simplest example: symbolic regression

Numeric regression

Symbolic regression

Input data:

- pairs of values: (arg, val)
- model which depends on some parameters(e.g.: linear model, quadratic model etc)
- Output: values of the model parameters

Input data:

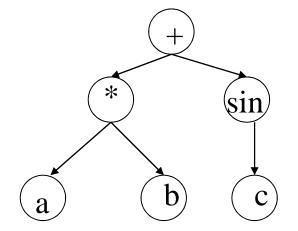
- pairs of values : (arg, val)
- alphabets of terminals (variables, constants) and nonterminals (operators, functions)
- Output: expression which describes the dependence between the output variable (predicted value) and the input variable (predictor)

Numerical regression	Symbolic regression
Input data: (1,3),(2,5),(3,7),(4,9)	Input data: (1,3),(2,5),(3,7),(4,9)
Model: f(x)=ax+b	Alphabet: +,*,-,/,constants,x
Result: a=2 b=1	Result: 2*x+1 or x+x+1 or any other equivalent expression
Search in the parameter space	Search in the space of expressions
	http://alphard.ethz.ch/gerber/approx/default.html

Encoding: the individuals are usually tree-like structures

Example 1: arithmetical expression a*b+sin(c)

Components:

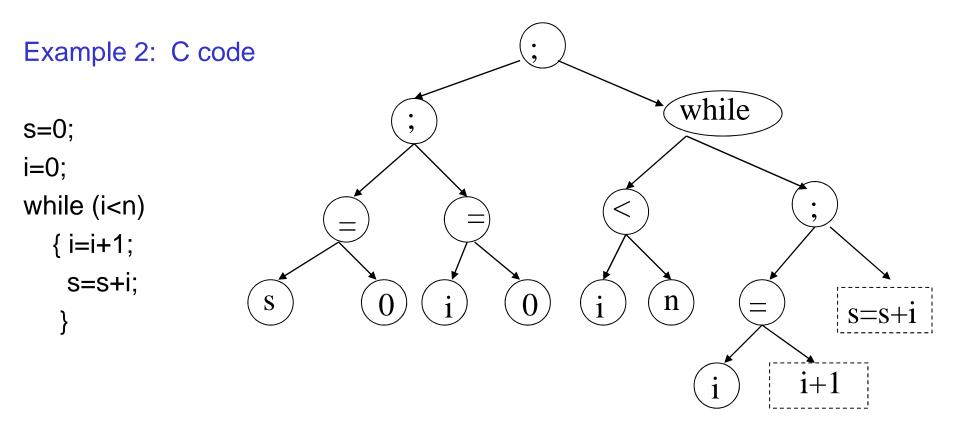


Nonterminals: operators and functions

Terminals: variables, constants (fixed or randomly generated – called ephemeral random constants), 0-arity functions

Prefixed form: +*a b sin c (preorder) Postfixed form: a b * c sin + (postorder)

Encoding: the individuals are usually tree-like structures



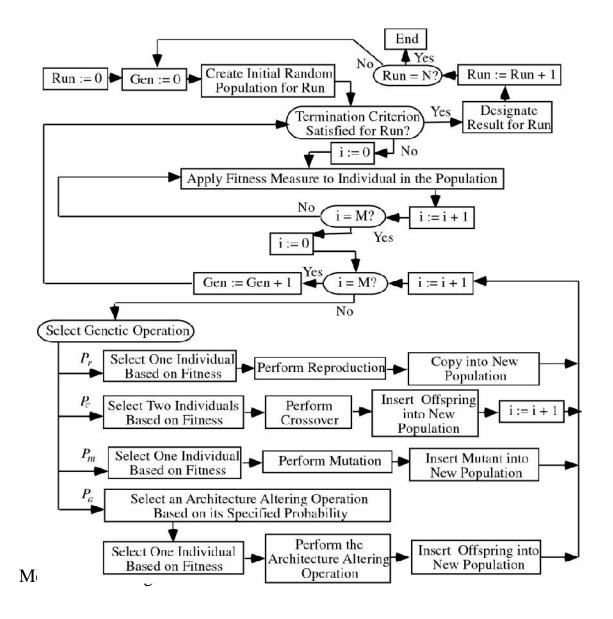
Problem: the tree representation can be complex even for simple programs

Summary: the terminals and nonterminals sets are chosen depending on the problem to be solved

Function Set Kind of Primitive Example(s)		Terminal Set Kind of Primitive Example(s)	
:	-		

Overall structure of a GP algorithm [Koza, 2003]

Remark: The evolutionary operators (selection, crossover, mutation) are applied alternatively, i.e. either crossover or mutation is applied



Implementation:

- classical variant: LISP
- lists corresponding to prefixed description of expressions

Difficulty: all elements should be syntactically correct

Generation function - parameters

T: terminals N: nonterminals A: tree depth Generate(T,N,A) IF A=0 THEN expr:=choose(T) **FLSE** fct:=choose(N) IF (unary(fct)) THEN arg:=generate(T,N,A-1) expr:=(fct,arg) IF (binary(fct)) THEN arg1:=generate(T,N,A-1) arg2:=generate(T,N,A-1) expr:=(fct,arg1,arg2) **RETURN** expr

Other types of population elements:

- Decision trees
- If-then rules
- Neural networks
- Logical expressions
- Binary decision diagrams
- Grammars \rightarrow Grammatical Evolution

Fitness computation:

- the expression (phenotype) corresponding to each chromosome (genotype) is evaluated for a test data set

- the fitness of a chromosome is higher if the value obtained by evaluating the expression is close to the desired value

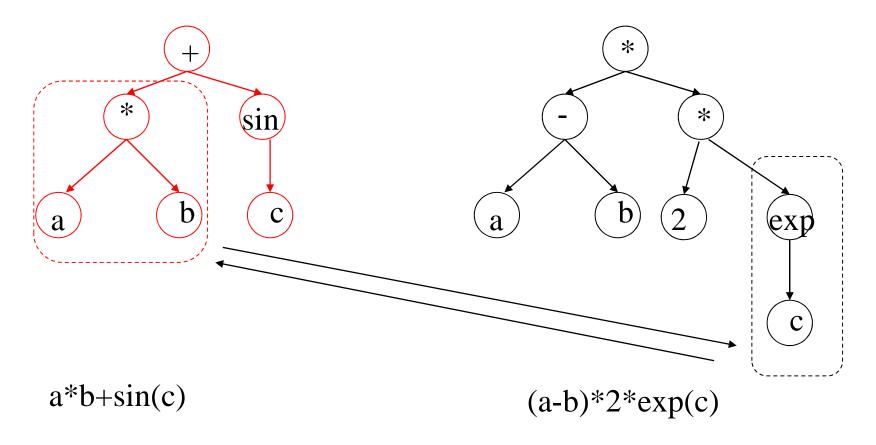
Evaluation:

Algorithm 3 Typical interpreter for GP

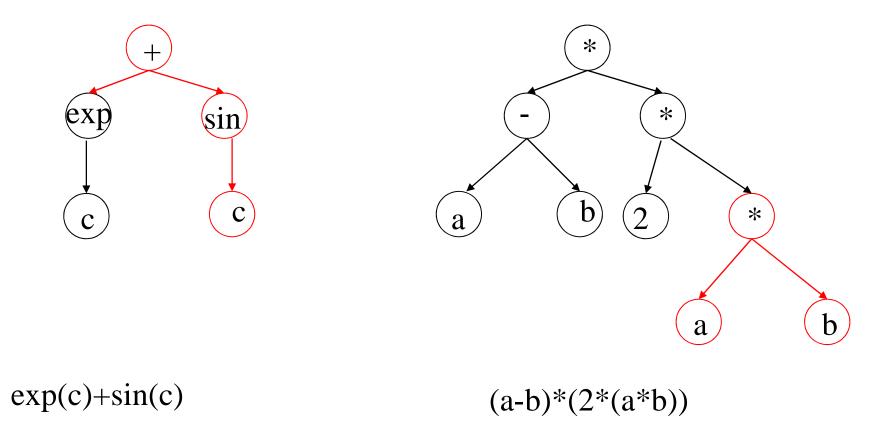
procedure: eval(expr)

```
1: if expr is a list then
       proc = expr(1) {Non-terminal: extract root}
2:
 3:
       if proc is a function then
          value = \operatorname{proc}(\operatorname{eval}(\exp(2)), \operatorname{eval}(\exp(3)), \dots) {Function: evaluate
 4:
          arguments}
 5:
       else
          value = \operatorname{proc}(\operatorname{expr}(2), \operatorname{expr}(3), \ldots) {Macro: don't evaluate arguments}
6:
 7: else
8:
       if expr is a variable or expr is a constant then
          value = expr {Terminal variable or constant: just read the value}
9:
10:
       else
          value = expr() {Terminal 0-arity function: execute}
11:
12: return value
----
```

Crossover: two parents (trees) generate two offspring (also trees) by swapping some subtrees



Crossover: two parents (trees) generate two offspring (also trees) by swapping some subtrees



Crossover:

Prefixed forms of parents and children

+ * a b sin c	* - a b * 2 <mark>exp c</mark>
+ exp c sin c	* - a b * 2 * <mark>a b</mark>

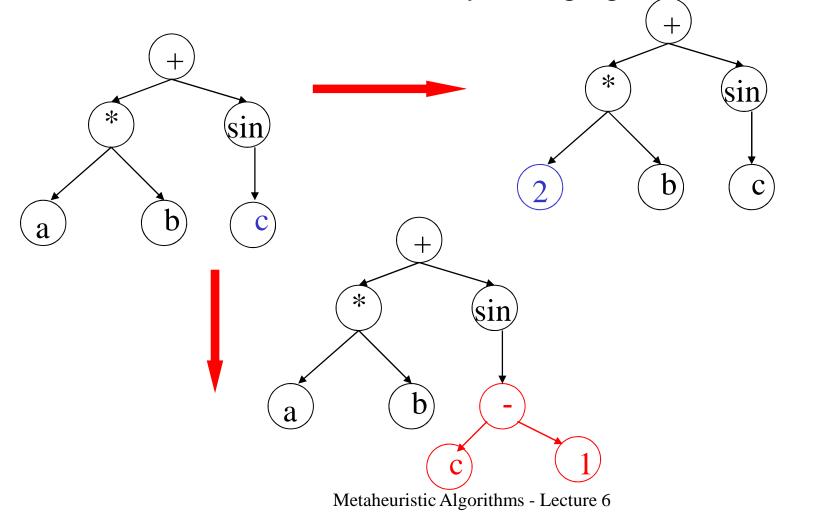
Remark. It is similar to the crossover used at GAs but the size for exchanged portions are usually different.

Mutation: consists of randomly changing some elements

- Change the symbol of a leaf node with another terminal symbol (in the case of constants this mutation could be as in the case of evolution strategies)
- Replace a leaf node with a tree (growing mutation)
- Replace the symbol corresponding to an internal node with another nonterminal from the same class (function with the same arity)
- Replace a subtree with a terminal node (pruning mutation)

Remark: the mutation can be implemented by a crossover with a randomly generated element

Mutation: consists of randomly changing some elements



Bloat problem: the complex structures become dominant in the population

Solutions:

- Use a threshold for the structure complexity (e.g. tree depth) and reject all structures larger (deeper) than the threshold
- Use a penalty term depending on the structure complexity in the fitness computation; this term will penalize the complex structures

GP related approaches:

- Linear Genetic Programming
- Gene Expression Programming [http://www.gene-expressionprogramming.com/]
- Cartesian Genetic Programming [http://www.cartesiangp.co.uk/]
- Multi-expression Programming [http://www.mep.cs.ubbcluj.ro/]
- Grammatical Evolution [http://www.grammatical-evolution.org/]

Linear Genetic Programming [Brameier, Banzhaf, 2003]

```
void gp(r)
  double r[8];
£
   . . .
   r[0] = r[5] + 71;
// r[7] = r[0] - 59;
   if (r[1] > 0)
   if (r[5] > 2)
     r[4] = r[2] * r[1];
// r[2] = r[5] + r[4];
   r[6] = r[4] * 13;
   r[1] = r[3] / 2;
// if (r[0] > r[1])
// r[3] = r[5] * r[5];
   r[7] = r[6] - 2;
// r[5] = r[7] + 15;
   if (r[1] <= r[6])
     r[0] = sin(r[7]);
}
```

Particularities:

- Used to generate programs as sequences of lines (e.g. like in assembling languages)
- The operations involves registers
- Instructions: if and goto
- The commented lines correspond to processing steps which do not influence the final result (similar to noncoding portions of DNA – the so-called introns)
- Crossover: uses a variant of single point crossover adapted for chromosomes with different lengths (the program is a chromosome, each line is a gene)

GEP - Gene Expression Programming (C. Ferreira, 2001):

+*

sin

Chromosome:

- Consists of several genes of fixed length
- Each gene has a head and a tail
 - The head contains h symbols (both terminals and nonterminals); the tail contains only terminals; the number of elements in the tail is h*(n-1)+1, n=the maximal arity of functions/operators which appears in the head

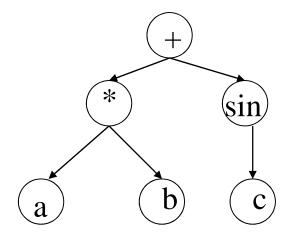
Example: gene of length $13 = 6 + (6^{*}(2-1)+1) = h + (h^{*}(n-1)+1)$

+ * sin a b c b a c c b a a

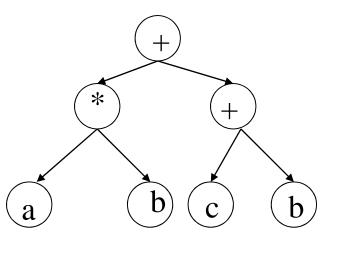
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- The first 6 elements correspond with the expression (breadth first search of the tree)
- All other elements are terminals (unused in the genotype-phenotype conversion) Metaheuristic Algorithms - Lecture 6 33

GEP: allow to generate syntactically correct expressions by extending the head over the symbols in the tail



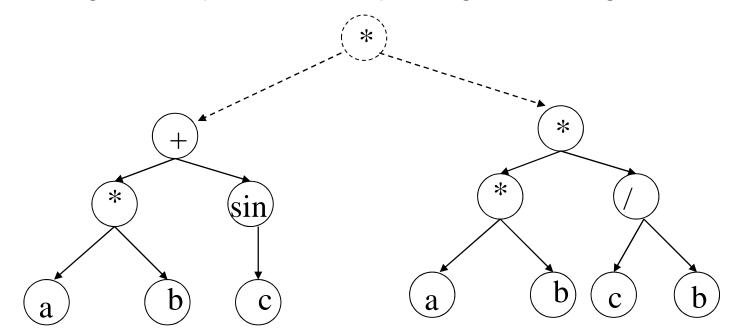
+ * sin a b c b a c c b a a



+ * + a b c b a c c b a a

GEP: chromosome consisting of two genes: + * sin a b c b a c c b a a * * / a b c b a c c b a a

The phenotype corresponding to the chromosome is obtained by combining the components corresponding to the two genes



Applications:

- Extracting models from data (e.g. predictive models)
- Extracting rules from data
- Electrical circuits design
- Robust systems synthesis
- Evolvable hardware

- parallel applications design
- cellular automata design
- signal/image processing filters design
- generation of multi-agent strategies
- generation of game strategies
- generation of quantum algorithms

See http://www.human-competitive.org/awards for GA, ES and GP applications in generating solutions competitive with those obtained by human experts

Genetic Programming Software:

- Java: ECJ, TinyGP,
- Matlab: GPLab, GPTips
- C/C++: MicroGP
- Python: DEAP, PyEvolve