

# Genetic Programming

## Lecture 6

- ▶ Developed: USA in the 1990's
- ▶ Early names: J. Koza
- ▶ Typically applied to:
  - machine learning tasks (prediction, classification...)
- ▶ Attributed features:
  - competes with neural nets and alike
  - needs huge populations (thousands)
  - slow
- ▶ Special:
  - non-linear chromosomes: trees, graphs
  - mutation possible but not necessary (disputed!)

# GP technical summary tableau

Representation	Tree structures
Recombination	Exchange of subtrees
Mutation	Random change in trees
Parent selection	Fitness proportional
Survivor selection	Generational replacement

# Introductory example: credit scoring

- ▶ Bank wants to distinguish good from bad loan applicants
- ▶ Model needed that matches historical data

ID	No of children	Salary	Marital status	OK?
ID-1	2	45000	Married	0
ID-2	0	30000	Single	1
ID-3	1	40000	Divorced	1
...				

# Introductory example: credit scoring

- ▶ A possible model:

IF (NOC = 2) AND (S > 80000) THEN good ELSE bad

- ▶ In general:

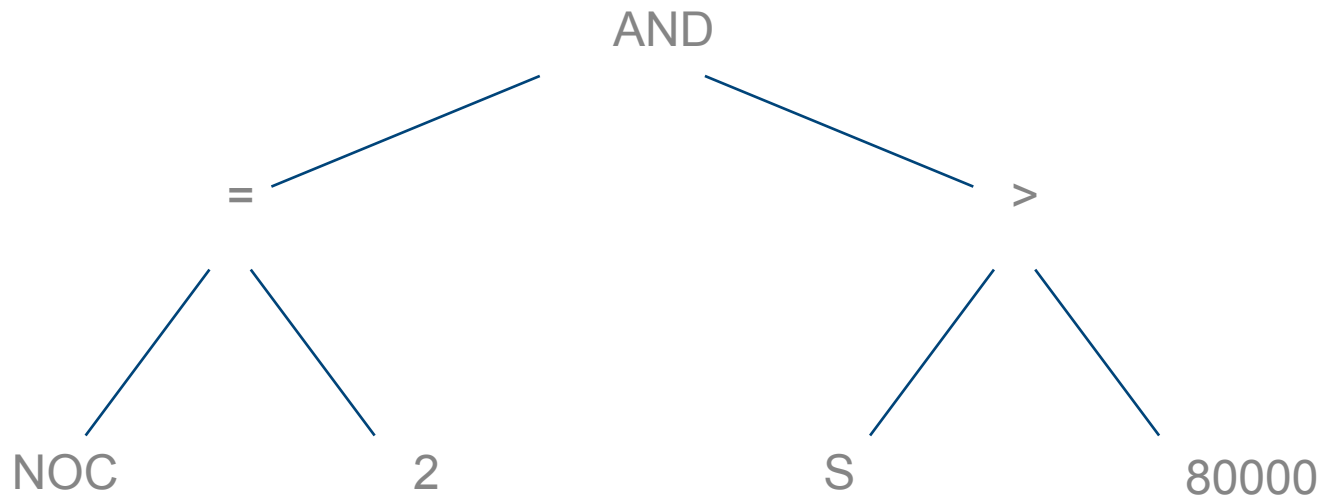
IF formula THEN good ELSE bad

- ▶ Only unknown is the right formula, hence
- ▶ Our search space (phenotypes) is the set of formulas
- ▶ Natural fitness of a formula: percentage of well classified cases of the model it stands for
- ▶ Natural representation of formulas (genotypes) is: parse trees

# Introductory example: credit scoring

IF (NOC = 2) AND (S > 80000) THEN good ELSE bad

can be represented by the following tree



# Tree based representation

- ▶ Trees are a universal form, e.g. consider

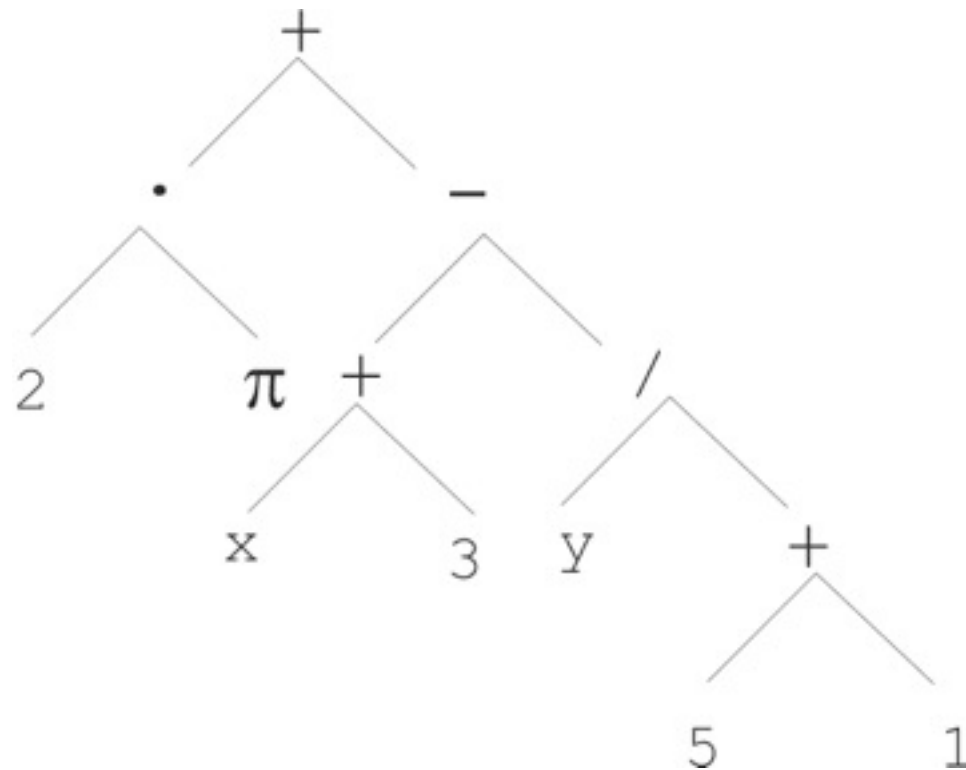
- Arithmetic formula  $2 \cdot \pi + \left( (x + 3) - \frac{y}{5 + 1} \right)$

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

- Program

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

# Tree based representation

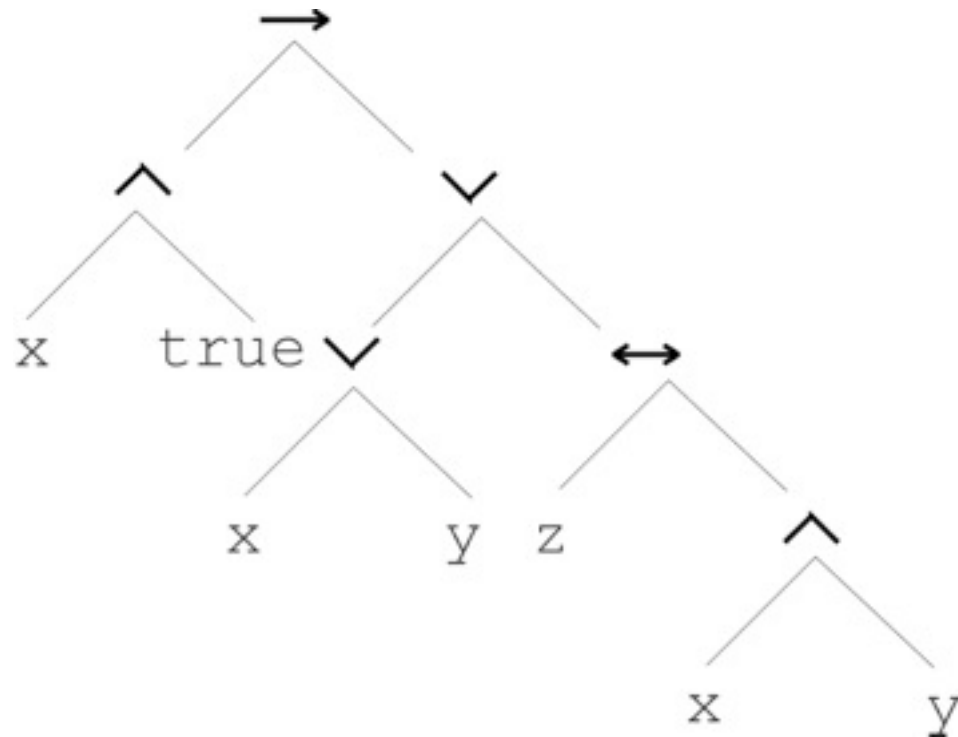


$$2 \cdot \pi + \left( (x + 3) - \frac{y}{5 + 1} \right)$$

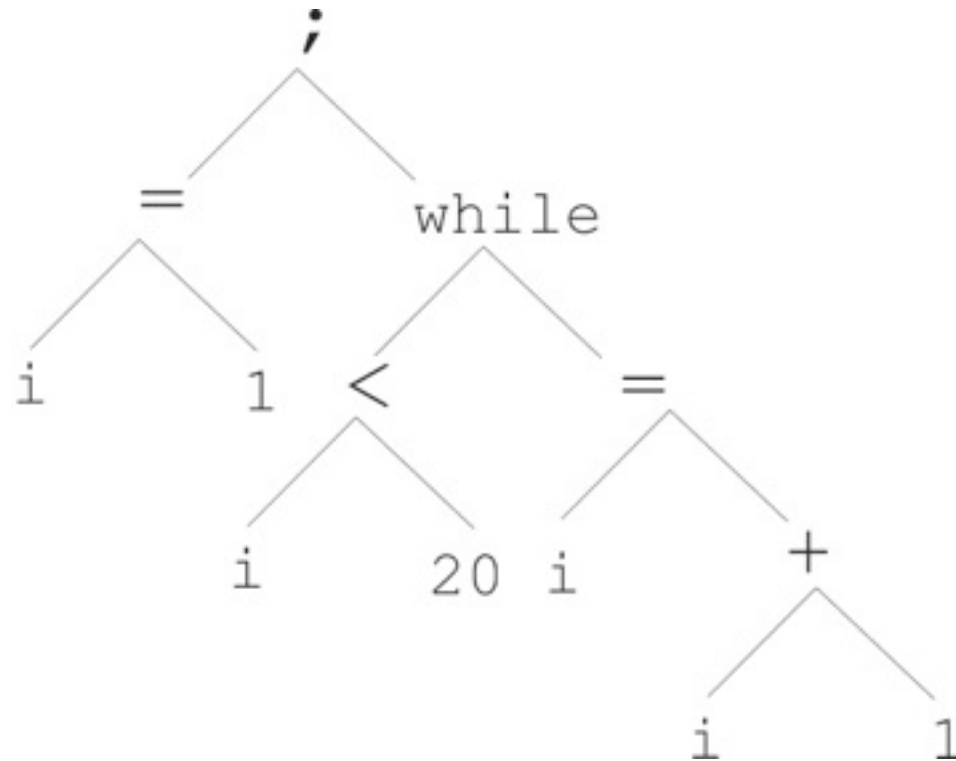


# Tree based representation

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



# Tree based representation



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

# Tree based representation

- ▶ In GA, ES, EP chromosomes are linear structures (bit strings, integer string, real-valued vectors, permutations)
- ▶ Tree shaped chromosomes are non-linear structures
- ▶ In GA, ES, EP the size of the chromosomes is fixed
- ▶ Trees in GP may vary in depth and width

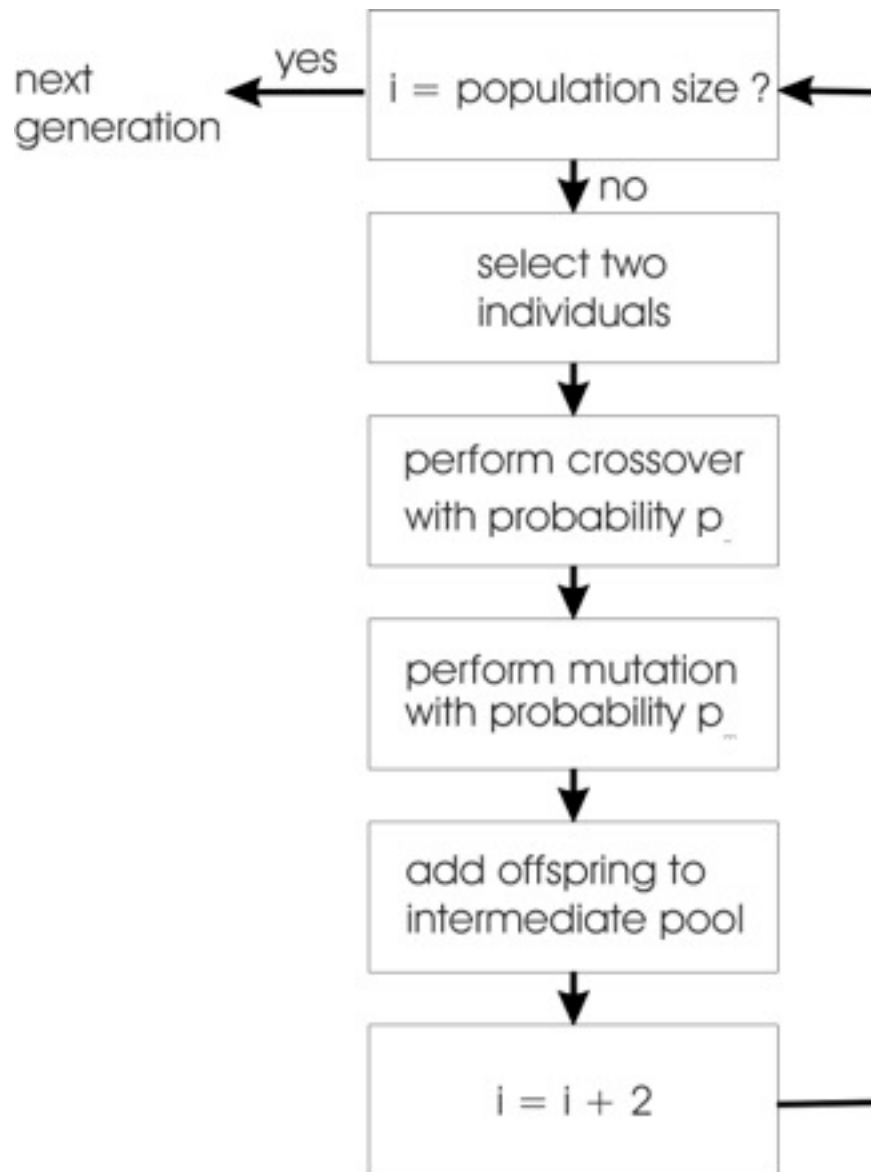
# Tree based representation

- Symbolic expressions can be defined by
  - Terminal set  $T$
  - Function set  $F$  (with the arities of function symbols)
- Adopting the following general recursive definition:
  1. Every  $t \in T$  is a correct expression
  2.  $f(e_1, \dots, e_n)$  is a correct expression if  $f \in F$ ,  $\text{arity}(f)=n$  and  $e_1, \dots, e_n$  are correct expressions
  3. There are no other forms of correct expressions
- In general, expressions in GP are not typed (closure property: any  $f \in F$  can take any  $g \in F$  as argument)

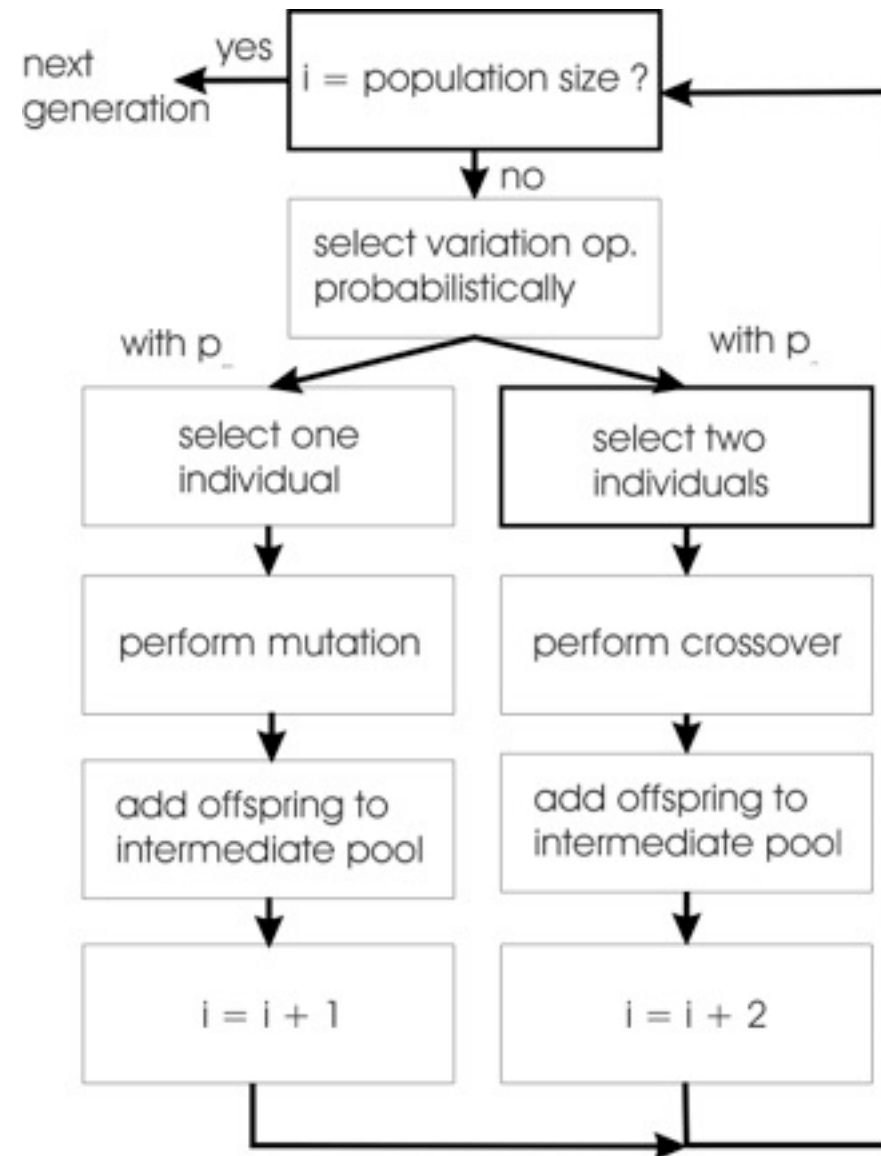
# Offspring creation scheme

Compare

- ▶ GA scheme using crossover AND mutation sequentially (be it probabilistically)
- ▶ GP scheme using crossover OR mutation (chosen probabilistically)



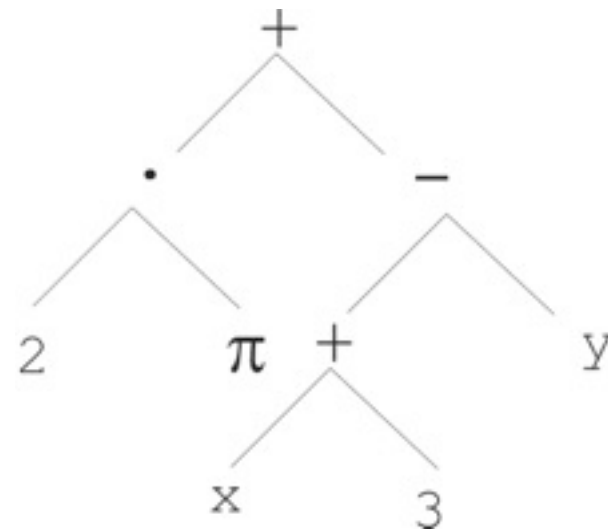
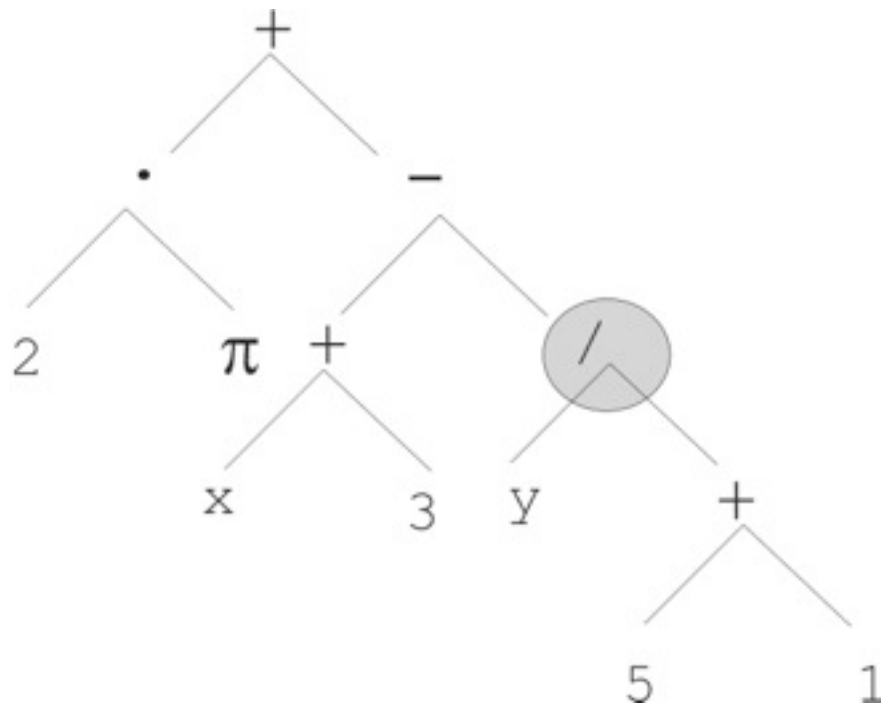
GA flowchart



GP flowchart

# Mutation

- ▶ Most common mutation: replace randomly chosen subtree by randomly generated tree



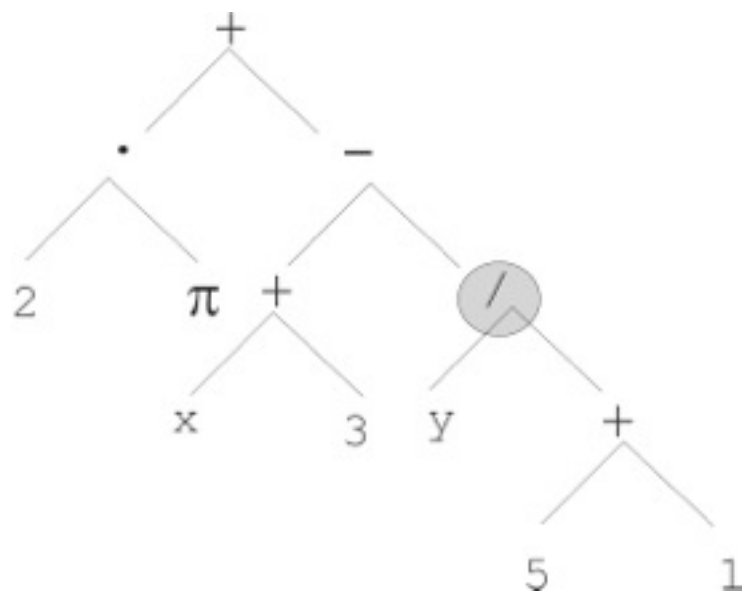
# Mutation cont'd

- ▶ Mutation has two parameters:
  - Probability  $p_m$  to choose mutation vs. recombination
  - Probability to choose an internal point as the root of the subtree to be replaced
- ▶ Remarkably  $p_m$  is advised to be 0 (Koza'92) or very small, like 0.05 (Banzhaf et al. '98)
- ▶ The size of the child can exceed the size of the parent

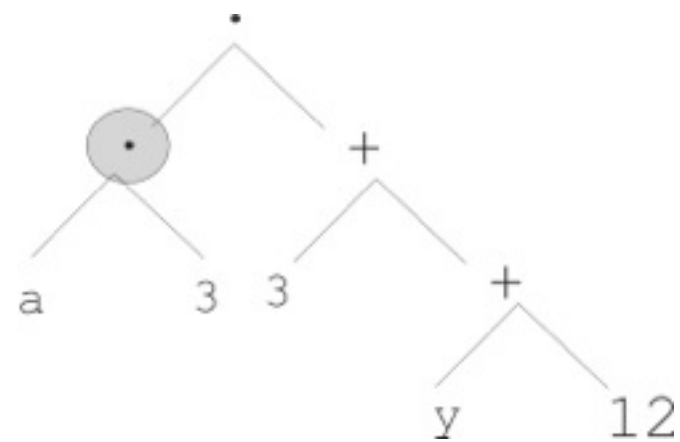


# Recombination

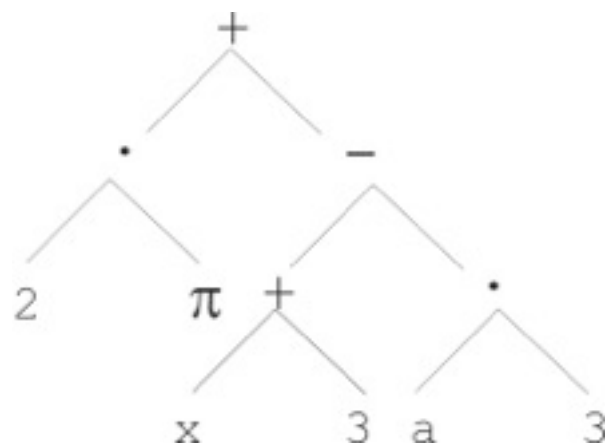
- ▶ Most common recombination: exchange two randomly chosen subtrees among the parents
- ▶ Recombination has two parameters:
  - Probability  $p_c$  to choose recombination vs. mutation
  - Probability to choose an internal point within each parent as crossover point
- ▶ The size of offspring can exceed that of the parents



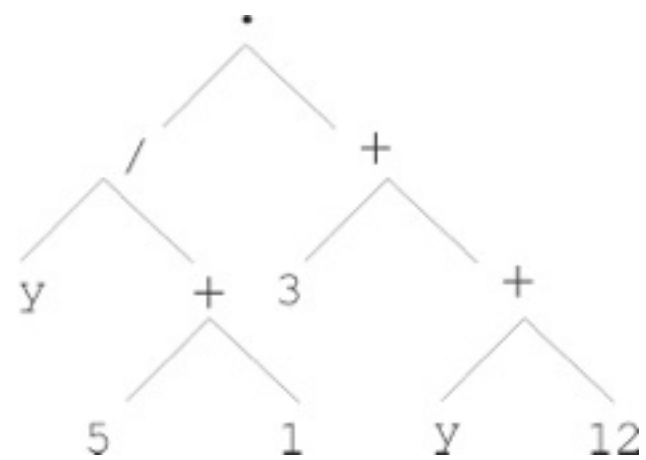
Parent 1



Parent 2



Child 1



Child 2

# Selection

- ▶ Parent selection typically fitness proportionate
- ▶ Over-selection in very large populations
  - rank population by fitness and divide it into two groups:
  - group 1: best  $x\%$  of population, group 2 other  $(100-x)\%$
  - 80% of selection operations chooses from group 1, 20% from group 2
  - for pop. size = 1000, 2000, 4000, 8000  $x = 32\%, 16\%, 8\%, 4\%$
  - motivation: to increase efficiency, %'s come from rule of thumb
- ▶ Survivor selection:
  - Typical: generational scheme (thus none)
  - Recently steady-state is becoming popular for its elitism

# Initialisation

- ▶ Maximum initial depth of trees  $D_{\max}$  is set
- ▶ Full method (each branch has depth =  $D_{\max}$ ):
  - nodes at depth  $d < D_{\max}$  randomly chosen from function set  $F$
  - nodes at depth  $d = D_{\max}$  randomly chosen from terminal set  $T$
- ▶ Grow method (each branch has depth  $\leq D_{\max}$ ):
  - nodes at depth  $d < D_{\max}$  randomly chosen from  $F \cup T$
  - nodes at depth  $d = D_{\max}$  randomly chosen from  $T$
- ▶ Common GP initialisation: ramped half-and-half, where grow & full method each deliver half of initial population

# Bloat

- ▶ Bloat = “survival of the fattest”, i.e., the tree sizes in the population are increasing over time
- ▶ Ongoing research and debate about the reasons
- ▶ Needs countermeasures, e.g.
  - Prohibiting variation operators that would deliver “too big” children
  - Parsimony pressure: penalty for being oversized

# Problems involving “physical” environments

- ▶ Trees for data fitting vs. trees (programs) that are “really” executable
- ▶ Execution can change the environment → the calculation of fitness
- ▶ Example: robot controller
- ▶ Fitness calculations mostly by simulation, ranging from expensive to extremely expensive (in time)
- ▶ But evolved controllers are often to very good

# Example application:

- ▶ Given some points in  $\mathbf{R}^2$ ,  $(x_1, y_1), \dots, (x_n, y_n)$
- ▶ Find function  $f(x)$  s.t.  $\forall i = 1, \dots, n : f(x_i) = y_i$
- ▶ Possible GP solution:
  - Representation by  $F = \{+, -, /, \sin, \cos\}$ ,  $T = \mathbf{R} \cup \{x\}$
  - Fitness is the error  $err(f) = \sum_{i=1}^n (f(x_i) - y_i)^2$
  - All operators standard
  - pop.size = 1000, ramped half-half initialisation
  - Termination: n “hits” or 50000 fitness evaluations reached (where “hit” is if  $|f(x_i) - y_i| < 0.0001$ )

# Discussion

Is GP:

The art of evolving computer programs ?

Means to automated programming of computers?

GA with another representation?