

# Genetic Algorithms

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[Read Chapter 9]  
[Exercises 9.1, 9.2, 9.3, 9.4]

- Evolutionary computation
- Prototypical GA
- An example: GABIL
- Genetic Programming
- Individual learning and population evolution

# Evolutionary Computation

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1. Computational procedures patterned after biological evolution
2. Search procedure that probabilistically applies search operators to set of points in the search space

# Biological Evolution

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Lamarck and others:

- Species “transmute” over time

Darwin and Wallace:

- Consistent, heritable variation among individuals in population
- Natural selection of the fittest

Mendel and genetics:

- A mechanism for inheriting traits
- genotype  $\rightarrow$  phenotype mapping

GA(*Fitness*, *Fitness\_threshold*, *p*, *r*, *m*)

- *Initialize*:  $P \leftarrow p$  random hypotheses
- *Evaluate*: for each  $h$  in  $P$ , compute  $Fitness(h)$
- While  $[\max_h Fitness(h)] < Fitness\_threshold$ 
  1. *Select*: Probabilistically select  $(1 - r)p$  members of  $P$  to add to  $P_s$ .

$$\Pr(h_i) = \frac{Fitness(h_i)}{\sum_{j=1}^p Fitness(h_j)}$$

2. *Crossover*: Probabilistically select  $\frac{r \cdot p}{2}$  pairs of hypotheses from  $P$ . For each pair,  $\langle h_1, h_2 \rangle$ , produce two offspring by applying the Crossover operator. Add all offspring to  $P_s$ .
  3. *Mutate*: Invert a randomly selected bit in  $m \cdot p$  random members of  $P_s$
  4. *Update*:  $P \leftarrow P_s$
  5. *Evaluate*: for each  $h$  in  $P$ , compute  $Fitness(h)$
- Return the hypothesis from  $P$  that has the highest fitness.

# Representing Hypotheses

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Represent

$$(Outlook = Overcast \vee Rain) \wedge (Wind = Strong)$$

by

$$\begin{array}{cc} Outlook & Wind \\ 011 & 10 \end{array}$$

Represent

$$\text{IF } Wind = Strong \text{ THEN } PlayTennis = yes$$

by

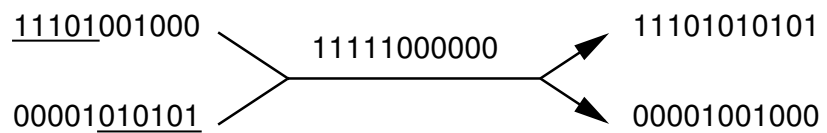
$$\begin{array}{ccc} Outlook & Wind & PlayTennis \\ 111 & 10 & 10 \end{array}$$

# Operators for Genetic Algorithms

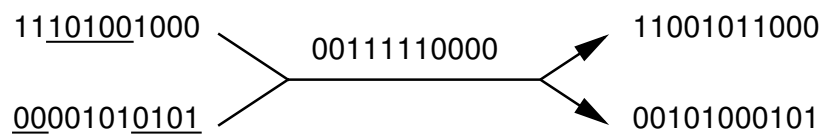
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<i>Initial strings</i>	<i>Crossover Mask</i>	<i>Offspring</i>
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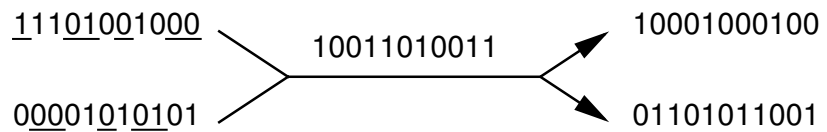
*Single-point crossover:*



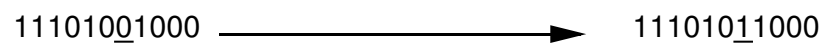
*Two-point crossover:*



*Uniform crossover:*



*Point mutation:*



# Selecting Most Fit Hypotheses

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Fitness proportionate selection:

$$\Pr(h_i) = \frac{Fitness(h_i)}{\sum_{j=1}^p Fitness(h_j)}$$

... can lead to *crowding*

Tournament selection:

- Pick  $h_1, h_2$  at random with uniform prob.
- With probability  $p$ , select the more fit.

Rank selection:

- Sort all hypotheses by fitness
- Prob of selection is proportional to rank

# GABIL [DeJong et al. 1993]

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Learn disjunctive set of propositional rules,  
competitive with C4.5

**Fitness:**

$$Fitness(h) = (correct(h))^2$$

**Representation:**

IF  $a_1 = T \wedge a_2 = F$  THEN  $c = T$ ; IF  $a_2 = T$  THEN  $c = F$   
represented by

$a_1$	$a_2$	$c$	$a_1$	$a_2$	$c$
10	01	1	11	10	0

**Genetic operators: ???**

- want variable length rule sets
- want only well-formed bitstring hypotheses



# Crossover with Variable-Length Bitstrings

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Start with

$$h_1 : \begin{array}{ccc} a_1 & a_2 & c \\ 10 & 01 & 1 \end{array} \quad \begin{array}{ccc} a_1 & a_2 & c \\ 11 & 10 & 0 \end{array}$$

$$h_2 : \begin{array}{ccc} 01 & 11 & 0 \end{array} \quad \begin{array}{ccc} 10 & 01 & 0 \end{array}$$

1. choose crossover points for  $h_1$ , e.g., after bits 1, 8
2. now restrict points in  $h_2$  to those that produce bitstrings with well-defined semantics, e.g.,  $\langle 1, 3 \rangle$ ,  $\langle 1, 8 \rangle$ ,  $\langle 6, 8 \rangle$ .

if we choose  $\langle 1, 3 \rangle$ , result is

$$h_3 : \begin{array}{ccc} a_1 & a_2 & c \\ 11 & 10 & 0 \end{array}$$

$$h_4 : \begin{array}{ccc} a_1 & a_2 & c \\ 00 & 01 & 1 \end{array} \quad \begin{array}{ccc} a_1 & a_2 & c \\ 11 & 11 & 0 \end{array} \quad \begin{array}{ccc} a_1 & a_2 & c \\ 10 & 01 & 0 \end{array}$$

# GABIL Extensions

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Add new genetic operators, also applied probabilistically:

1. *AddAlternative*: generalize constraint on  $a_i$  by changing a 0 to 1
2. *DropCondition*: generalize constraint on  $a_i$  by changing every 0 to 1

And, add new field to bitstring to determine whether to allow these

$a_1$	$a_2$	$c$	$a_1$	$a_2$	$c$	$AA$	$DC$
01	11	0	10	01	0	1	0

So now the learning strategy also evolves!

# GABIL Results

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Performance of GABIL comparable to symbolic rule/tree learning methods C4.5, ID5R, AQ14

Average performance on a set of 12 synthetic problems:

- GABIL without *AA* and *DC* operators: 92.1% accuracy
- GABIL with *AA* and *DC* operators: 95.2% accuracy
- symbolic learning methods ranged from 91.2 to 96.6

# Schemas

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How to characterize evolution of population in GA?

Schema = string containing 0, 1, \* (“don’t care”)

- Typical schema:  $10^{**}0^{*}$
- Instances of above schema: 101101, 100000, ...

Characterize population by number of instances representing each possible schema

- $m(s, t)$  = number of instances of schema  $s$  in pop at time  $t$

## Consider Just Selection

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- $\bar{f}(t)$  = average fitness of pop. at time  $t$
- $m(s, t)$  = instances of schema  $s$  in pop at time  $t$
- $\hat{u}(s, t)$  = ave. fitness of instances of  $s$  at time  $t$

Probability of selecting  $h$  in one selection step

$$\begin{aligned}\Pr(h) &= \frac{f(h)}{\sum_{i=1}^n f(h_i)} \\ &= \frac{f(h)}{n\bar{f}(t)}\end{aligned}$$

Probability of selecting an instance of  $s$  in one step

$$\begin{aligned}\Pr(h \in s) &= \sum_{h \in s \cap p_t} \frac{f(h)}{n\bar{f}(t)} \\ &= \frac{\hat{u}(s, t)}{n\bar{f}(t)} m(s, t)\end{aligned}$$

Expected number of instances of  $s$  after  $n$  selections

$$E[m(s, t + 1)] = \frac{\hat{u}(s, t)}{\bar{f}(t)} m(s, t)$$

# Schema Theorem

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$$E[m(s, t+1)] \geq \frac{\hat{u}(s, t)}{\bar{f}(t)} m(s, t) \left(1 - p_c \frac{d(s)}{l-1}\right) (1-p_m)^{o(s)}$$

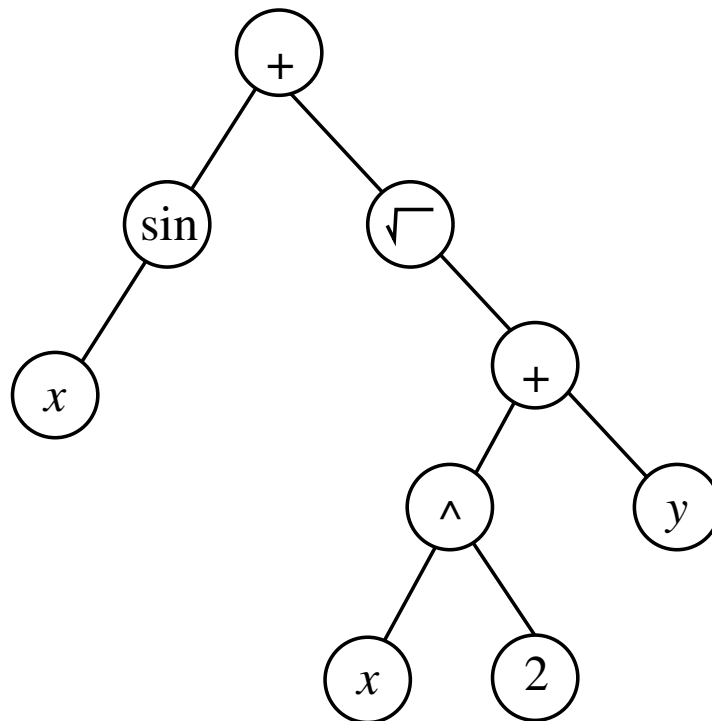
- $m(s, t)$  = instances of schema  $s$  in pop at time  $t$
- $\bar{f}(t)$  = average fitness of pop. at time  $t$
- $\hat{u}(s, t)$  = ave. fitness of instances of  $s$  at time  $t$
- $p_c$  = probability of single point crossover operator
- $p_m$  = probability of mutation operator
- $l$  = length of single bit strings
- $o(s)$  number of defined (non “\*”) bits in  $s$
- $d(s)$  = distance between leftmost, rightmost defined bits in  $s$

# Genetic Programming

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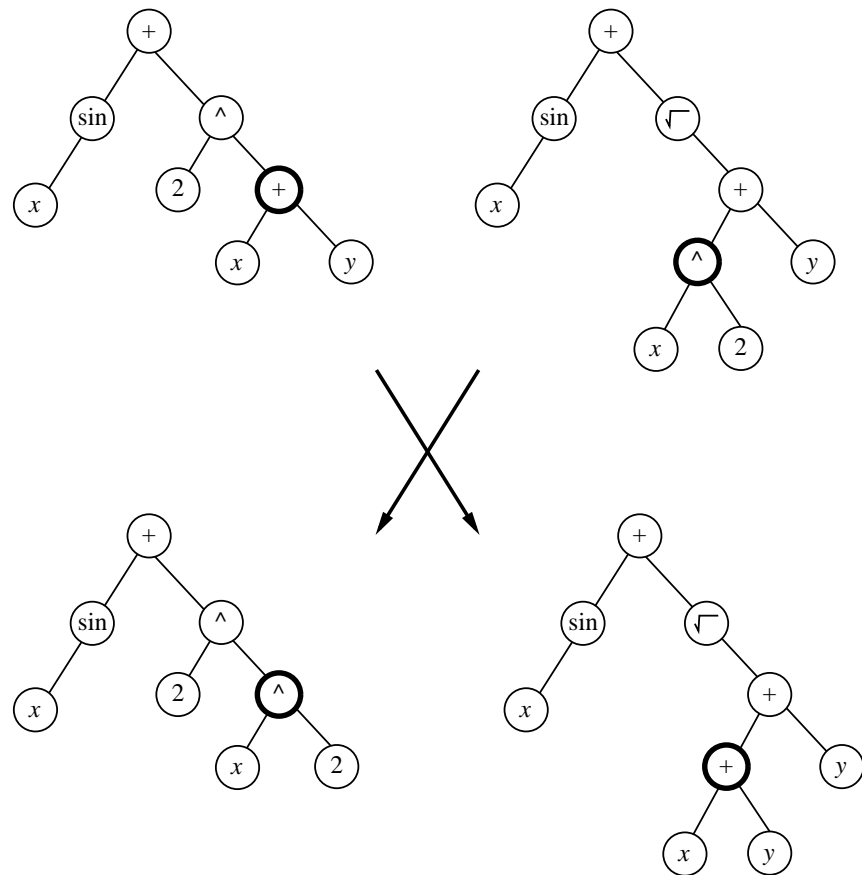
Population of programs represented by trees

$$\sin(x) + \sqrt{x^2 + y}$$



# Crossover

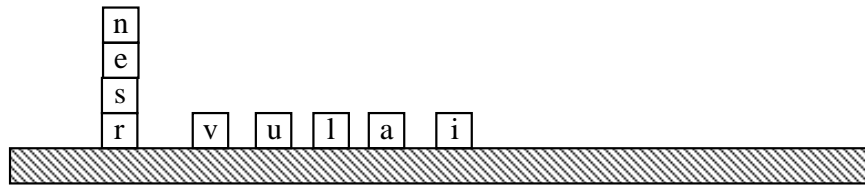
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# Block Problem

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Goal: spell UNIVERSAL

Terminals:

- CS (“current stack”) = name of the top block on stack, or  $F$ .
- TB (“top correct block”) = name of topmost correct block on stack
- NN (“next necessary”) = name of the next block needed above TB in the stack

Primitive functions:

- (MS  $x$ ): (“move to stack”), if block  $x$  is on the table, moves  $x$  to the top of the stack and returns the value  $T$ . Otherwise, does nothing and returns the value  $F$ .
- (MT  $x$ ): (“move to table”), if block  $x$  is somewhere in the stack, moves the block at the top of the stack to the table and returns the value  $T$ . Otherwise, returns  $F$ .
- (EQ  $x\ y$ ): (“equal”), returns  $T$  if  $x$  equals  $y$ , and returns  $F$  otherwise.
- (NOT  $x$ ): returns  $T$  if  $x = F$ , else returns  $F$
- (DU  $x\ y$ ): (“do until”) executes the expression  $x$  repeatedly until expression  $y$  returns the value  $T$

# Learned Program

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Trained to fit 166 test problems

Using population of 300 programs, found this after 10 generations:

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(EQ (DU (MT CS)(NOT CS)) (DU (MS NN)(NOT NN)) )
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# Genetic Programming

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More interesting example: design electronic filter circuits

- Individuals are programs that transform beginning circuit to final circuit, by adding/subtracting components and connections
- Use population of 640,000, run on 64 node parallel processor
- Discovers circuits competitive with best human designs