

Genetic Programming

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| Genetic programming (GP)

GP is a **machine learning** method that can **automatically design**

- **computer programs** (e.g., classifiers, predictors, neural networks, ...), but also
- **other structures** that can be represented in a computer (such as electronic circuits, molecules, antennas, artefacts ...)

without requiring the user to know or specify the form or structure of the solution in advance.

GP belongs to the class of **Evolutionary Algorithms** that are inspired in Darwin's theory of evolution and Neo-Darwinism.



1 Specification

Requirements
Constraints
Training data
Similar solutions
etc.



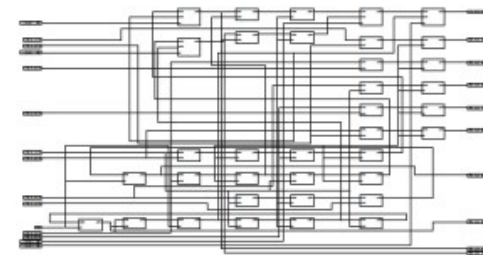
2 Design system



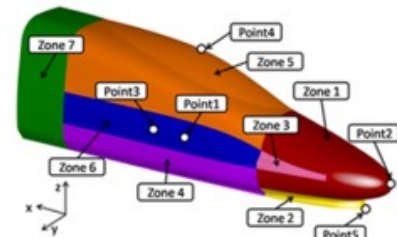
<https://scitechdaily.com/researchers-discover-how-the-human-brain-separates-stores-and-retrieves-memories/>



3 Solution



```
movf    NUM_2,w  
call    SQR  
movf    SQUARE+1,w  
addwf   SUM+1,f  
btfsc   STATUS,C  
    incf    SUM,f  
movf    SQUARE,w  
addwf   SUM,f
```

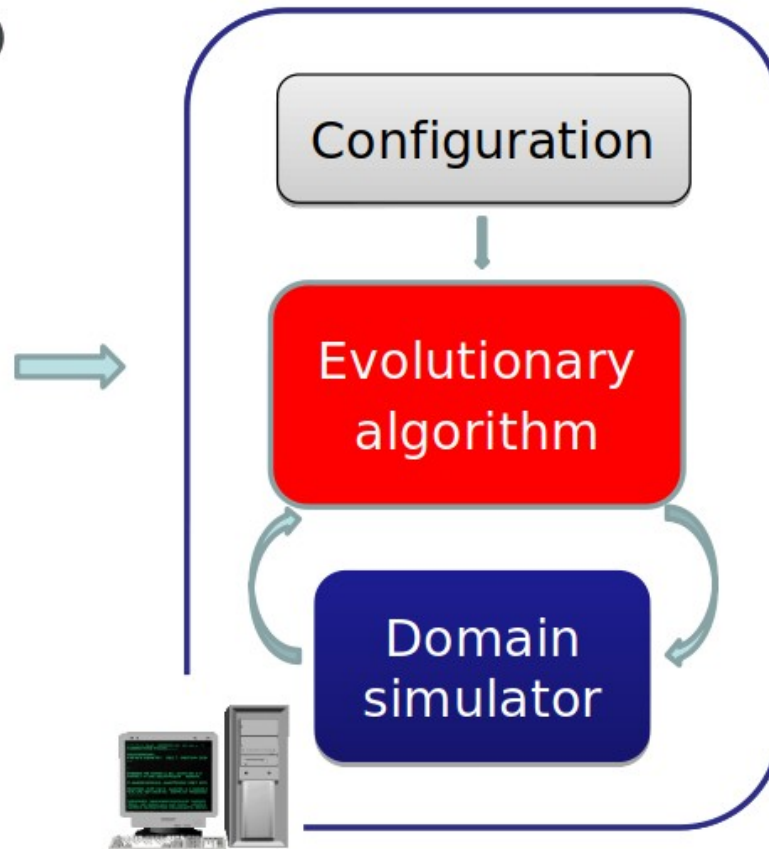


Automated system design

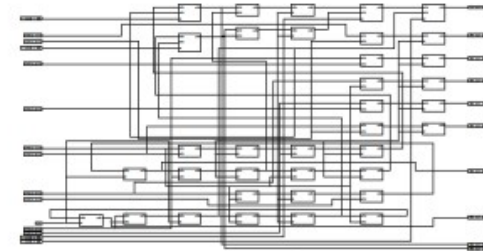
1 Specification

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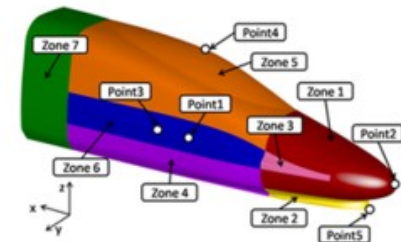
2 Design system



3 Solution



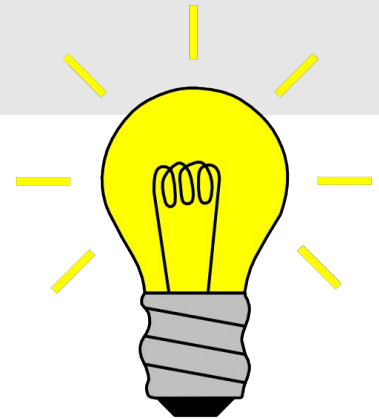
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call    SQR  
movf    SQUARE+1,w  
addwf   SUM+1,f  
btfsc   STATUS,C  
incf    SUM,f  
movf    SQUARE,w  
addwf   SUM,f
```



By Robert Plotkin

The Automation of Invention

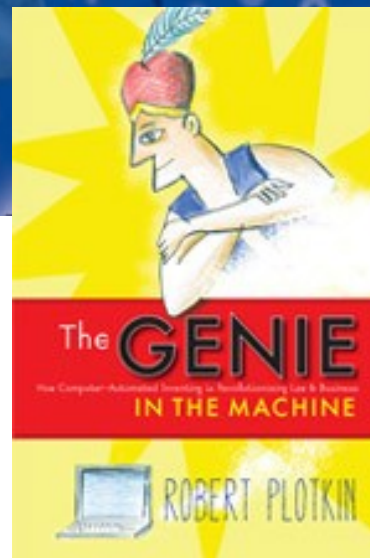
Cybernetic genies are designing and engineering new products and creating technological breakthroughs.



How It Works

Technologies that automate the process of inventing do so in two basic ways: (1) by generating, evaluating, and modifying potential inventions repeatedly until a satisfactory solution is found; and (2) by following a set of rules to design a product or process that achieves a goal.

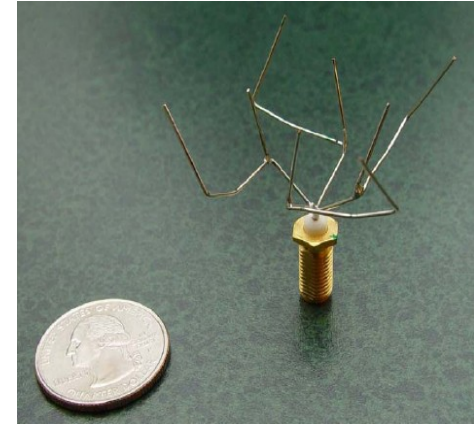
The former technique, which I call “inventing by searching,” is often performed manually by human inventors. Thomas Edison invented an improved light bulb by hiring a team of experts to search the globe for better filament material. After testing more than 6,000 alternatives, he settled on carbonized bamboo.





Benefits

- New (unexpected) and useful designs and solutions can be obtained.
- Human effort is reduced.
- The design can be performed online and autonomously, in target applications such as adaptive, learning and self-repairing systems.



NASA AMES, ST5 Mission



Problems

- Scalability
- Computation time (search method + simulator)
- New ethical issues

- Introduction
- **Evolutionary algorithms**
- Genetic programming
- Cartesian genetic programming
- Applications from FIT
- Summary

Darwinian Evolution

All species of life have descended from common ancestors.

Four Postulates

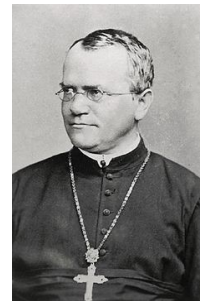
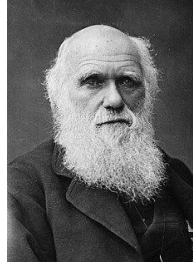
1. Individuals within species are variable
2. Some of the variations are passed on to offspring
3. In every generation, more offspring are produced than can survive
4. The survival and reproduction of individuals are not random: The individuals who survive and go on to reproduce, or who reproduce the most, are those with the most favourable variations. They are **naturally selected**.



On the Origin of Species by Means of Natural Selection (Darwin, 1859)

Neo-Darwinism is generally used to describe the integration of Darwin's theory of evolution by natural selection with Mendel's theory of genetics.

Charles Darwin
(1809 - 1882)



Gregor J. Mendel
(1822 - 1884)

Brno

I Evolutionary algorithms (EA)

- **Evolutionary Algorithms** are inspired in Darwin's theory of evolution and Neo-Darwinism.
- The term **Evolutionary Algorithm (EA)** covers various *search* algorithms that have the following common features:
 - There is a **population** (a set) of individuals (so-called **candidate solutions**).
 - Each candidate solution gets a **fitness value** by evaluating its quality by means of an **objective function** (or **fitness function**). **This function is application specific.**
 - In each iteration (a **generation**) of EA, new candidate solutions are created by
 - **selection** of parent individuals (**those with better fitness are preferred = natural selection**),
 - offspring are created from parents by **crossover** and **mutation** operators (this process is called **reproduction**).
 - The goal is to **optimize** (minimize or maximize) the fitness, i.e. to find a solution which fulfils the requirements of the fitness function as best as possible.
 - EA is a **stochastic algorithm**; nothing is guaranteed :(
- **Observe the analogy with biology:** the offspring may inherit the features of parents (= crossover), sometimes with small random changes (= mutation) → new features may emerge, the offspring **evolve** during many generations.

| When can EA be used?

- If it is possible to **encode** a candidate solution (so-called **phenotype**, e.g., a digital circuits or NN) into a suitable computer representation (so-called **genotype**, e.g. a bit string, a sequence of “real” numbers...) and
- If it is possible to **evaluate** the quality of each candidate solution using a fitness function (e.g. fitness score: 0%...worst – 100%...best).
- EA are typically used to solve **hard optimization** and **design problems** for which common methods do not provide sufficient solutions.
- Main branches of EA:
 - Evolutionary Programming - EP (Fogel ~1962)
 - Evolution Strategies – ES (Rechenberg and Schwefel ~1964)
 - Genetic Algorithms – GA (Holland ~1973)
 - **Genetic Programming** – GP (Cramer ~1985, Schmidhuber ~1987, **Koza, ~1989**)
 - and some others variants currently exist...

Example 1: evolutionary design of a logic circuit

- We must know what we want to design, i.e. the function of the circuit – this is typically specified as a truth table, e.g.:

a	b	c	d	y
0	0	0	0	0
0	0	0	1	1
0	0	1	0	1
0	0	1	1	1
0	1	0	0	1
0	1	0	1	1
0	1	1	0	1
0	1	1	1	1
1	0	0	0	1
1	0	0	1	1
1	0	1	0	1
1	0	1	1	1
1	1	0	0	1
1	1	0	1	1
1	1	1	0	1
1	1	1	1	1

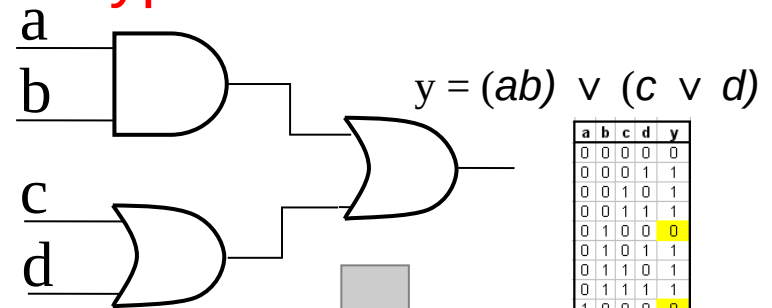
- EA then performs a search in the space of possible candidate solutions using a (small) subset of candidate solutions and bio-inspired operators in order to maximize the fitness score.

Evaluation of a candidate sol.:

genotype (a chromosome)

00110011110010101000011000101

phenotype



a	b	c	d	y
0	0	0	0	0
0	0	0	1	1
0	0	1	0	1
0	0	1	1	1
0	1	0	0	0
0	1	0	1	1
0	1	1	0	1
0	1	1	1	1
1	0	0	0	0
1	0	0	1	1
1	0	1	0	1
1	0	1	1	1
1	1	0	0	1
1	1	0	1	1
1	1	1	0	1
1	1	1	1	1

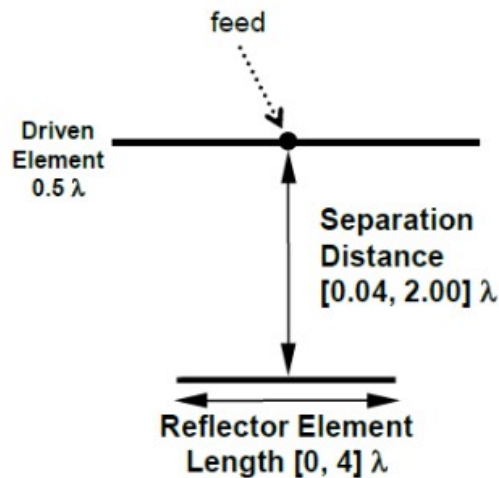
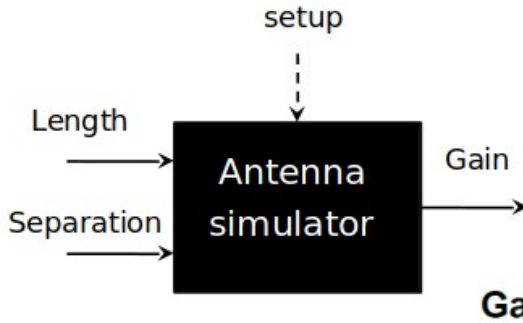
fitness score = 14

where the fitness function is the number of correct bits with respect to the truth table:

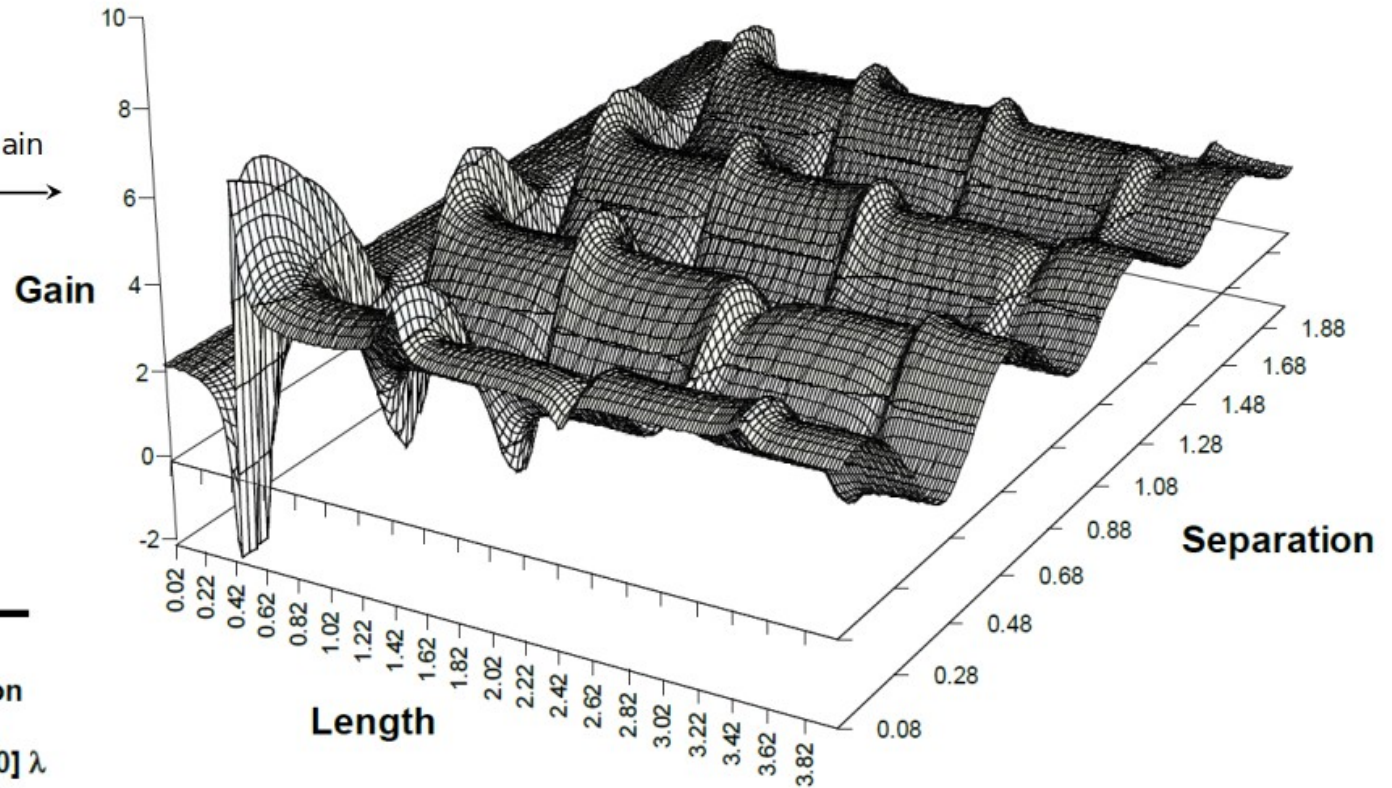
$$y(a, b, c, d) = a \vee b \vee c \vee d.$$

(max. fitness is 16)

Example 2: an antenna optimization, a search space illustration



Linden, 1997



What are we looking for?

Length: 7 bits

Separation: 7 bits

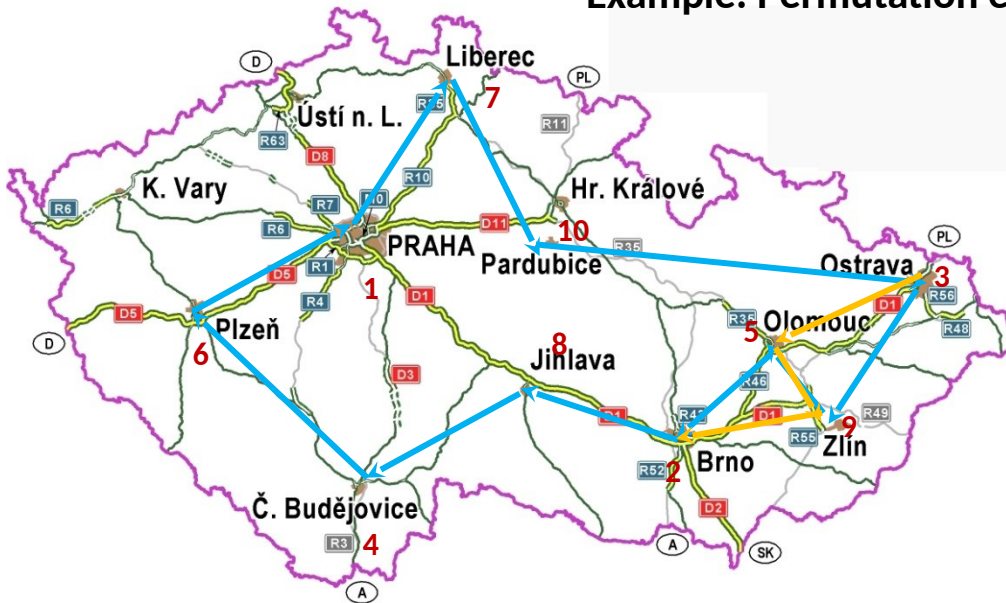
Search space size: $2^{14} = 16384$

The **Gain** is our **objective (fitness) function** – to be maximized.

In real applications, the search space is usually much larger and multidimensional. It cannot be visualized, its form is unknown!

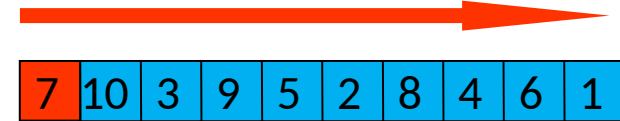
| Example 3: encoding a route for Traveling Salesman Problem

Example: Permutation encoding for TSP



- 1 - Praha
- 2 - Brno
- 3 - Ostrava
- 4 - České Budějovice
- 5 - Olomouc
- 6 - Plzeň
- 7 - Liberec
- 8 - Jihlava
- 9 - Zlín
- 10 - Pardubice

Encoded
route



mutation



Notice: genetic operators have to be devised specifically for the given representation (e.g. the mutation for permutations: swap two randomly selected integers; **the crossover operator is usually not used here because often the offspring are not valid permutations!**).

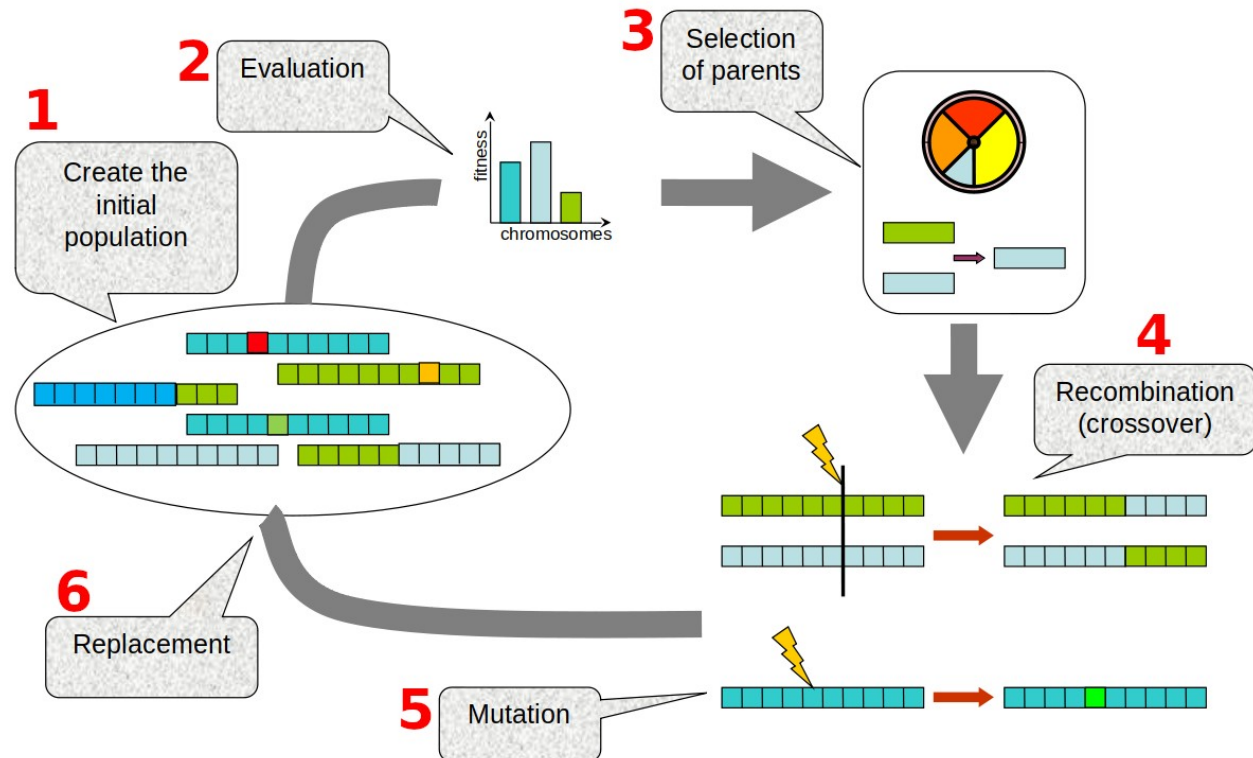
- Examples of the most common encodings (in general):
 - a vector of integers, characters, real numbers... (in computer always have a bin. form)
 - a permutation - see the TSP example above
 - a syntax tree - in GP
 - an acyclic directed graph - in CGP

The main loop of EA

1. Generate the initial population (randomly or heuristically).
2. Evaluate each candidate solution using the fitness function.

WHILE termination_condition_is_not_satisfied DO

3. Select parents (according to the fitness).
- 4-5. Create offspring from parents using crossover and mutation.
6. Create a new population by replacing some/all old individuals by the offspring individuals.
Evaluate the new population.



| Selection of parents

- Solutions with better fitness have higher probability to be selected.
- Major methods:
 - **Weighted roulette wheel selection (fitness proportional)** – a relative fitness is calculated for every individual: $f_r(i) = f(i) / \sum f(i)$, $i = 1 \dots \text{pop_size}$
Individuals are selected proportionally to f_r .
 - Problem: If there is an individual with very high f_r and remaining ones have very low f_r , the best one will be selected all the time \Rightarrow degeneration of the population
 - **Rank selection** – similar to the roulette, but the candidate solutions are sorted according to the fitness first, then a “rank” values are assigned to them based on the ordering (not on the absolute fitness values).
 - **Tournament selection** – randomly chooses K individuals from the population and selects one with the best fitness score as a parent. Often $K = 2$. **Simple, efficient and most widely used.**
- The selection procedure is repeated until the required number of parent individuals is obtained.
- An individual can be selected multiple times.
- The selection of the currently best-scored individual is not guaranteed.

Example 4: selection using the weighted roulette wheel

A sample populations & fitness:

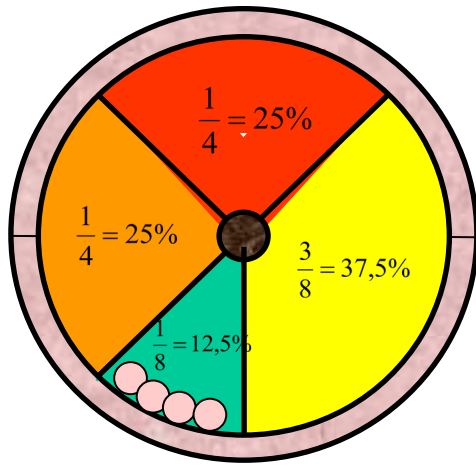
1 1 0 0	-> fitness (12) = 6
1 0 0 0	-> fitness (8) = 4
0 1 0 1	-> fitness (5) = 4
0 0 1 1	-> fitness (3) = 2

The number of portions

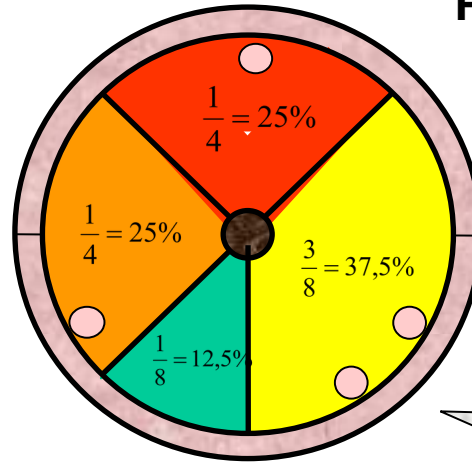
$$6 + 4 + 4 + 2 = 16$$

Calculated portions for the individuals on the wheel

1 1 0 0	$\rightarrow \frac{6}{16} = \frac{3}{8} = 37,5\%$
1 0 0 0	$\rightarrow \frac{4}{16} = \frac{1}{4} = 25\%$
0 1 0 1	$\rightarrow \frac{4}{16} = \frac{1}{4} = 25\%$
0 0 1 1	$\rightarrow \frac{2}{16} = \frac{1}{8} = 12,5\%$



very unlikely



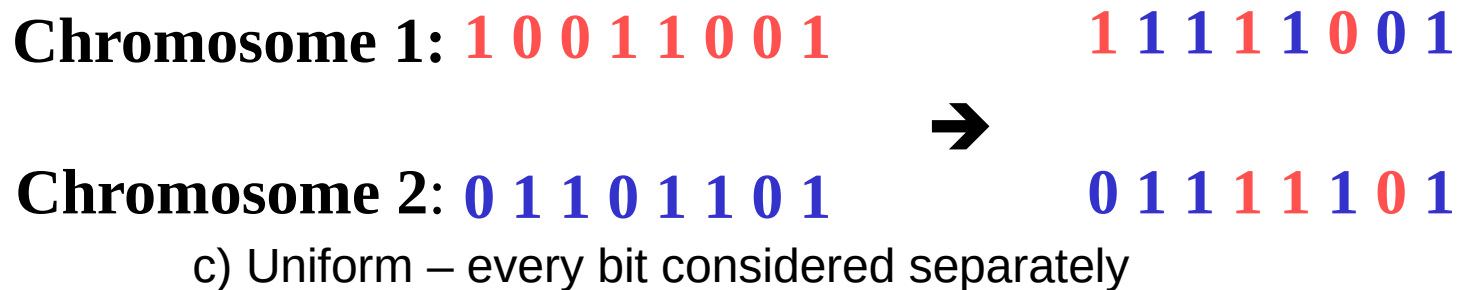
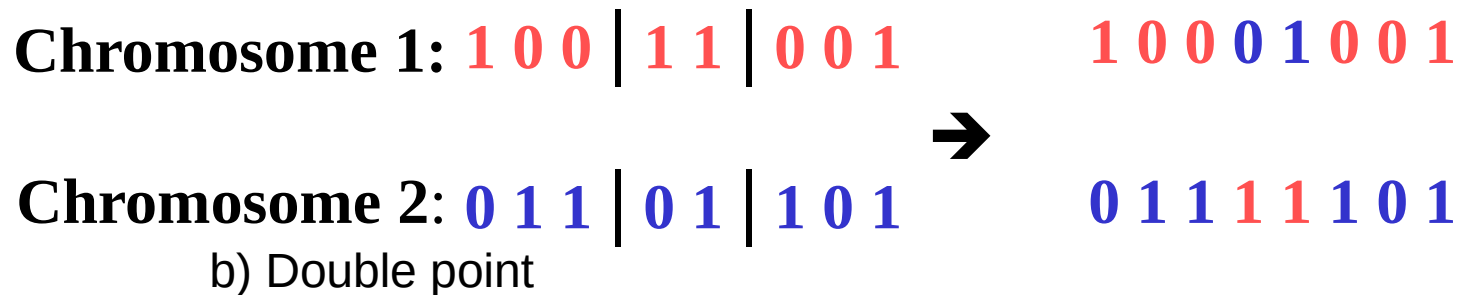
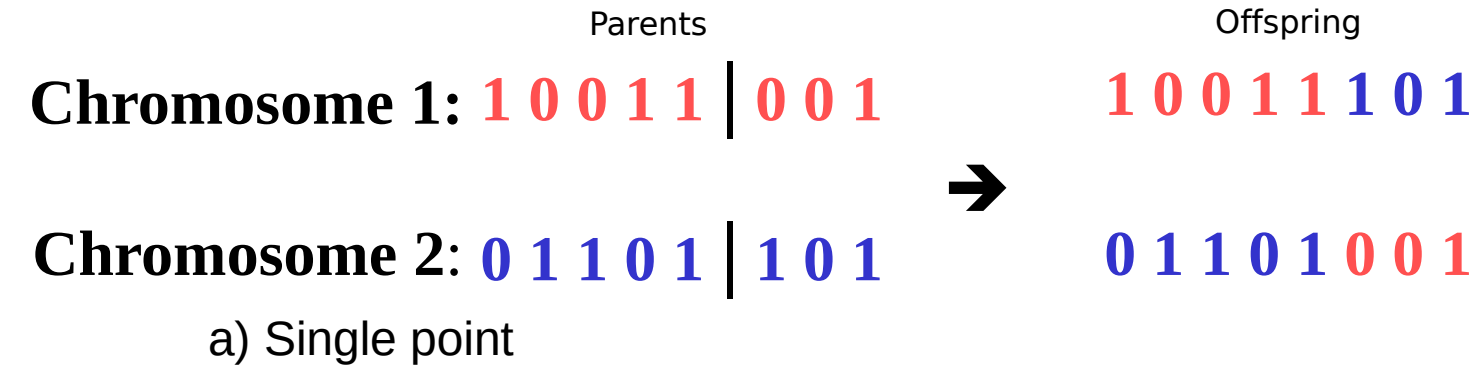
typical result

Resulting parents:

1 1 0 0
1 1 0 0
0 1 0 1
1 0 0 0

In order to obtain 4 individuals, the roulette is run 4 times.

Genetic operators – crossover



In EA the crossover is usually applied with a probability, e.g. ~0.7.

Genetic operators – mutation

- Mutation is typically a small change in a single chromosome – e.g. a bit inversion:

Chromosome: 1 0 0 1 1 0 0 1 0 1 1 0 0 1

↓ ↓

After mutation: 1 0 0 0 1 0 0 1 0 1 1 0 1 1

In EA the mutation is usually applied with a low probability, e.g. ~ 0.05 .

The mutation probability may be higher (sometimes even $\rightarrow 1,0$) if the mutation is used as a single genetic operator. However, the mutation always should modify only a very small portion of the chromosome.

Replacement and termination criteria

- Replacement mechanisms

- **Steady-state EA:** The new population of P individuals will contain N individuals from the old population and M new offspring individuals ($P = N + M$).
- **Generational EA:** The new population contains only new offspring individuals.

- **Elitism** – the best e individuals of the previous population are always copied to the new population (usually $e = 1$).

- Typical termination criteria

- A solution with fitness = 100% was discovered.
- The maximum number of generations or time was exhausted.
- There has not been any progress in the fitness in the last k generations.

| Summary of EA utilization

- In order to apply EA to solve a given problem, the designer needs to design:
 - problem encoding
 - genetic operators
 - fitness function
- Then some parameters have to be set up:
 - population size
 - max. number of generations
 - probabilities of mutation, crossover...
 - the way of initialization of the population
 - the termination condition
- As EA are stochastic algorithms, the results need to be evaluated statistically:
 - for each EA settings run the EA several times (at least 20x, better 100x)
 - report the best / average / median / standard deviation of the results
- **Tune the parameters or repeat the entire process if EA does not work.**

In case of real-world applications, there is no relevant theory for solving these issues!
The design and tuning of EA is primarily an experimental work.

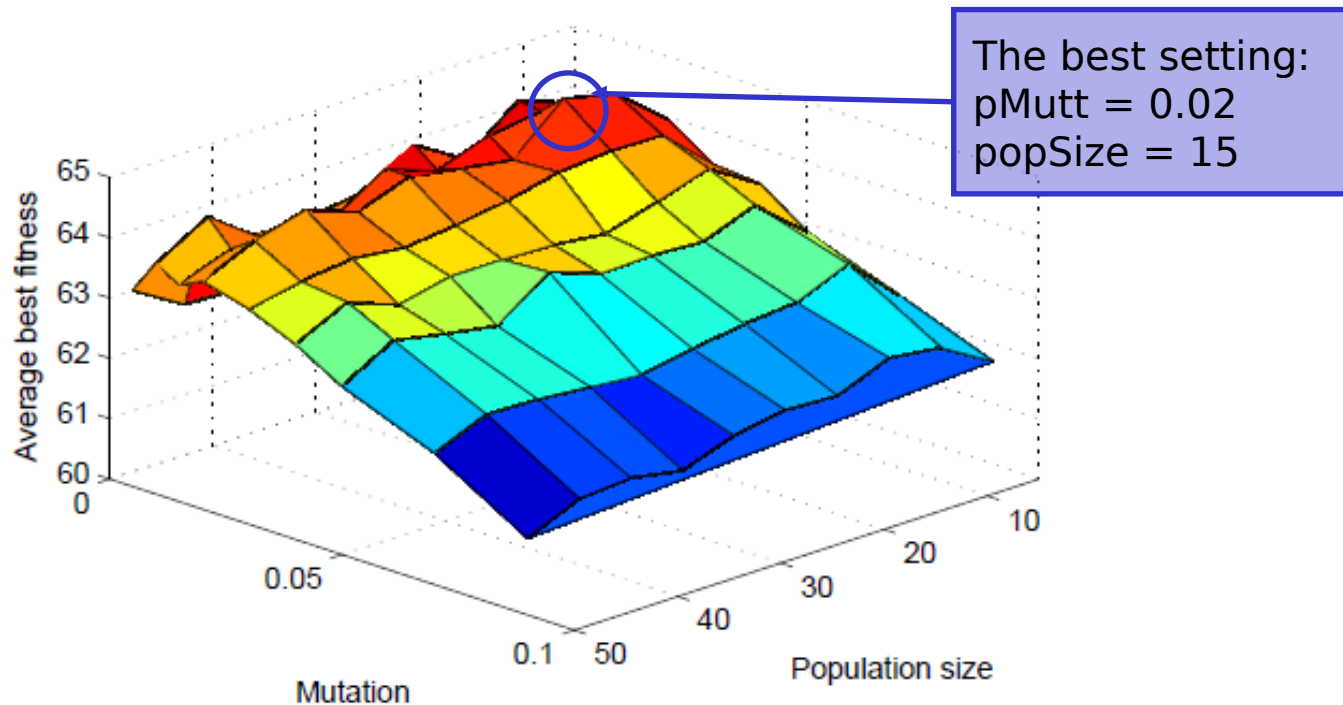
| Is EA1 better than EA2?

- For EA1: $P1$ is the population size and $G1$ is the max. number of generations, analogically $P2$ and $G2$ for the EA2.
- Then $P1 * G1$ (resp. $P2 * G2$) is the number of candidate solutions evaluated within a single EA run.
- In order to adequately compare EA1 and EA2, **$P1 * G1 = P2 * G2$ must hold!**
Why? Because if more individuals can be evaluated for an EA (i.e. a bigger part of the search space can be explored), then this EA has a higher chance to provide better results (the comparison would be inadequate).
- **A single run is not meaningful** (may be a lucky chance)!
- The quality of EA can be statistically measured as (for r independent runs):
 - the average (median) number of evaluations (or generations) needed to find an acceptable solution (given by the termination condition),
 - the number of successful runs (when 100% fitness was achieved) out of r runs,
 - the average (median) fitness of the best solutions after the max. generations,
 - and many other methods exists.

Example 5: visualization of statistical tuning of some EA parameters

Principles of the statistical evaluation - an example:

For 10 different values of P , the corresponding values $G=40000/P$, and 10 values of the mutation probability, we performed 50 independent runs **for each (P , G , P_{mut})**. All other EA parameters remain fixed. In total, $50 \times 10 \times 10$ independent runs are performed. The average best fitness out of 50 independent runs for each settings is visualized.



- Introduction
- Evolutionary algorithms
- **Genetic programming**
- Cartesian genetic programming
- Applications from FIT
- Summary

I Genetic Programming (GP)

- Genetic Programming is an EA for the evolutionary design of computer programs.
- The goal is to find a **program** (close to) optimally working for arbitrary input data and providing the desired results.
- In GP:
 - **The EA is almost identical with that on page 15.**
 - Candidate programs are represented using **syntax trees** (or other suitable structures), machine code instructions or strings of integers... all representing an executable (calculable) function.
 - Crossover and mutation are adapted adequately, sometimes more complex genetic operators can be used, e.g. subprogram definition.
 - In order to obtain the fitness score, a candidate program is executed using some given training data.
 - The result of GP (i.e., the best-performing program) is evaluated using the test data.

- In addition to that stated on page 21, we need to define:

A set of **terminals (T)**, typically: constants, input variables, functions of 0-arity (e.g. time(), rand(), halt() etc.) - the terminals will appear only in the leafs of the trees.

A set of **functions (F)** - non-terminals, typically: +, -, *, /, AND, NOT, ... *any function with at least 1 input – these functions will appear in non-leaf nodes of the trees.*

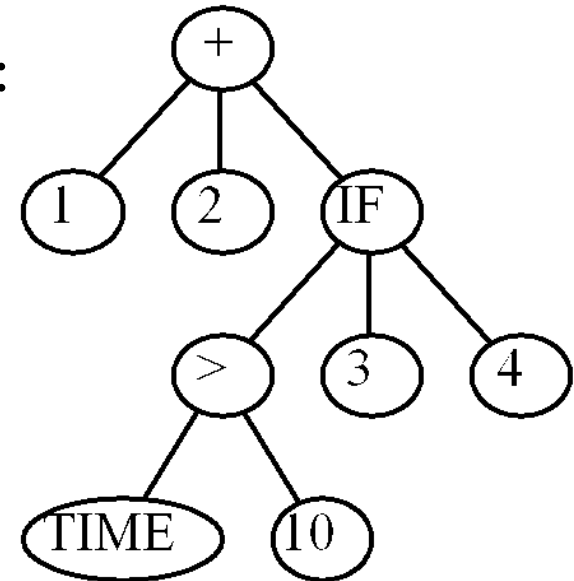
Note: some operations need to be defined as **protected functions** in order to avoid program crash during execution, e.g. division:

$$\text{div}(a, b) := \text{if } b \neq 0 \text{ then return } a/b \text{ else return } 1$$

Example 6: a sample GP tree

- Consider $T = \{ 0, \dots, 9, \text{TIME} \}$, $F = \{ +, -, >, <, \text{IF} \}$
- A syntax tree may look like this, e.g.:
- In GP, however, the trees are represented in a suitable language, e.g. **LISP**:

(+ 1 2 (IF (> TIME 10) 3 4))



- Or the same may be written in C:

```
int a;  
if (TIME() > 10) a = 3 else a = 4;  
return 1 + 2 + a;
```

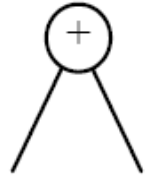
| GP – random program generation

- Given a maximum tree depth = h , the generation starts from the root node.
- Methods for generating the initial population of trees:
 - **Grow** – randomly choose items from T or F until h is reached, then use only T.
 - **Full** – randomly choose items only from F until h is reached, then use only T.
 - **Ramped Half-and-Half** – for each depth $1...h$ generate one half of individuals in the population by the Grow method and the other half of individuals by the Full method.

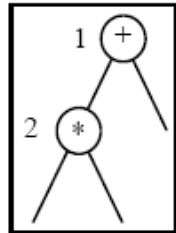
Example 7: a random tree generation

- Terminal set $T = \{A, B, C\}$
- Function set $F = \{+, -, *, \%, \text{IFLTE}\}$

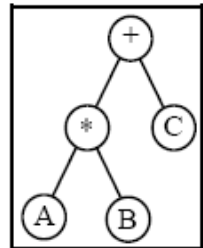
BEGIN WITH TWO-ARGUMENT +



CONTINUE WITH TWO-ARGUMENT *

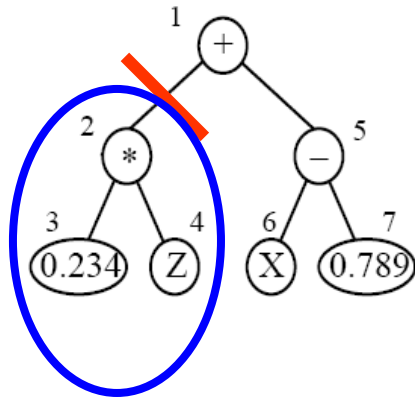


FINISH WITH TERMINALS A, B, AND C

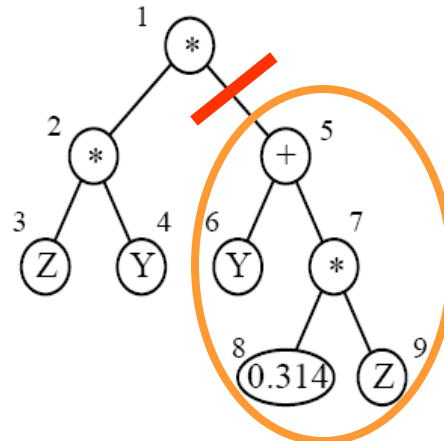


- The result is a syntactically valid executable program (provided the set of functions is closed)

2 parents



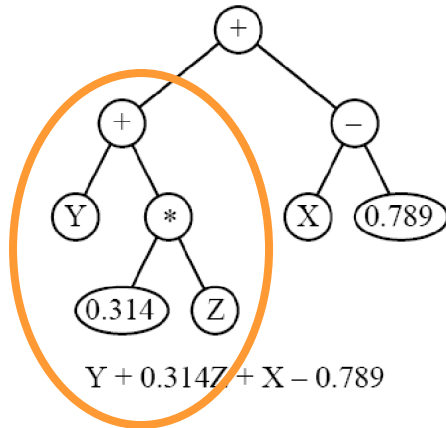
$$0.234Z + X - 0.789$$



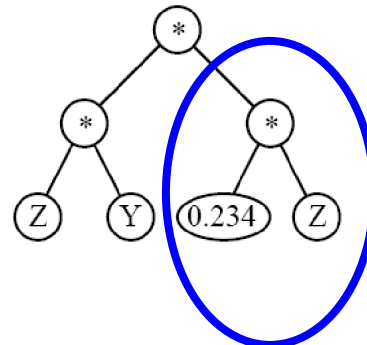
$$ZY(Y + 0.314Z)$$

Crossover swaps randomly selected sub-trees.

2 offspring

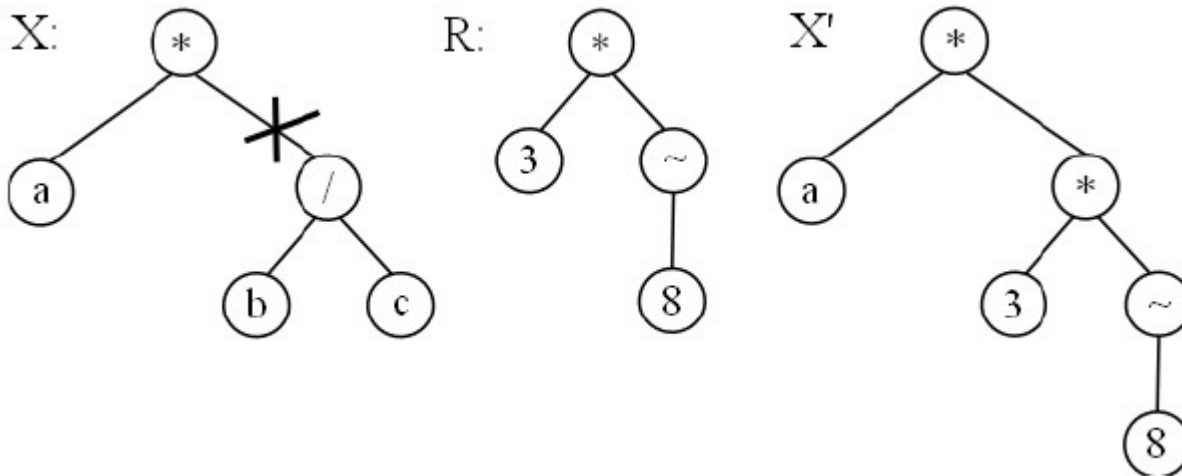


$$Y + 0.314Z + X - 0.789$$



$$0.234Z^2Y$$

- Randomly select a node (a corresponding sub-tree is marked) in X and replace the sub-tree by a randomly generated sub-tree R to get X'.



A typical GP application: Symbolic Regression (~ data fitting)

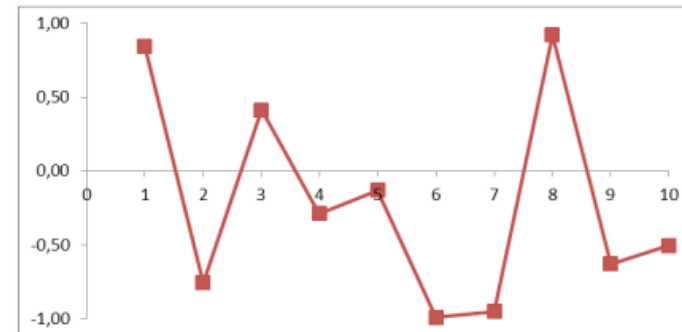
Given some data (usually from a measurement), the goal of symbolic regression is to discover a mathematical prescription that fits the data as best as possible. For example, for 10 test cases given by the table, we are trying to minimize the function f (the sum of absolute differences):

$$f = \sum_{i=1}^{K=10} |y_i - w_i|$$

$K=10$ ← The number of test cases

y_i ↑ Desired result

w_i ↑ Candidate program result



Test cases

x	y
1	0,84
2	-0,76
3	0,41
4	-0,29
5	-0,13
6	-0,99
7	-0,95
8	0,92
9	-0,63
10	-0,51

Why to use GP?

- An arbitrary function set can be used (no assumptions in terms of mathematical properties).
- Can be extended to n-input/m-output functions and many objectives can be optimized together (error, size of the tree, speed...).
- Constraints can easily be handled.

Disadvantages: computationally expensive, stochastic

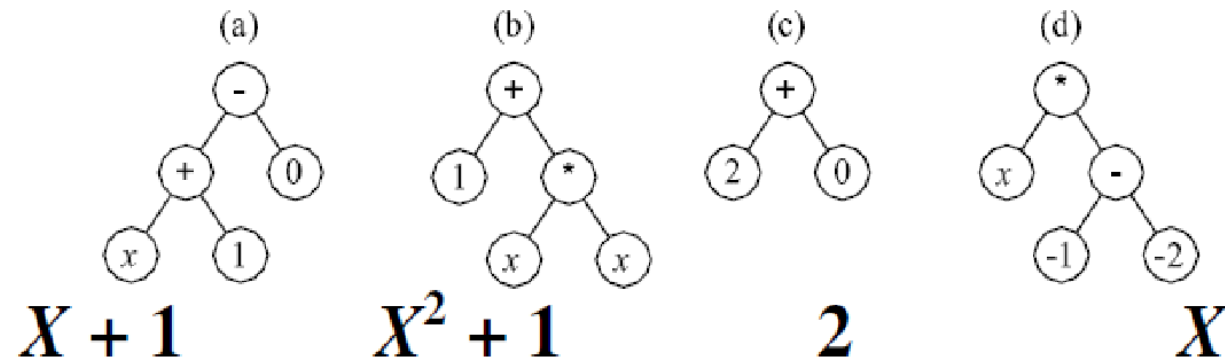
Example 8: a simple example of symbolic regression using GP

Independent variable X	Dependent variable Y
-1.00	1.00
-0.80	0.84
-0.60	0.76
-0.40	0.76
-0.20	0.84
0.00	1.00
0.20	1.24
0.40	1.56
0.60	1.96
0.80	2.44
1.00	3.00

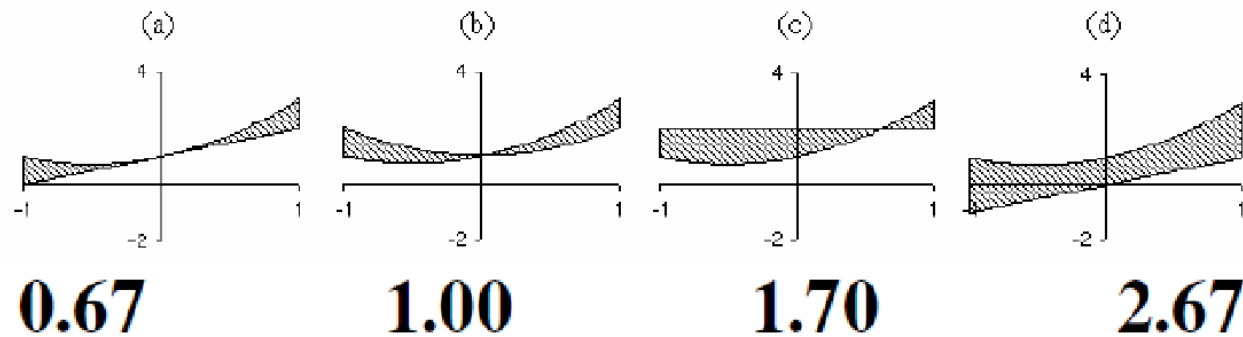
	Objective:	Find a computer program with one input (independent variable x), whose output equals the value of the quadratic polynomial $x^2 + x + 1$ in range from -1 to +1.
1	Terminal set:	$T = \{X\}$
2	Function set:	$F = \{+, -, *, \%\}$ NOTE: The protected division function $\%$ returns a value of 1 when division by 0 is attempted (including 0 divided by 0)
3	Fitness:	The sum of the absolute value of the differences (errors), computed (in some way) over values of the independent variable x from -1.0 to +1.0, between the program's output and the target quadratic polynomial $x^2 + x + 1$.
4	Parameters:	Population size $M = 4$.
5	Termination:	An individual emerges whose sum of absolute errors is less than 0.1

Example 8: a simple example of symbolic regression using GP

GENERATION 0

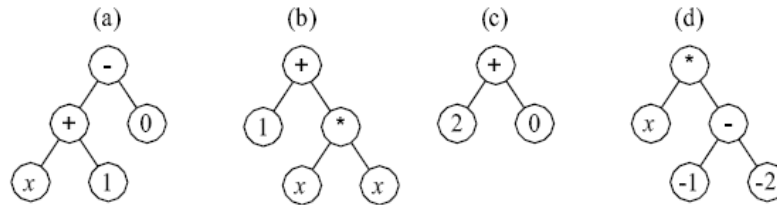


FITNESS

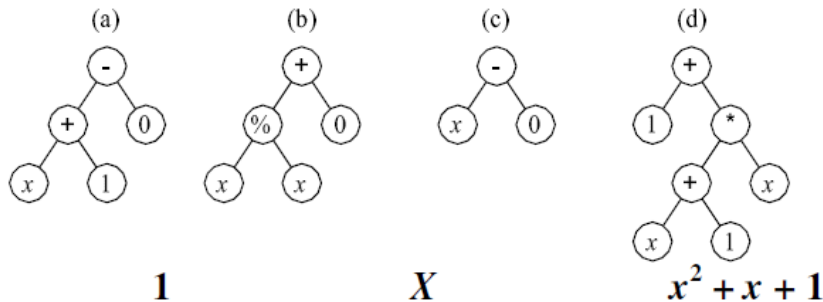


Example 8: a simple example of symbolic regression using GP

GENERATION 0



GENERATION 1



Copy of (a)	Mutant of (c)	First offspring of crossover of (a) and (b)	Second offspring of crossover of (a) and (b)
	—picking “2” as mutation point	—picking “+” of parent (a) and left-most “x” of parent (b) as crossover points	—picking “+” of parent (a) and left-most “x” of parent (b) as crossover points

Example 9: a more complex example of symbolic regression...

SYMBOLIC REGRESSION OF QUARTIC POLYNOMIAL $X^4+X^3+X^2+X$ (WITH 21 FITNESS CASES)

Independent variable (Input)	Dependent Variable (Output)
X	Y
-1.0	0.0000
-0.9	-0.1629
-0.8	-0.2624
-0.7	-0.3129
-0.6	-0.3264
-0.5	-0.3125
-0.4	-0.2784
-0.3	-0.2289
-0.2	-0.1664
-0.1	-0.0909
0	0.0
0.1	0.1111
0.2	0.2496
0.3	0.4251
0.4	0.6496
0.5	0.9375
0.6	1.3056
0.7	1.7731
0.8	2.3616
0.9	3.0951
1.0	4.0000

TABLEAU—SYMBOLIC REGRESSION OF QUARTIC POLYNOMIAL $X^4+X^3+X^2+X$

Objective:	Find a function of one independent variable, in symbolic form, that fits a given sample of 21 (x_i, y_i) data points
Terminal set:	x (the independent variable).
Function set:	$+$, $-$, $*$, $\%$, SIN, COS, EXP, RLOG
Fitness cases:	The given sample of 21 data points (x_i, y_i) where the x_i are in interval $[-1, +1]$.
Raw fitness:	The sum, taken over the 21 fitness cases, of the absolute value of difference between value of the dependent variable produced by the individual program and the target value y_i of the dependent variable.
Standardized fitness:	Equals raw fitness.
Hits:	Number of fitness cases (0 – 21) for which the value of the dependent variable produced by the individual program comes within 0.01 of the target value y_i of the dependent variable.
Wrapper:	None.
Parameters:	Population size, $M = 500$. Maximum number of generations to be run, $G = 51$.
Success Predicate:	An individual program scores 21 hits.

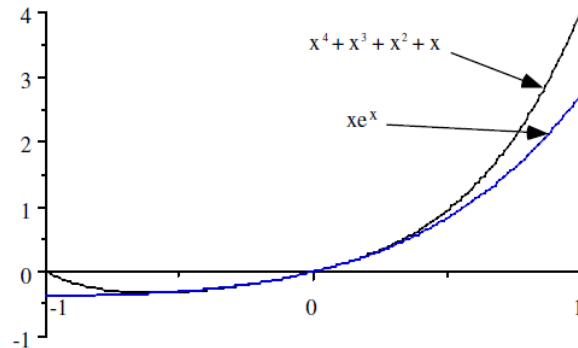
Example 9: a more complex example of symbolic regression...

**BEST-OF-GENERATION INDIVIDUAL IN
GENERATION 0 WITH RAW FITNESS OF
4.47 (AVERAGE ERROR OF 0.2)**

$(* X (+ (+ (- (\% X X) (\% X X))$
 $(SIN (- X X))) (RLOG (EXP (EXP$
 $X)))))$

Equivalent to

$x e^x$



**CREATION OF GENERATION 1 FROM
GENERATION 0**

- In the so-called "generational" model for genetic algorithms, a new population is created that is equal in size to the old population
 - 1% mutation (i.e., 5 individuals out of 500)
 - 9% reproduction (i.e., 45 individuals)
 - 90% crossover (i.e., 225 pairs of parents — yielding 450 offspring)
- All participants in mutation, reproduction, and crossover are chosen from the current population **PROBABILISTICALLY, BASED ON FITNESS**
 - Anything can happen
 - Nothing is guaranteed
 - The search is heavily (but not completely) biased toward high-fitness individuals
 - The best is not guaranteed to be chosen
 - The worst is not necessarily excluded
 - Some (but not much) attention is given even to low-fitness individuals

Example 9: a more complex example of symbolic regression...

**BEST-OF-GENERATION INDIVIDUAL IN
GENERATION 2 WITH RAW FITNESS OF
2.57 (AVERAGE ERROR OF 0.1)**

```
(+ (* (* (+ X (* X (* X (% (% X
X) (+ X X))))))
    (+ X (* X X))) X) X)
```

Equivalent to...

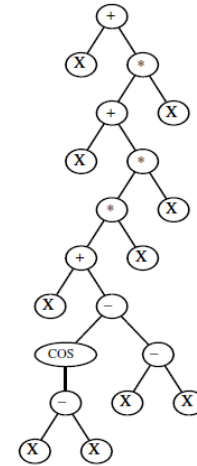
$$x^4 + 1.5x^3 + 0.5x^2 + x$$

**BEST-OF-RUN INDIVIDUAL IN
GENERATION 34 WITH RAW FITNESS
OF 0.00 (100%-CORRECT)**

```
(+ X (* (+ X (* (* (+ X (- (COS
(- X X)) (- X X))) X) X)) X))
```

Equivalent to

$$x^4 + x^3 + x^2 + x$$



Notice: $\text{COS}(X - X)$, $(X - X)$ etc. are examples of the so-called **intron** – a piece of code which does not contribute to the fitness, but increases the size of candidate solution.

$\text{COS}(X - X) = 1$, $(X - X) = 0$ for any X etc.

Bloat – a phenomenon in which the program size grows during the evolution, but the fitness remains unchanged. Various techniques were developed to eliminate the bloat.

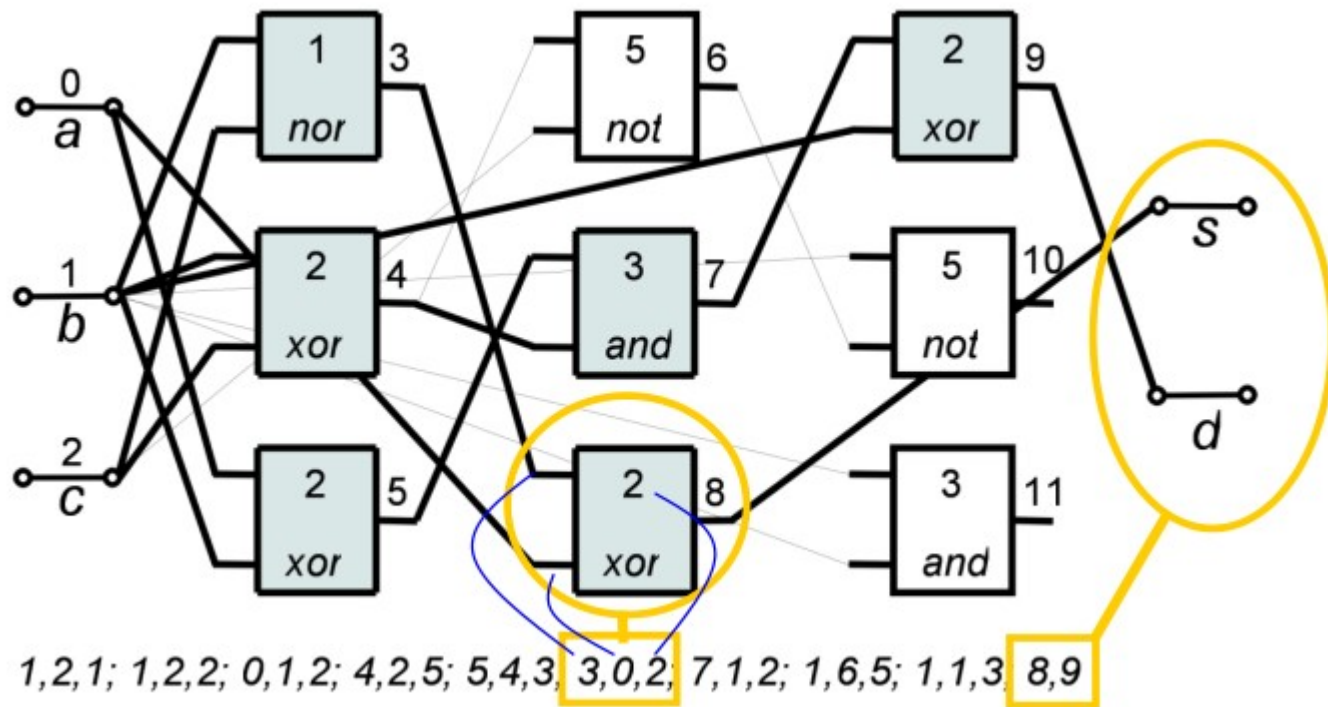
- Introduction
- Evolutionary algorithms
- Genetic programming
- **Cartesian genetic programming**
- Applications from FIT
- Summary

- So far we discussed so-called “tree-based” GP.
- There are also other types (Linear GP, developmental GP...).
- CGP is a type of GP in which the programs are represented as *oriented acyclic graphs* the nodes of which form a matrix.
- CGP encodes these graphs as **fixed length integer vectors**. This representation is particularly suitable for designing digital circuits.
- CGP parameters include:
 - The number of rows, n_r
 - The number of columns, n_c
 - The number of primary inputs, n_i
 - The number of primary outputs, n_o
 - The maximum arity of a block function, n_a
 - Level-back parameter (L-back), L
 - The set of block functions, Γ

In genotypes, there are n_a+1 integers per node (describing connection of node inputs and node function) and n_o integers for describing connection of primary outputs. Therefore, the genomes are of the length $n_c n_r (n_a + 1) + n_o$ integers.

Example 10: a logic circuit represented in CGP

- Consider $n_r=3$, $n_c=3$, $n_i=3$, $n_o=2$, $n_a=2$, $L=3$,
 $\Gamma = \{\text{NAND}^{(0)}, \text{NOR}^{(1)}, \text{XOR}^{(2)}, \text{AND}^{(3)}, \text{OR}^{(4)}, \text{NOT}^{(5)}\}$

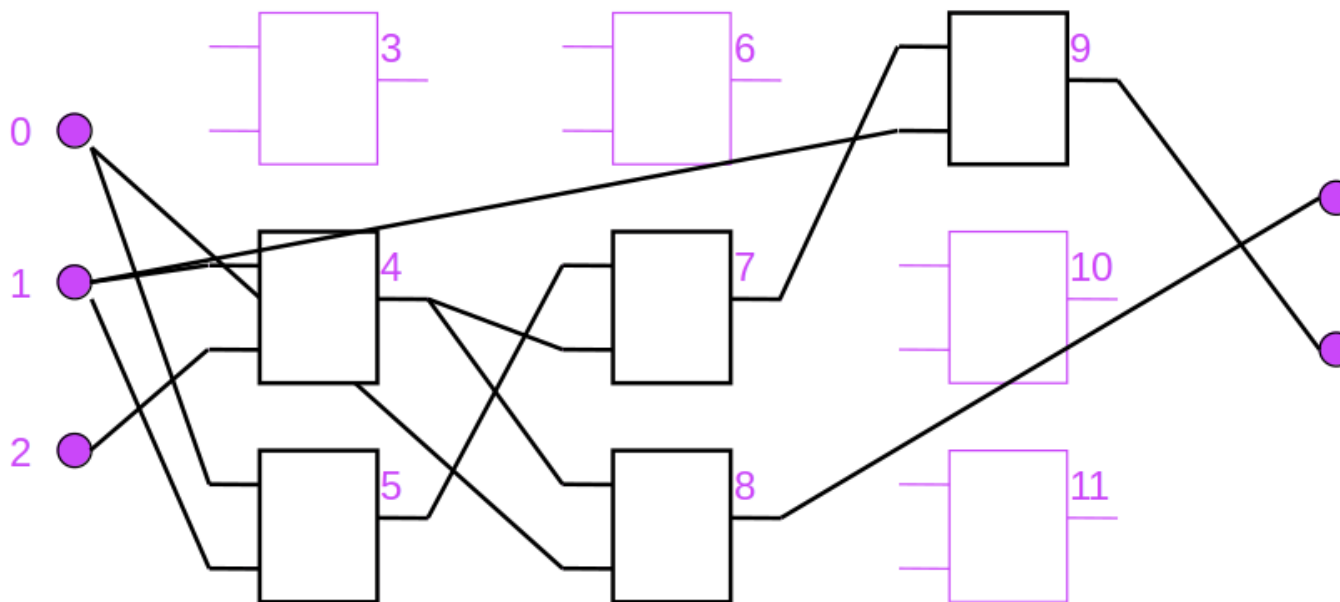


Circuit construction from a CGP chromosome

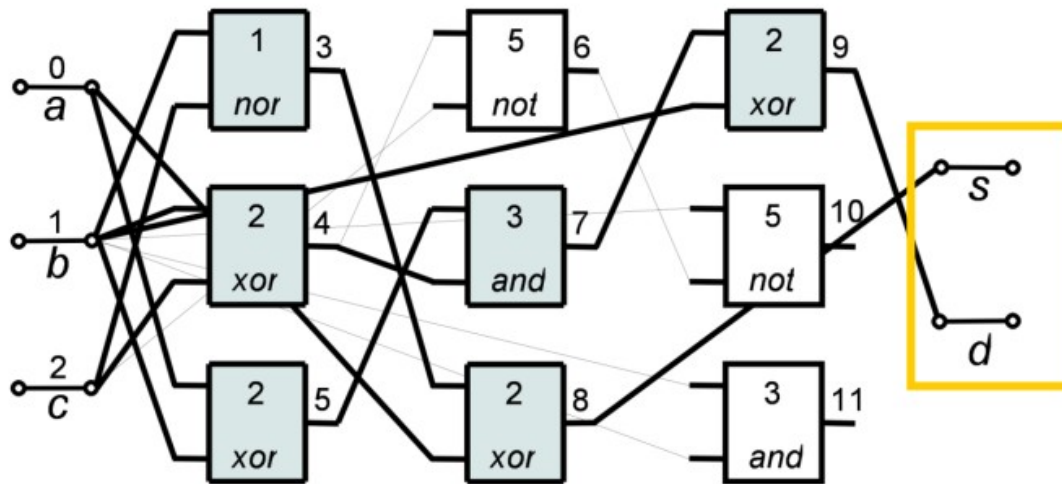
- (1) The processing of the chromosome goes from the outputs to the inputs. The nodes that need to be evaluated are stored in a stack.
- (2) Then, the functions are taken from the stack and evaluated, the results are stored in an array at the indexes given by the figure.

A chromosome (from the previous example):

(1,2,1) (1,2,2) (0,1,2) (4,2,5) (5,4,3) (4,0,2) (7,1,2) (1,6,5) (1,1,3) (8,9)



Fitness function for circuit design using CGP



1,2,1; 1,2,2; 0,1,2; 4,2,5; 5,4,3; 3,0,2; 7,1,2; 1,6,5; 1,1,3; 8,9

a	b	c	d	s
0	0	0	0	0
0	0	1	0	1
0	1	0	0	1
0	1	1	1	0
1	0	0	0	1
1	0	1	1	0
1	1	0	1	0
1	1	1	1	1

Specification
(1-bit adder),
target table:

=> fitness = 16

a	b	c	d	s
0	0	0	0	1
0	0	1	0	0
0	1	0	0	0
0	1	1	1	0
1	0	0	0	0
1	0	1	1	1
1	1	0	1	1
1	1	1	1	1

=> fitness = 10

Typical fitness function (to be maximized):

$$f = \text{Max} - \sum_{i=1}^K HD(y_i, w_i)$$

\nwarrow The number of test vectors
 \uparrow Hamming distance
 (between circuit response
 and desired response)

Additional objectives:

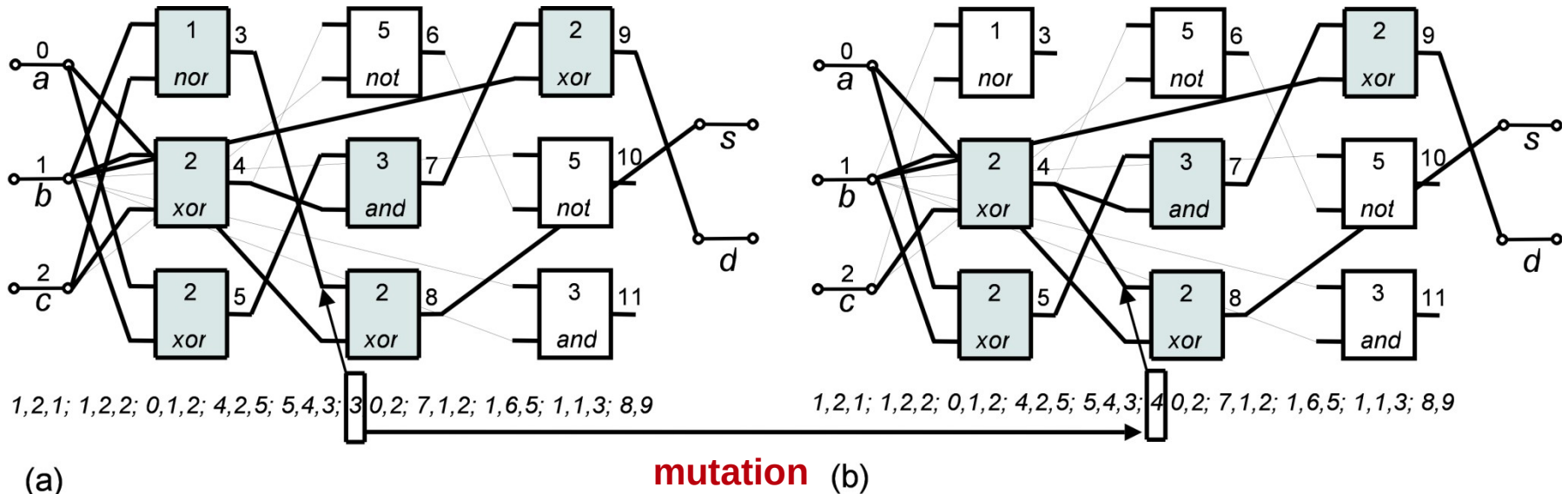
- area (the number of gates)
- delay
- power consumption etc.

Max = #outputs * $2^{\text{\#inputs}}$ (in our case $2 * 2^8 = 16$)

$K = 2^{\text{\#inputs}}$ for combinational circuits. Not scalable!!!

CGP uses mutation-based search (usually without crossover)

- Mutation: Randomly select *h* integers and replace them by randomly generated (but legal) values.
- Mutation can be useful, harmful or neutral.



a	b	c	d	s
0	0	0	0	1
0	0	1	0	0
0	1	0	0	0
0	1	1	1	0
1	0	0	0	0
1	0	1	1	1
1	1	0	1	1
1	1	1	1	1

=> fitness = 10

a	b	c	d	s
0	0	0	0	0
0	0	1	0	1
0	1	0	0	1
0	1	1	1	0
1	0	0	0	1
1	0	1	1	0
1	1	0	1	0
1	1	1	1	1

=> fitness = 16
(for full adder)

More on mutations in CGP -> Goldman B.W., Punch W.F.: Analysis of Cartesian Genetic Programming's Evolutionary Mechanisms. IEEE Trans. Evolutionary Computation 19(3): 359-373 (2015)

Algorithm 1: CGP

Input: CGP parameters, fitness function

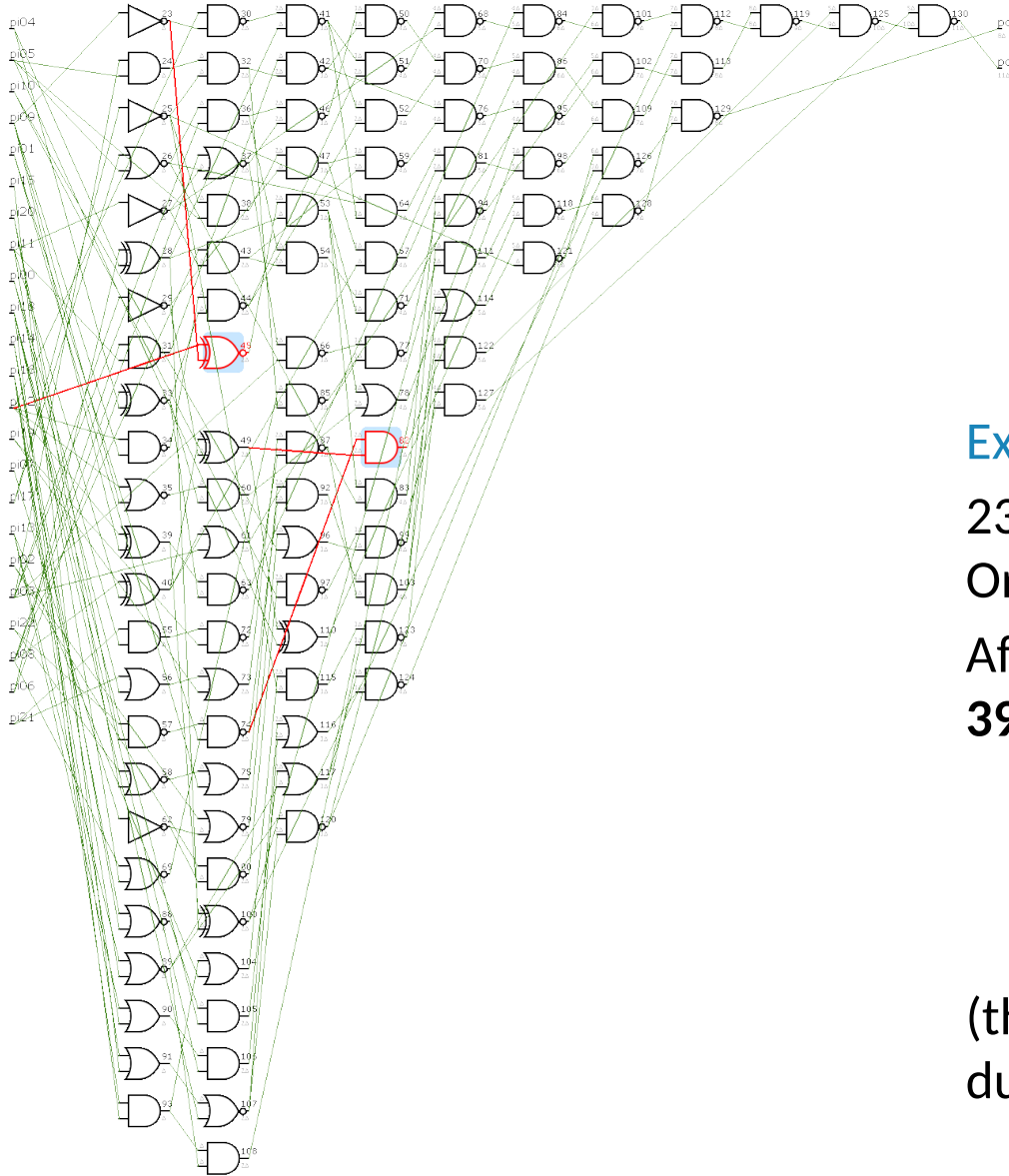
Output: The highest scored individual p and its fitness

```
1  $P \leftarrow$  randomly generate population; // or use conventional designs
2 EvaluatePopulation( $P$ );  $p \leftarrow$  highest-scored-individual( $P$ );
3 while  $\langle$ terminating condition not satisfied $\rangle$  do
4    $\alpha \leftarrow$  highest-scored-individual( $P$ );
5   if  $\text{fitness}(\alpha) \geq \text{fitness}(p)$  then
6      $p \leftarrow \alpha$ ;
7    $P \leftarrow$  create  $\lambda$  offspring of  $p$  using mutation;
8   EvaluatePopulation( $P$ );
9 return  $p$ ,  $\text{fitness}(p)$ ;
```

\geq is better than $>$ as it increases the diversity of the population and leads to better solutions.

- Introduction
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Optimization of the number of gates of existing circuits using CGP



Example: Cordic circuit

23 inputs, 2 outputs

Original: **106 gates** (LGSynth93)

After the CGP optimization:

39 gates

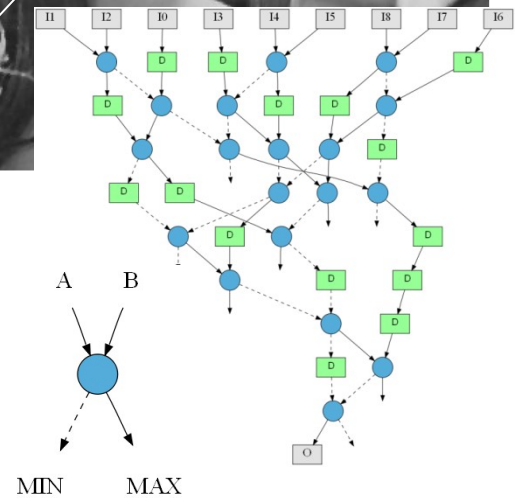
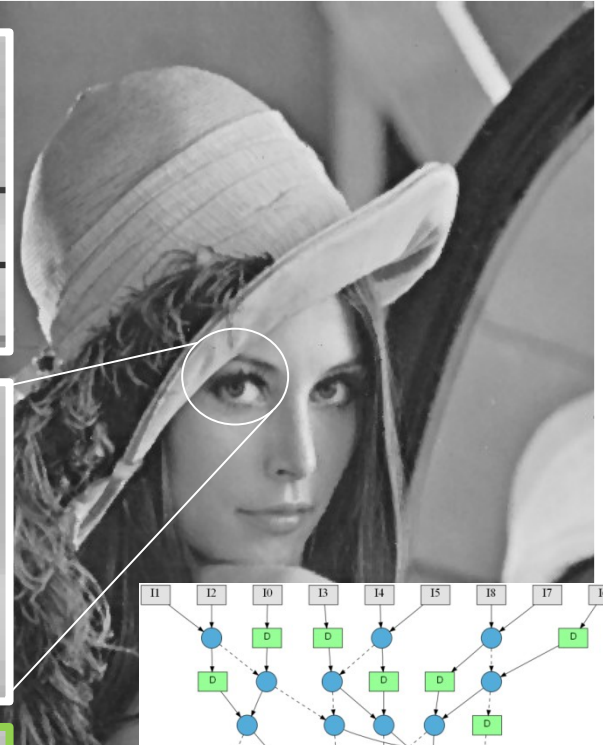
(the best circuits discovered
during a CGP run are shown)

Non-linear image filters

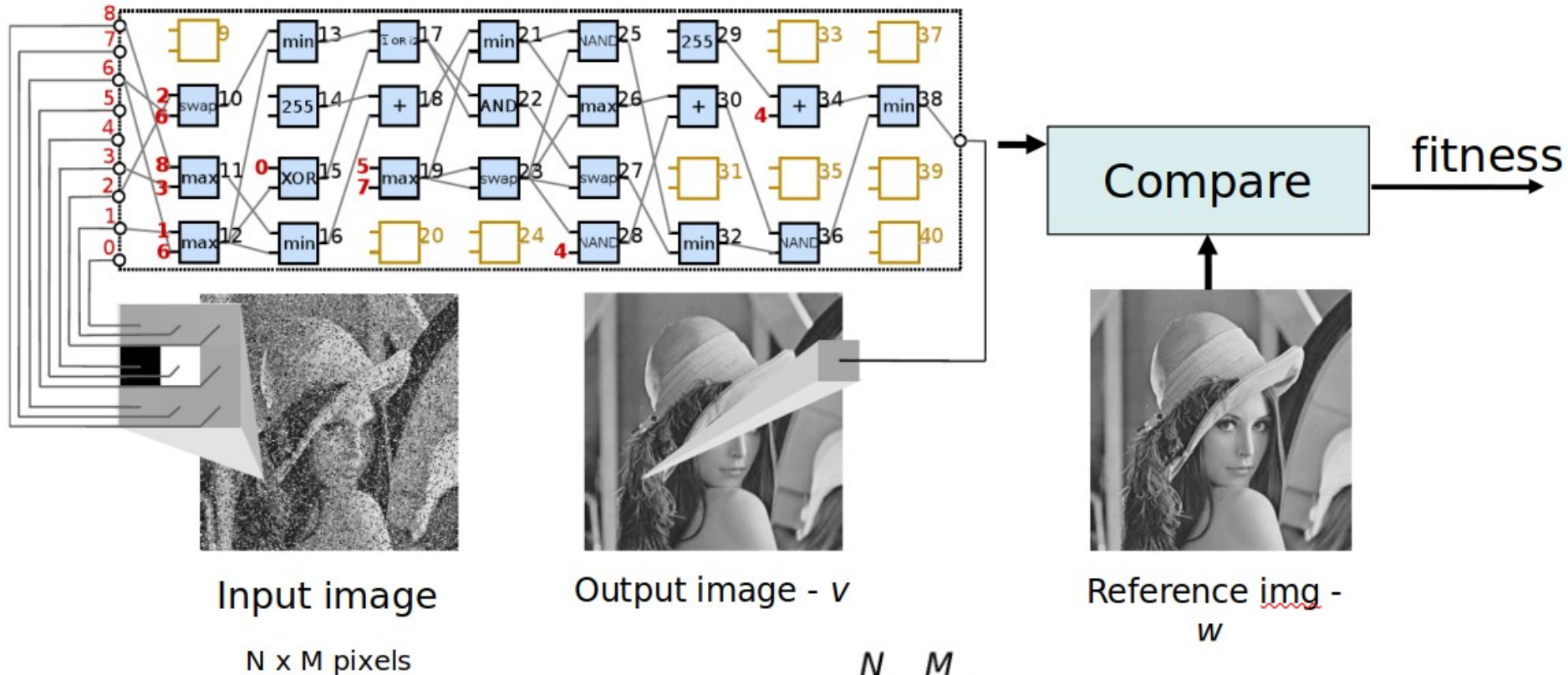
corrupted image
(10% pixels, impulse noise)



filtered image
(9-input median filter)



Evolutionary design of image filters using CGP



$$fitness = \sum_{i=1}^N \sum_{j=1}^M |v(i, j) - w(i, j)|$$

- $n_i=9$, $n_o=1$, $n_c=8$, $n_r=4$, $n_a=2$, $\lambda = 7$; 30,000 generations/run; mutation: max. 5%
- Function set (over 8 bits) = {+, min, max, swap, constants, average, log. functions}

CGP: Evolutionary design with 8-bit components (FPGA)

- a) Image corrupted by 5% salt-and-pepper noise

PSNR: 18.43 dB (peak signal to noise ratio)

- b) Original image

- c) Median filter (kernel 3x3)

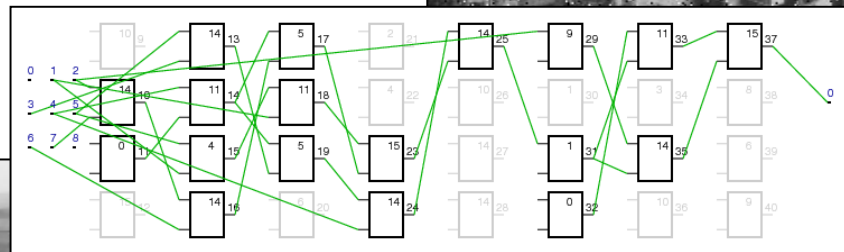
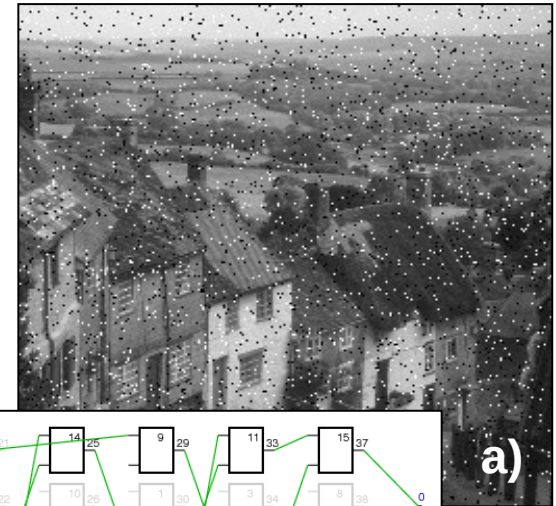
PSNR: 27.92 dB

268 FPGA slices; 305 MHz

- d) Evolved filter (kernel 3x3)

PSNR: 37.50 dB

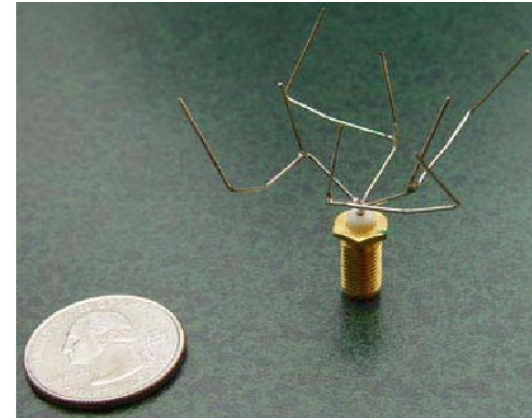
200 FPGA slices; 308 MHz

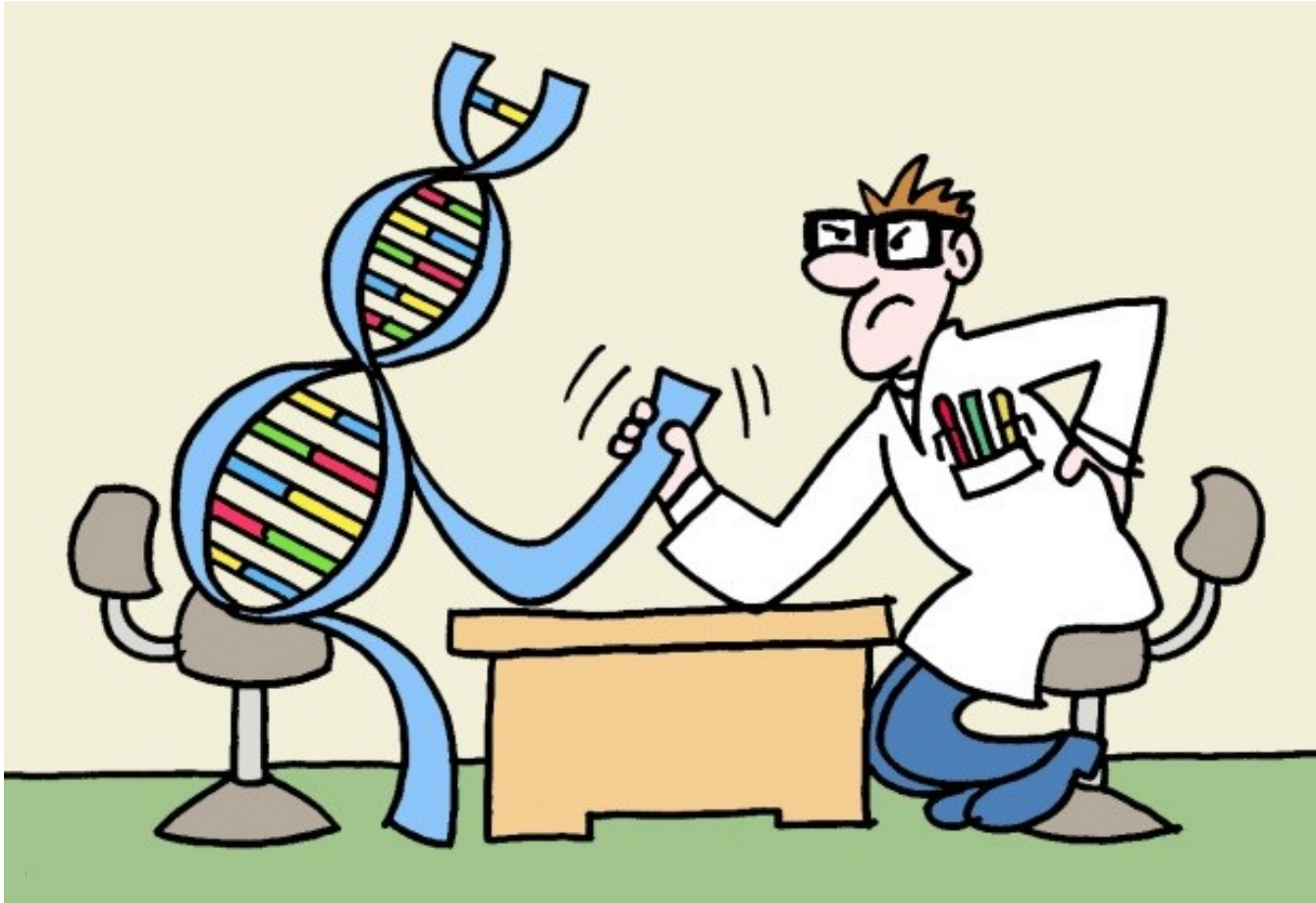


A Czech patent #304181 approved for evolved image filters using CGP ([Sekanina, Vašíček](#)).

| Other successful applications of GP

- Predictors, classifiers, schedulers, controllers, etc.
 - Not as scalable as DNNs, but the resulting solution can be optimized for error as well as the size and other criteria
 - Their behavior can often be explained.
- Game strategies
- Design of DNNs, circuits, regulators, antennas, quantum circuits, antennas, ...
- Automated design of program tests
- Automated software repair
- Genetic improvement of software
- Automated patent invention





Human-Competitive Results Produced By Genetic And Evolutionary Computation
<http://www.human-competitive.org/>

J. Koza: We say that *an automatically created result is “human-competitive” if it satisfies one or more of the eight criteria below.*

- (A) The result was patented as an invention in the past, is an improvement over a patented invention, or would qualify today as a patentable new invention.
- (B) The result is equal to or better than a result that was accepted as a new scientific result at the time when it was published in a peer-reviewed scientific journal.
- (C) The result is equal to or better than a result that was placed into a database or archive of results maintained by an internationally recognized panel of scientific experts.
- (D) The result is publishable in its own right as a new scientific result $\frac{3}{4}$ *independent* of the fact that the result was mechanically created.
- (E) The result is equal to or better than the most recent human-created solution to a long-standing problem for which there has been a succession of increasingly better human-created solutions.
- (F) The result is equal to or better than a result that was considered an achievement in its field at the time it was first discovered.
- (G) The result solves a problem of indisputable difficulty in its field.
- (H) The result holds its own or wins a regulated competition involving human contestants (in the form of either live human players or human-written computer programs).

Humies winners (2004 - 2020)

Year	Humies Winners	Institution
2004	An Evolved Antenna for Deployment on NASA's ST 5 Mission	NASA AMES, US
	Automatic Quantum Computer Programming: A GP Approach	Hampshire Coll., US
2005	Two-dimensional photonic crystals designed by evolutionary algorithms	Cornell U., US
	Shaped-pulse optimization of coherent soft-x-rays	Colorado S. Univ., US
2006	Catalogue of Variable Freq. and Single-Resistance-Controlled Oscillators	MIT, US
2007	Evol. Design of Single-Mode Microstructured Polymer Optical Fibers	UCL, UK
2008	Genetic Programming for Finite Algebras	Hampshire Coll., US
2009	A Genetic Programming Approach to Automated Software Repair	U. of New Mexico, US
2010	Evol. design of the energy function for protein structure prediction	U. of Nottingham, UK
2011	GA-FreeCell: Evolving Solvers for the Game of FreeCell	Ben-Gurion U., IL
2012	Evolutionary Game Design	ICL, UK
2013	Evolutionary Design of FreeCell Solvers	Ben-Gurion U., IL
	Search for a grand tour of the Jupiter Galilean moons	ESA, EU
2014	Genetic Algorithms for Evolving Computer Chess Programs	Bar-Ilan University, IL
2015	Evolutionary Approach to Approximate Digital Circuits Design	Brno U. of Tech., CZ*
2016	Automated Software Transplantation	UCL, UK
2017	Explaining quantum correlations through evolution of causal models	U. of Sydney, AU
2018	A New EA-Based Home Monitoring Device for Parkinson's Dyskinesia	U. of York, UK
2019	Automated Self-Optimization in Heterogeneous Wireless Comm. Networks	UCD, IE
2020	Low cost satellite constellations for nearly continuous global coverage	Aerospace corporation, US

FIT BUT – Silver medal (2008, 2011, 2016), Bronze medal (2014, 2018)

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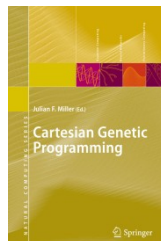
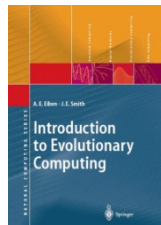
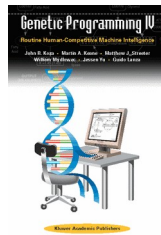
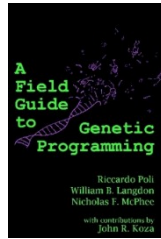
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DEAP: Distributed Evolutionary Algorithms in Python

<https://deap.readthedocs.io/en/master/>



Thank you for your attention!

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BISSIT, Brno, July 22, 2022