



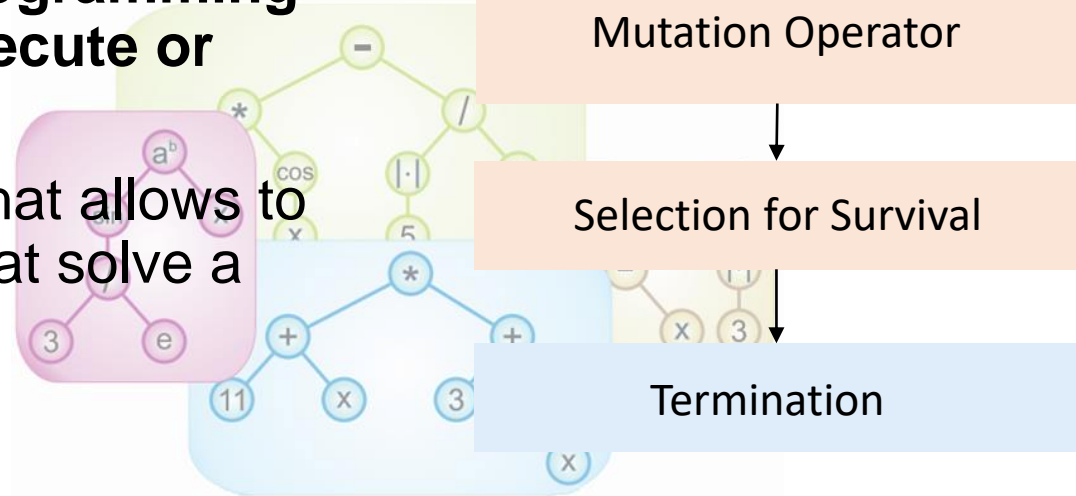
# Genetic Programming

Nuno Antunes Ribeiro

Assistant Professor

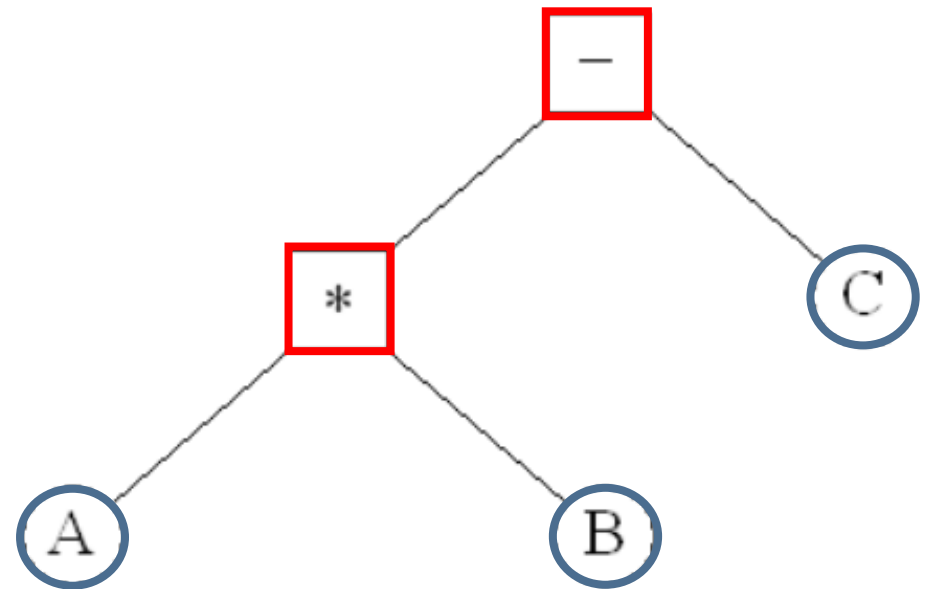
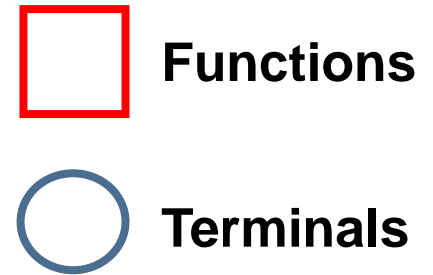
# Genetic Programming

- GP was developed by Jonh Koza (a PhD student of Jonh Holland) around 1992.
- It is a more recent evolutionary approach, which extends the generic model of learning to the space of programs
- Its major variation, with respect to other evolutionary families, is that the evolving individuals are themselves **computer programs (sequence of instructions in a programming language that a computer can execute or interpret)**
- GP is a form of program induction that allows to automatically generate programs that solve a given task



# Tree-based Encoding

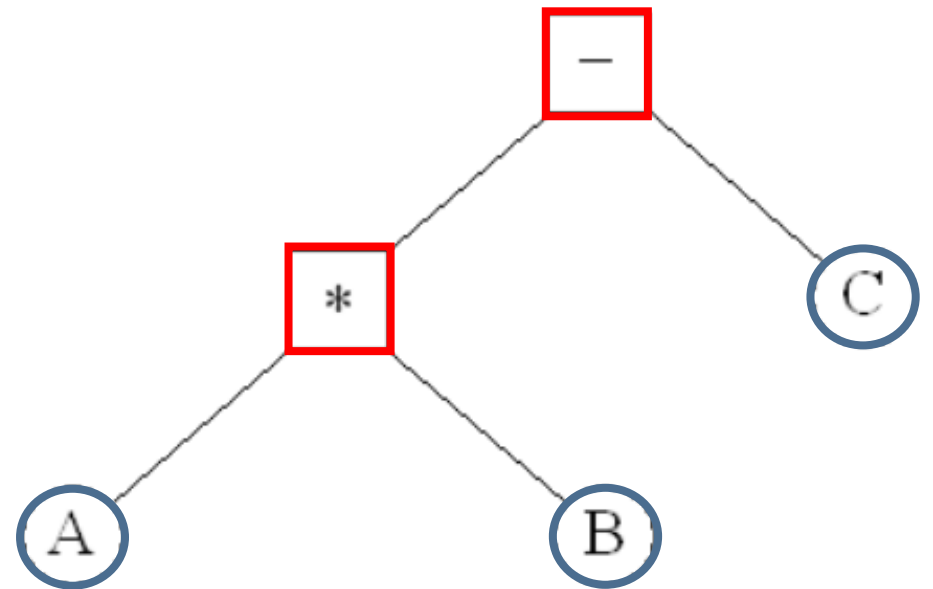
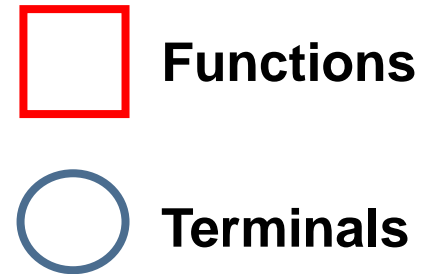
- **Any computer program is a sequence of operations (functions)** applied to values (arguments).
- **Tree encoding** is often used for hierarchical sequenced optimization problems. In tree encoding, a solution is represented by a tree of some operations /functions
- Computer programs can be coded as a hierarchical sequence of functions



# Tree-based Encoding

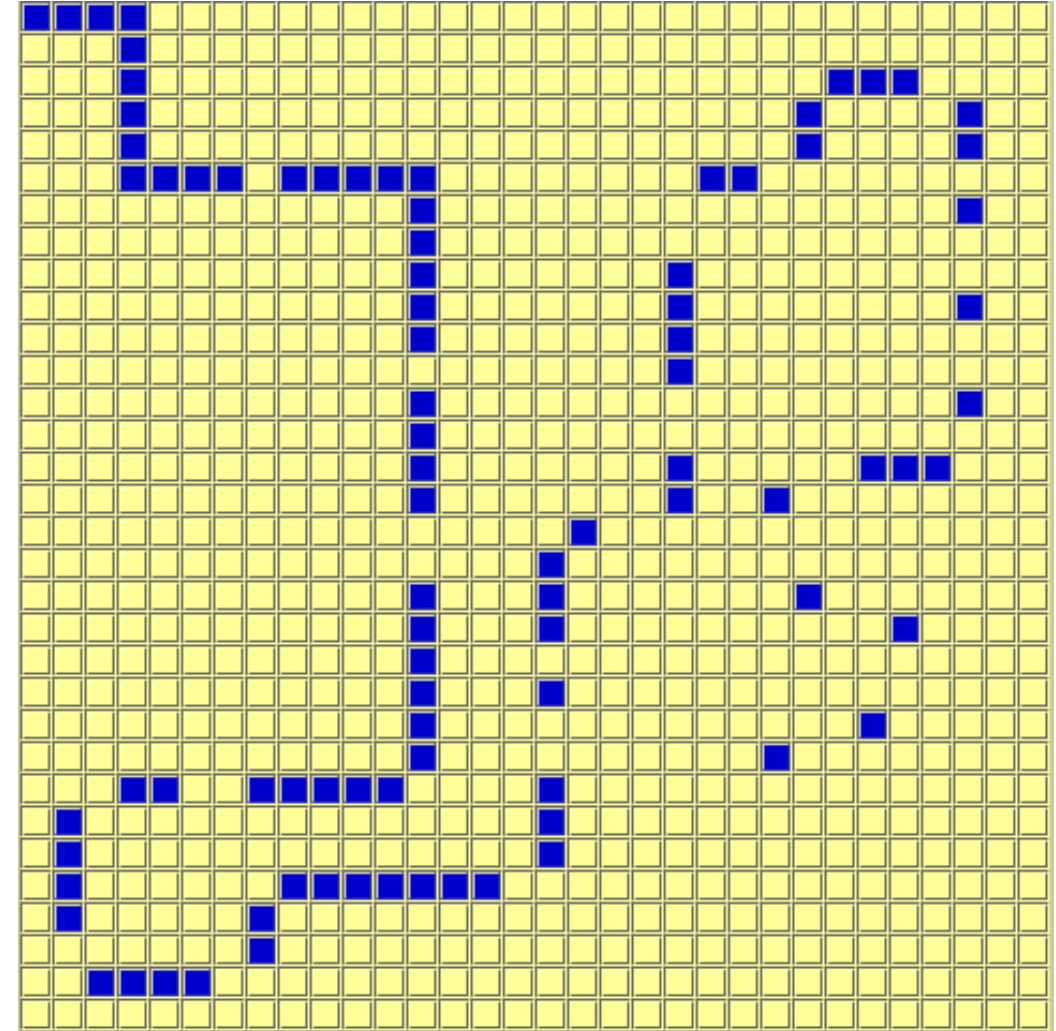
**Example:**  $(- (* A B) C) \longrightarrow (A \times B) - C$

- Calls for the application of the subtraction function (-) to two arguments, namely the list (\*A B) and the atom C.
- But, first, the multiplication function (\*) is applied to A and B.
- Once the list (\*A B) is evaluated, the tree applies the subtraction function (-) to the two arguments, and thus evaluates the entire list



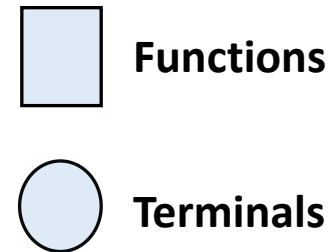
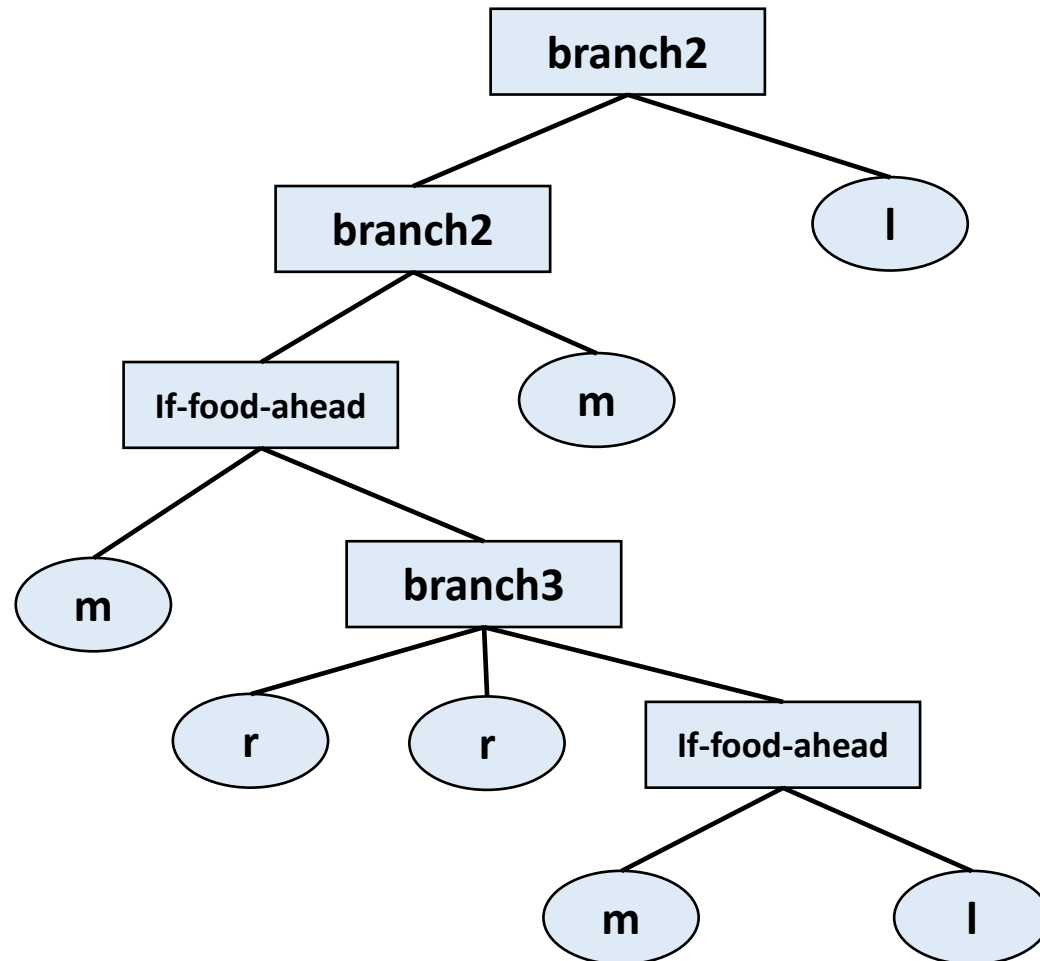
# Santa Fe Ant Trail

- The Santa Fe Trail problem is a genetic programming exercise in which artificial ants search for food pellets according to a programmed set of instructions
- Fitness will be measured by the number of food pellets the ant encounters.
- The basic problem considers 6 operators (3 functions and 2 operators):
  - **If-food-ahead** (if); **branch2** (2 branches); **branch3** (3 branches);
  - Terminal nodes: **move straight** (m); **left** (l); **right** (r)



# Random Tree

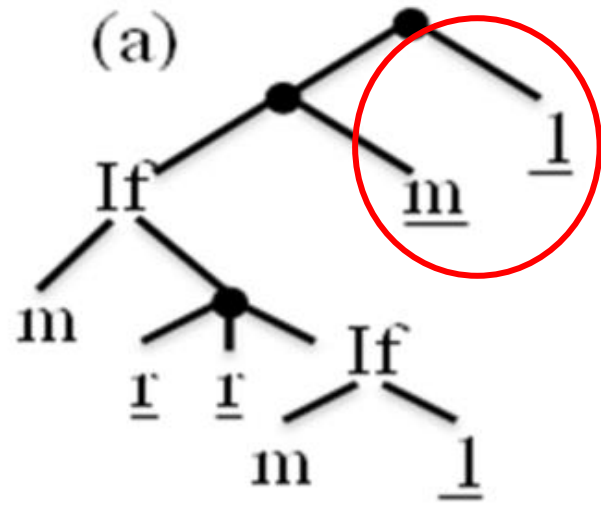
- Let's start by generating a random tree for the problem.



**Vector encoding**  
B2(B2(if(m,B3(r,r,if(m,l)))m)l)



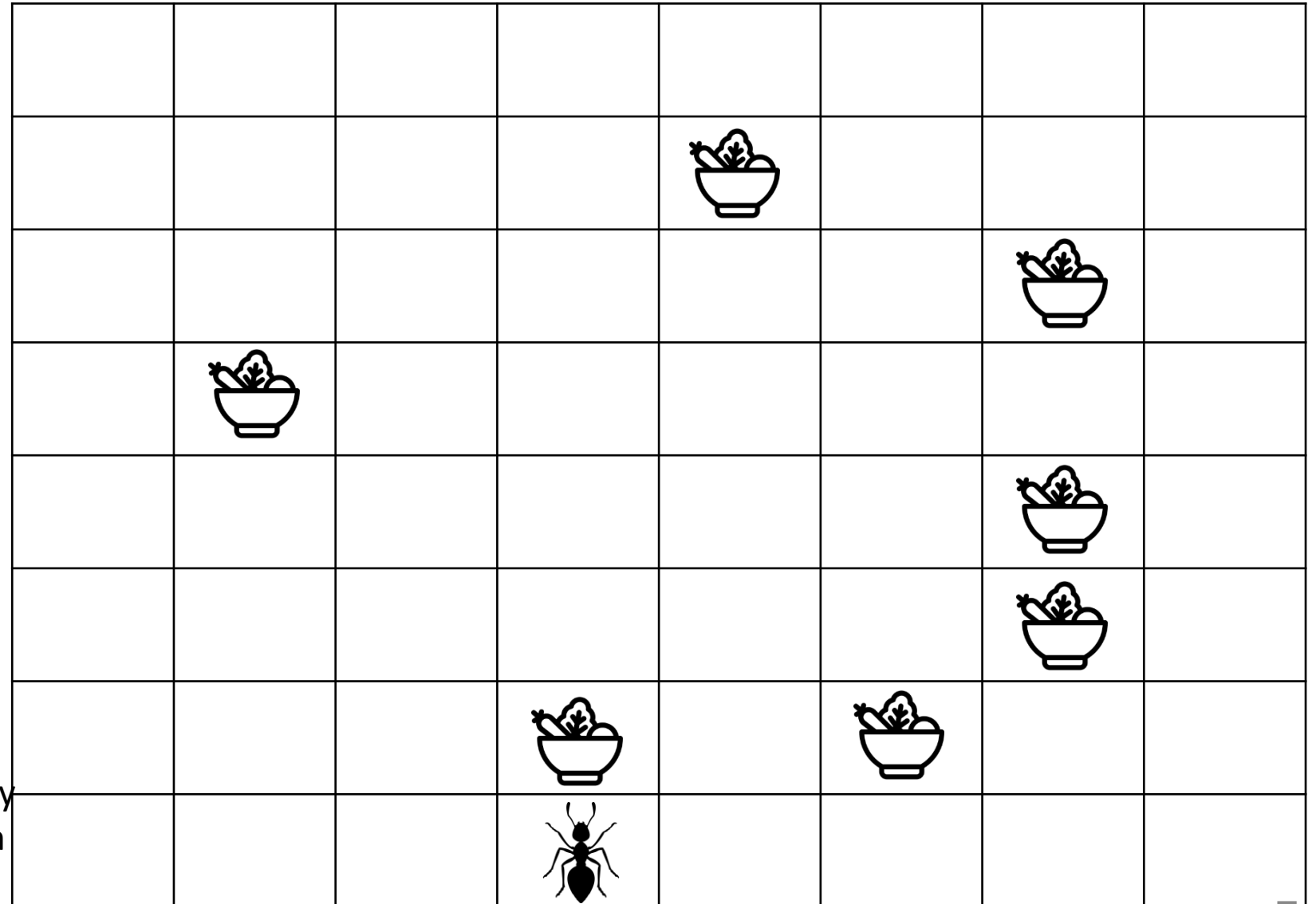
# Santa Fe Ant Trail



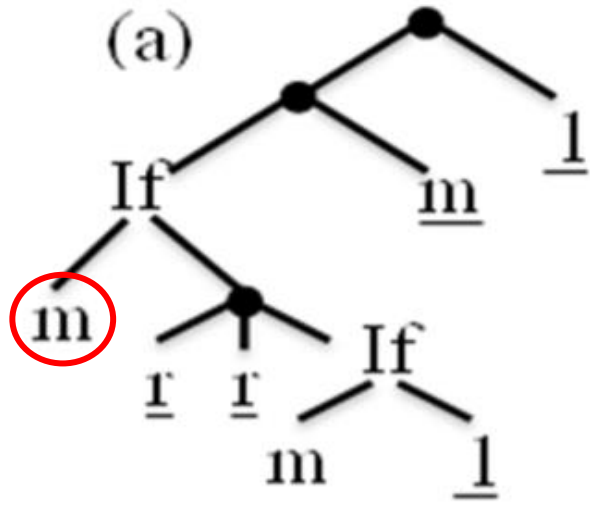
Move:

?ml

m and l moves are always applied. They are the last two moves to be applied in all iterations according to this program

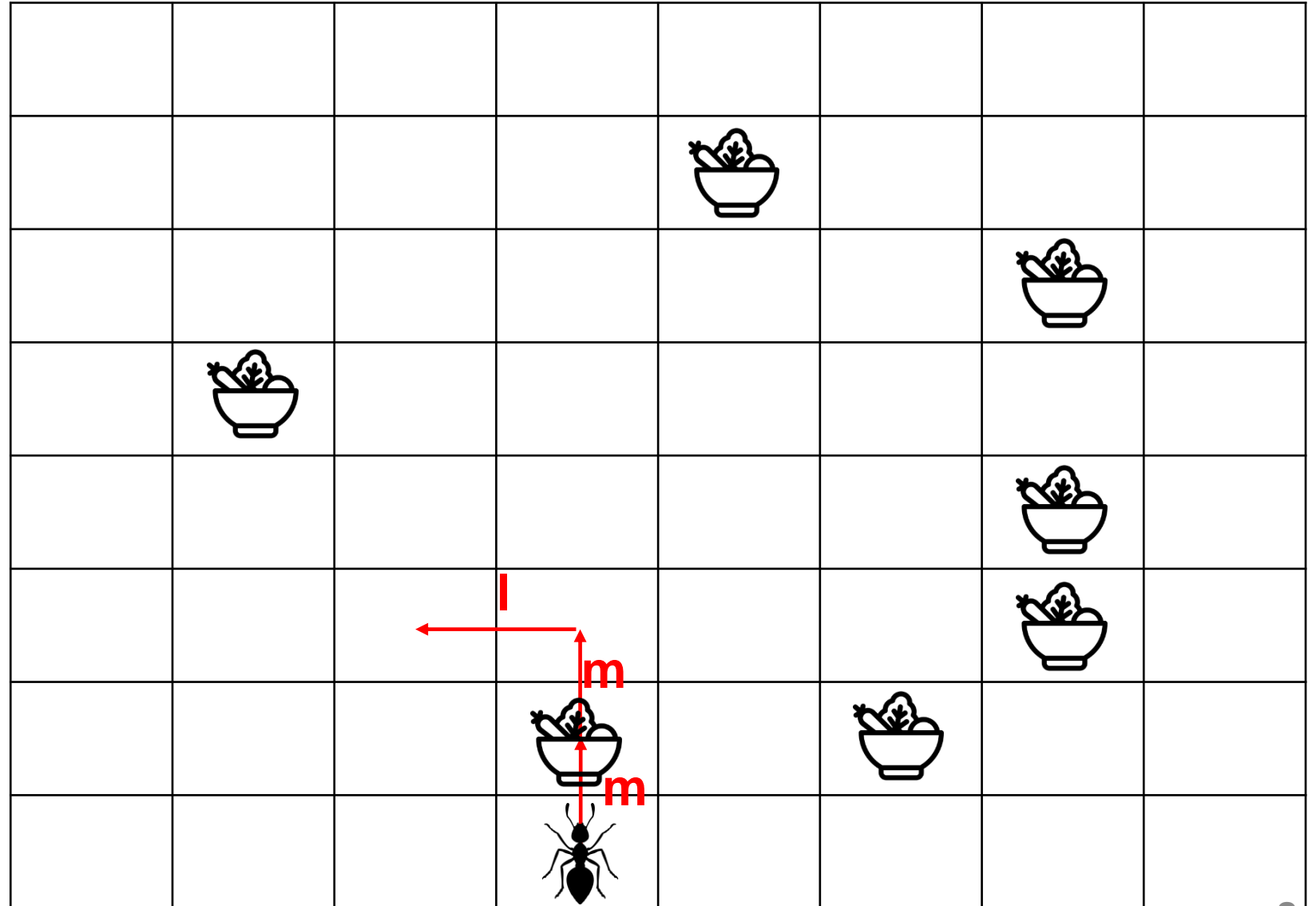


# Santa Fe Ant Trail



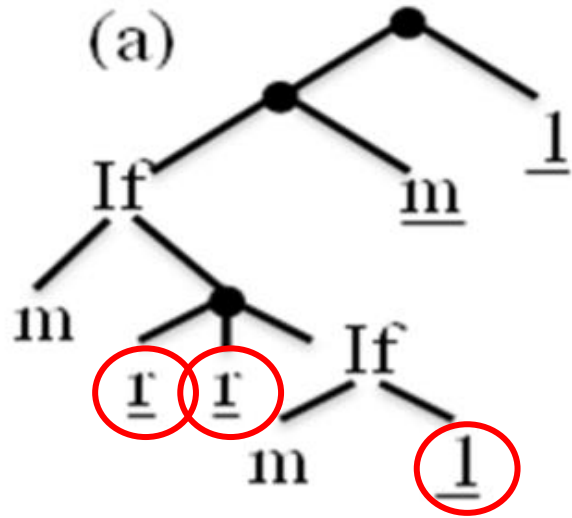
Move:

mml



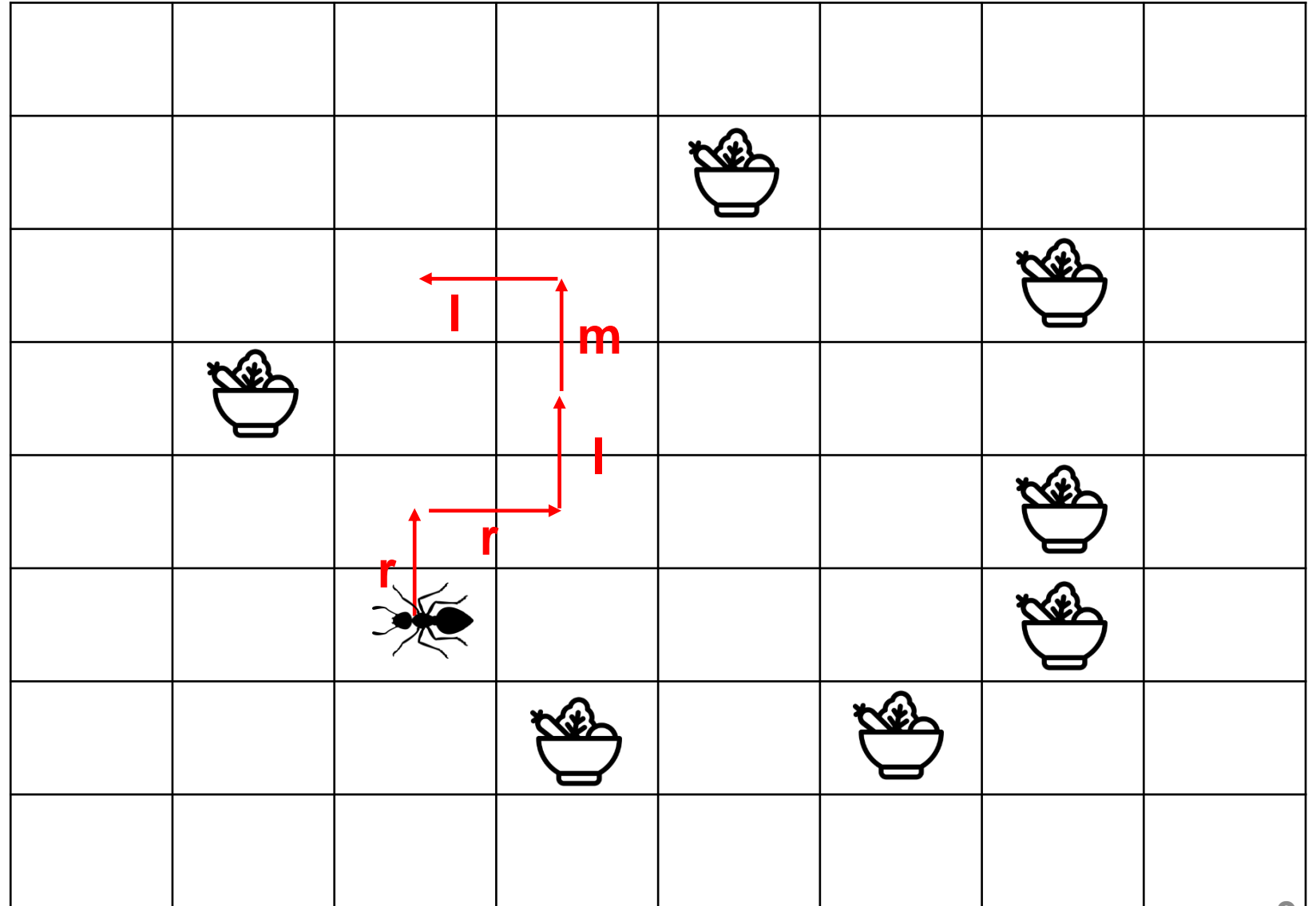


# Santa Fe Ant Trail

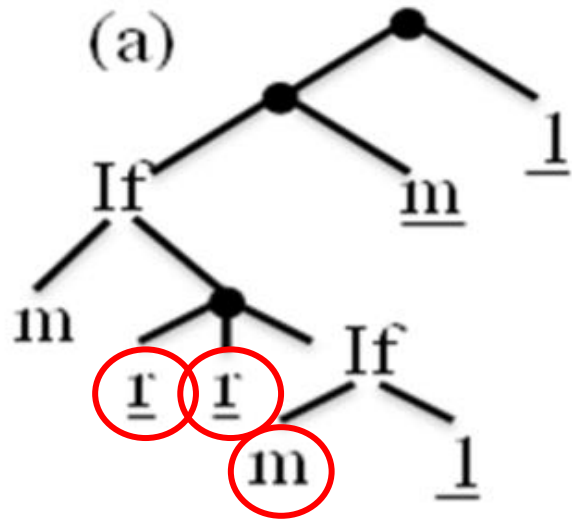


## Move:

rrlml

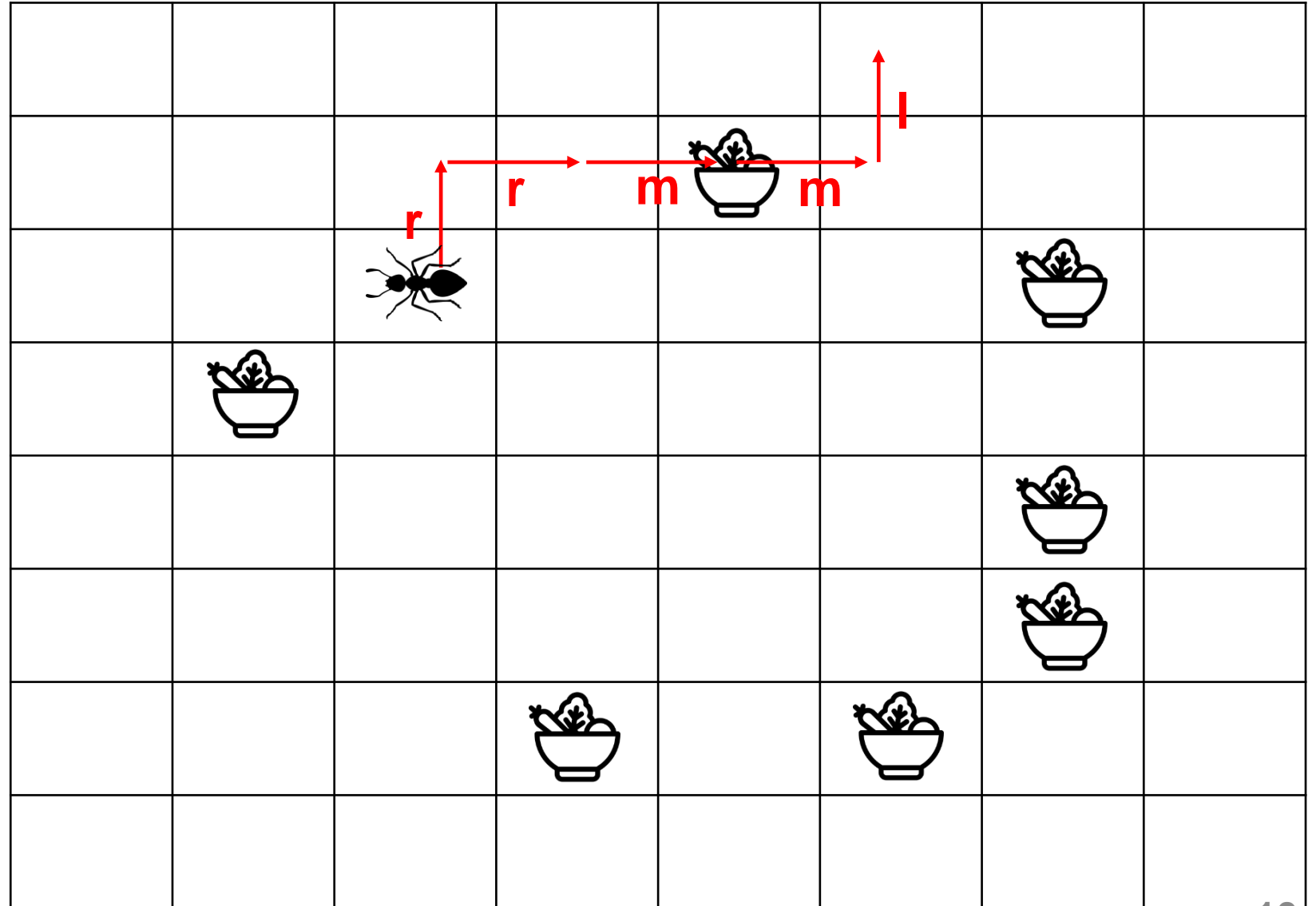


# Santa Fe Ant Trail

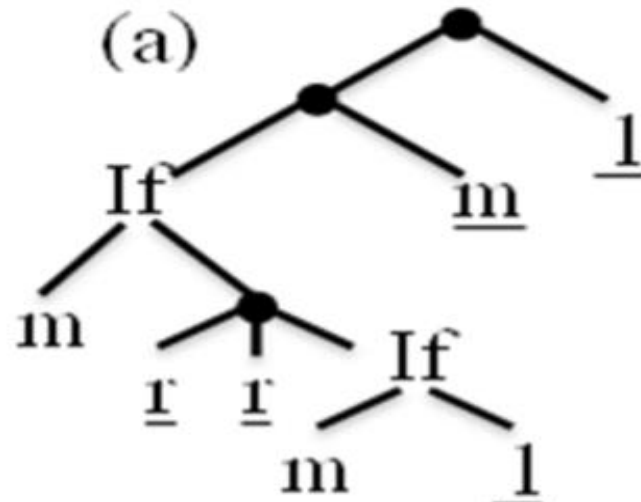


## Move:

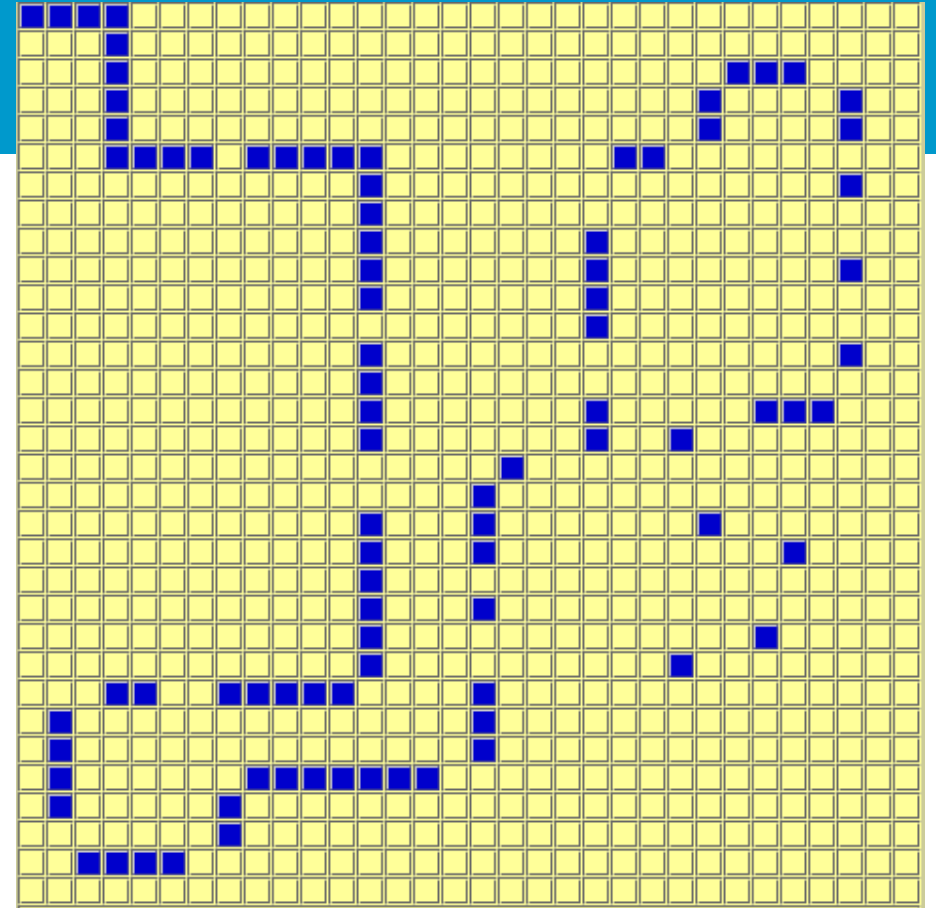
rrmml



# Santa Fe Ant Trail

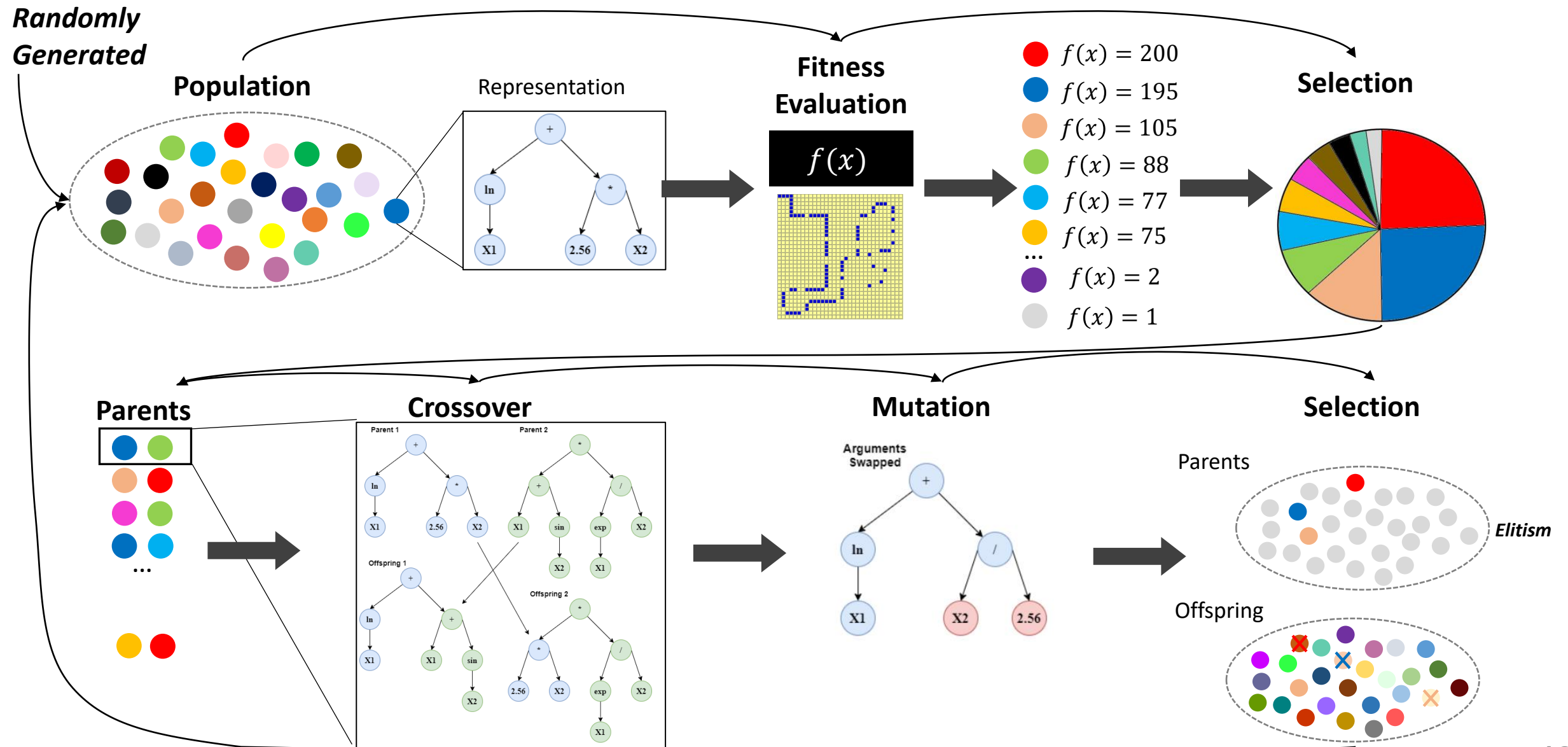


## All the Steps



mm1,rrlml,rrmm1,rrlml,rrlml,rrlml,mm1,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrmm1,rrlml,rrlml,rrlml  
,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrmm1,rrlml,rrlml,  
rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,mm1,rrlml,rrlml,rrlml,rrlml,mm1,rrlml,rrlml,rrlml,rrlml,mm1,r  
rlml,rrmm1,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,mm1,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml  
l,rrlml,rrlml,rrmm1,rrlml,rrlml,mm1,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrmm1,rrlml,rrlml,m  
ml,rrlml,rrmm1,rrlml,rrlml,rrlml,rrmm1,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrlml,rrmm  
l,rrlml,rrlml,rrlml,rrlml,mm1,rrlml,rrlml,mm1,rrlml,rrlml,rrmm1,rrlml,rrlml,rrmm1,rrlml,rrlml,mm1

# Evolve our Programme using GP



# Crossover

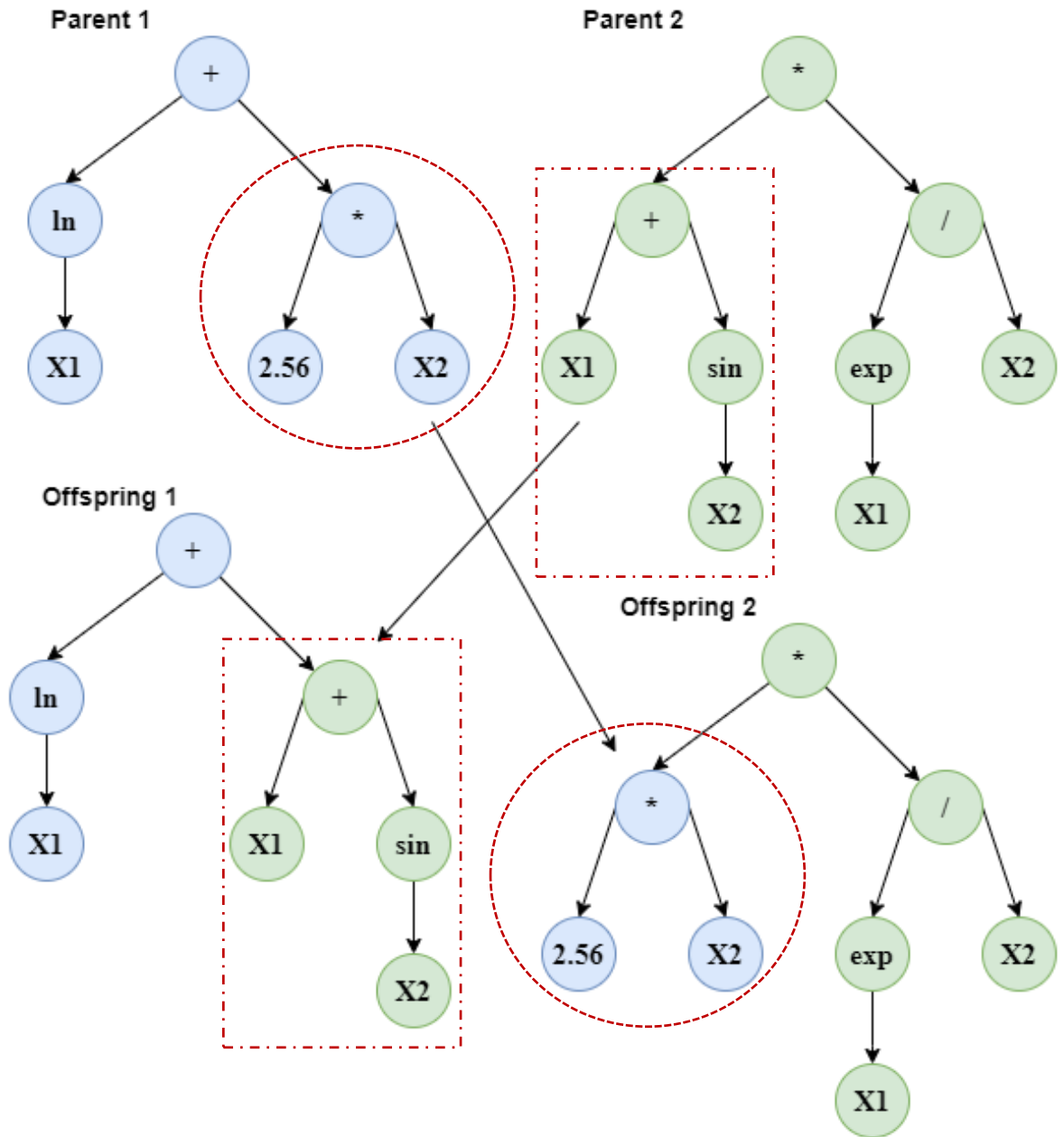
- Crossover operators in genetic programming are quite intuitive. As for other evolutionary algorithms, given two parents, we randomly select a point in each parent and crossover that subtree at that point to create an offspring.

**Parent 1**  $f(x_1, x_2) = \ln(x_1) + (2.56 \times x_2)$

**Parent 2**  $f(x_1, x_2) = (x_1 + \sin(x_2)) \times \left(\frac{e^{x_1}}{x_2}\right)$

**Child 1**  $f(x_1, x_2) = (2.56 \times x_2) \times \left(\frac{e^{x_1}}{x_2}\right)$

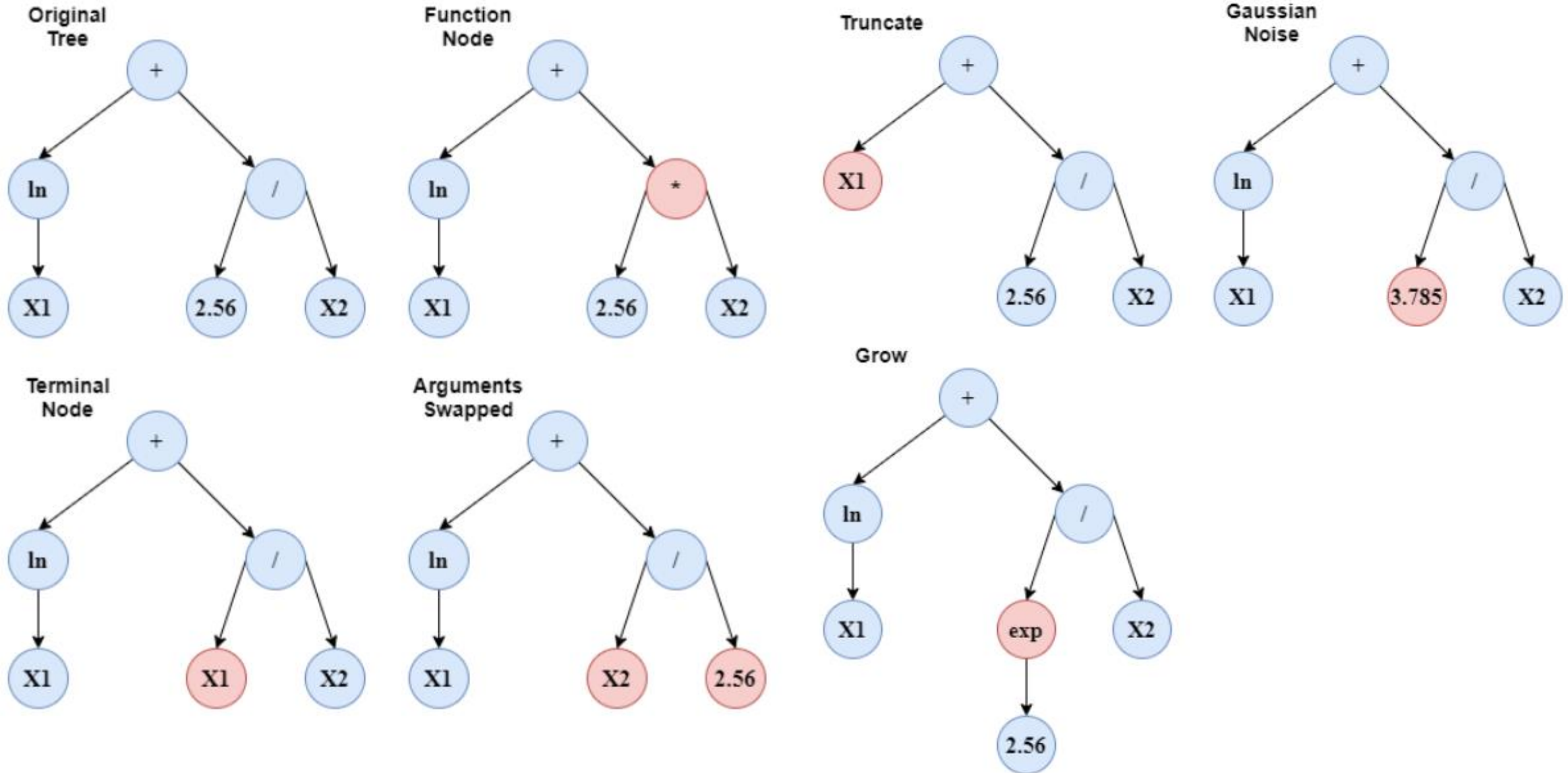
**Child 2**  $f(x_1, x_2) = \ln(x_1) + (x_1 + \sin(x_2))$



# Mutation

- Like crossover, mutation for genetic programming is extremely intuitive, and similar to other evolutionary algorithms.
  - **Function node switching** - switch a random function node to another viable node.
  - **Terminal node switching** - switch a random terminal node with another viable node
  - **Swapping arguments of a terminal** – swap to terminal nodes
  - **Gaussian Noise** - add random gaussian noise to existing numeric values.
  - **Grow** - grow our trees by randomly introducing a single new function node.
  - **Truncation** - shrink a tree by randomly deleting a single function node

# Mutation





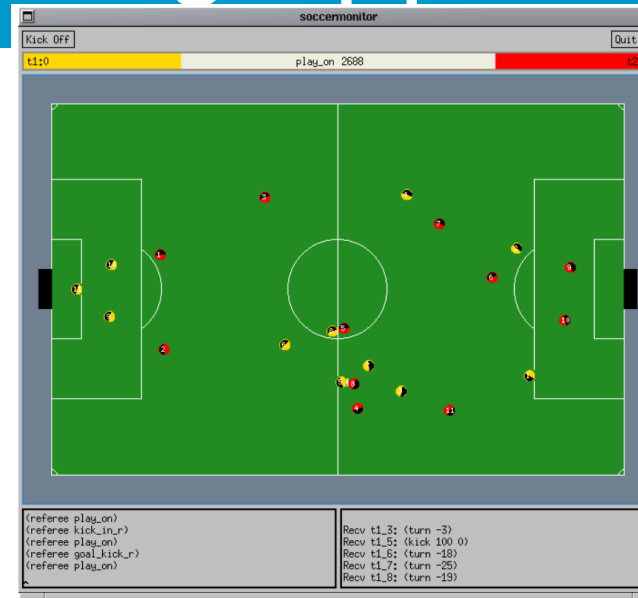
# Santa Fe Ant Trail

```
40: 4 ####
41: 6 #####
42: 3 ###
43: 1 #
44: 7 #####
45: 2 ##
46: 5 #####
47: 2 ##
48: 3 ###
49: 6 #####
50: 1 #
51: 2 ##
53: 4 ####
54: 1 #
55: 1 #
56: 1 #
57: 1 #
58: 3 ###
59: 1 #
60: 1 #
61: 1 #
62: 1 #
64: 2 ##
66: 1 #
67: 1 #
68: 1 #
71: 1 #
78: 1 #
85: 1 #
Average score: 4.576
```

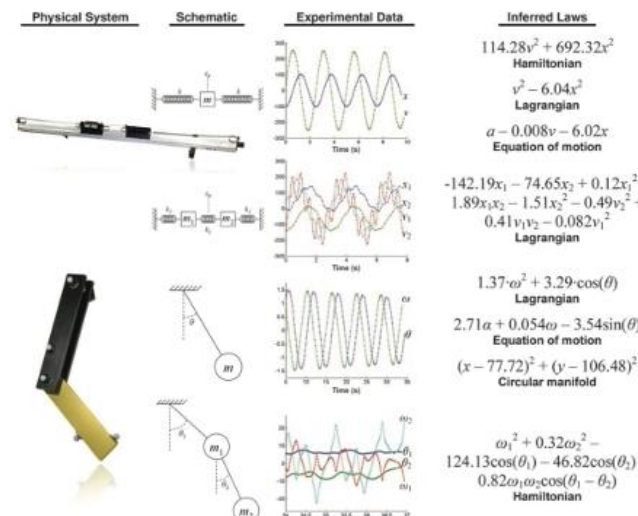
<https://www.youtube.com/watch?v=InpbbgpDQkg>

# Genetic Programming Applications

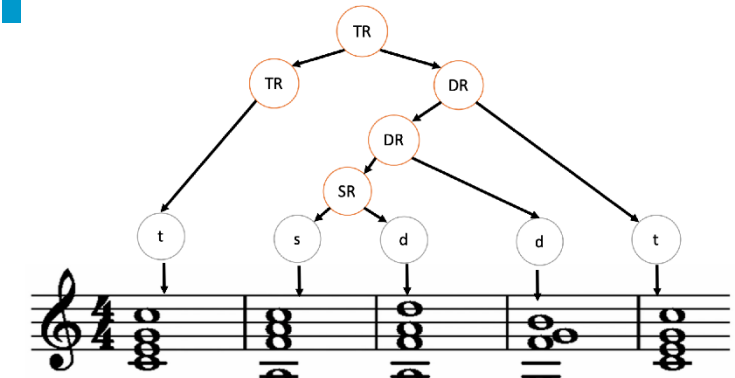
- Robotics ; Game Playing ; Control
- **Machine Learning (Regression and Classification Problems)**
- Maths and Physics
- Systems Security
- Economy and Finances
- Music Creation
- Video Editing
- Etc.



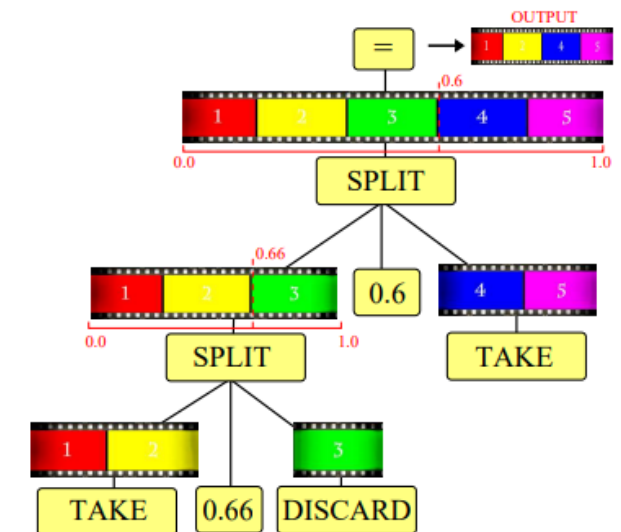
Source: <http://www.genetic-programming.com/hc/andretellersoccer.html>



Source: <https://www.science.org/doi/10.1126/science.1165893>



Source: <https://www.mdpi.com/2076-3417/10/17/6039/htm>



Source:

<https://citeseerx.ist.psu.edu/viewdoc/download?doi=10.1.1.370.9167&rep=rep1&type=pdf>



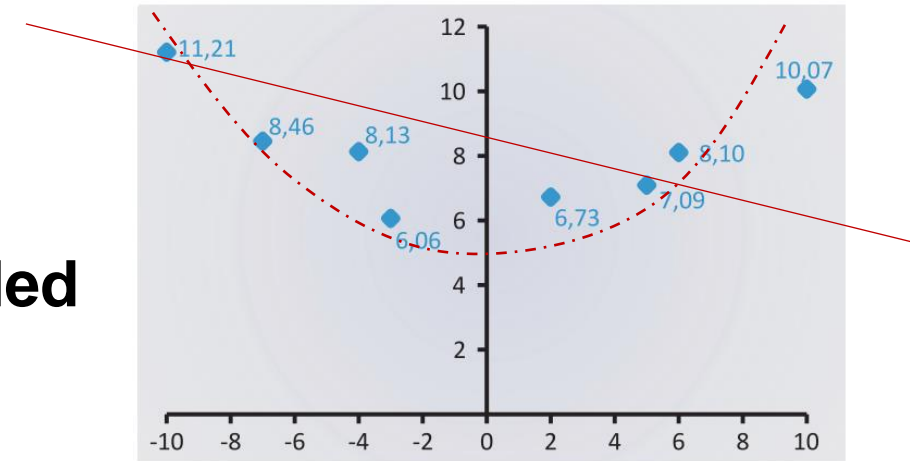
# Symbolic Regression

Nuno Antunes Ribeiro

Assistant Professor

# Genetic Programming - Symbolic Regression

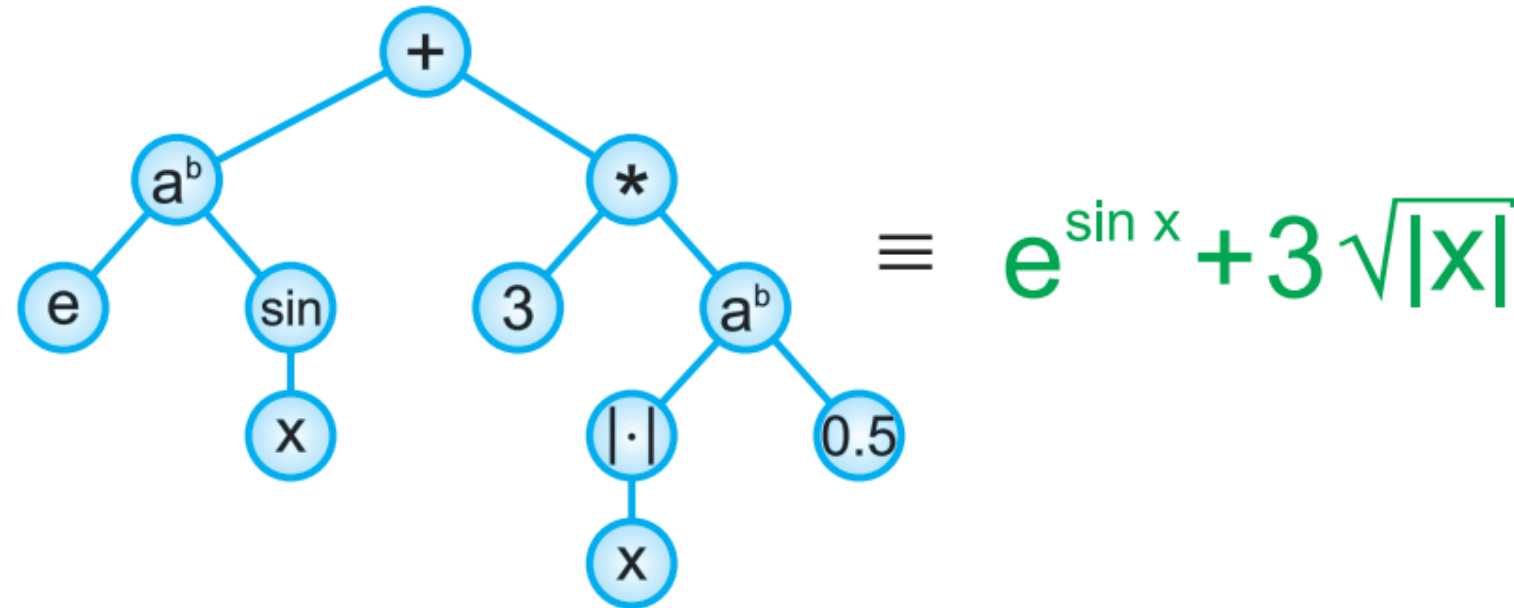
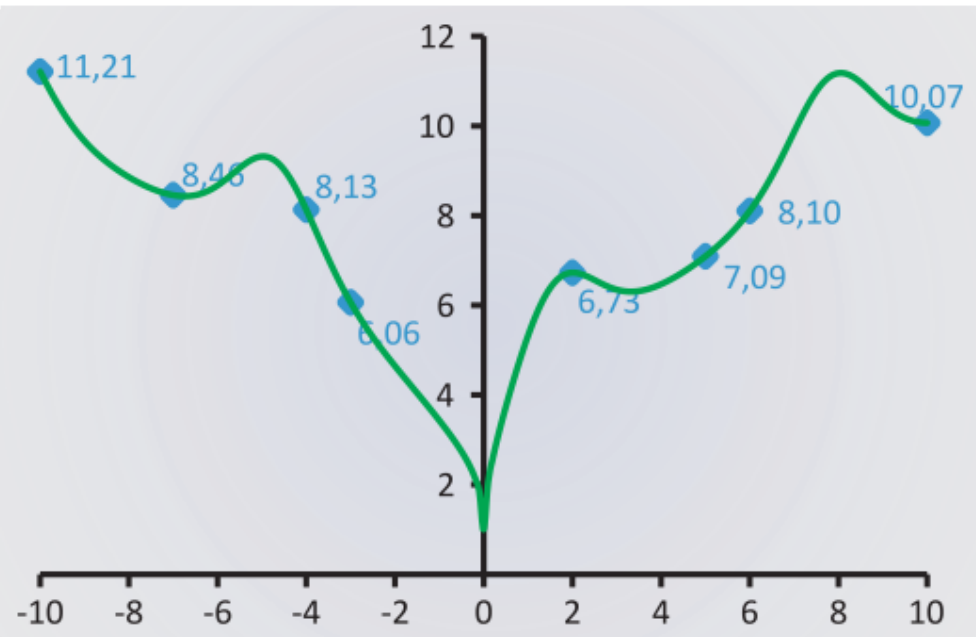
- Linear Regression
- Other regression methods
  - **Assumption about formula blueprint needed**



- **Symbolic regression** - new kind of optimization problem:
- We have
  - Given set of points (observations) -  $S = \{(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)\}$
  - A Function Set with elementary functions (e.g.  $F = \{+, -, \times, \sqrt{\phantom{x}}, \sin, \exp, \dots\}$ )
  - A Terminal Set, i.e. input variables (e.g.  $T = \{x \text{ and real constants}\}$ )
- We want to find the best formulation (combining  $F$  and  $T$ ) that best fits the observations  $S$

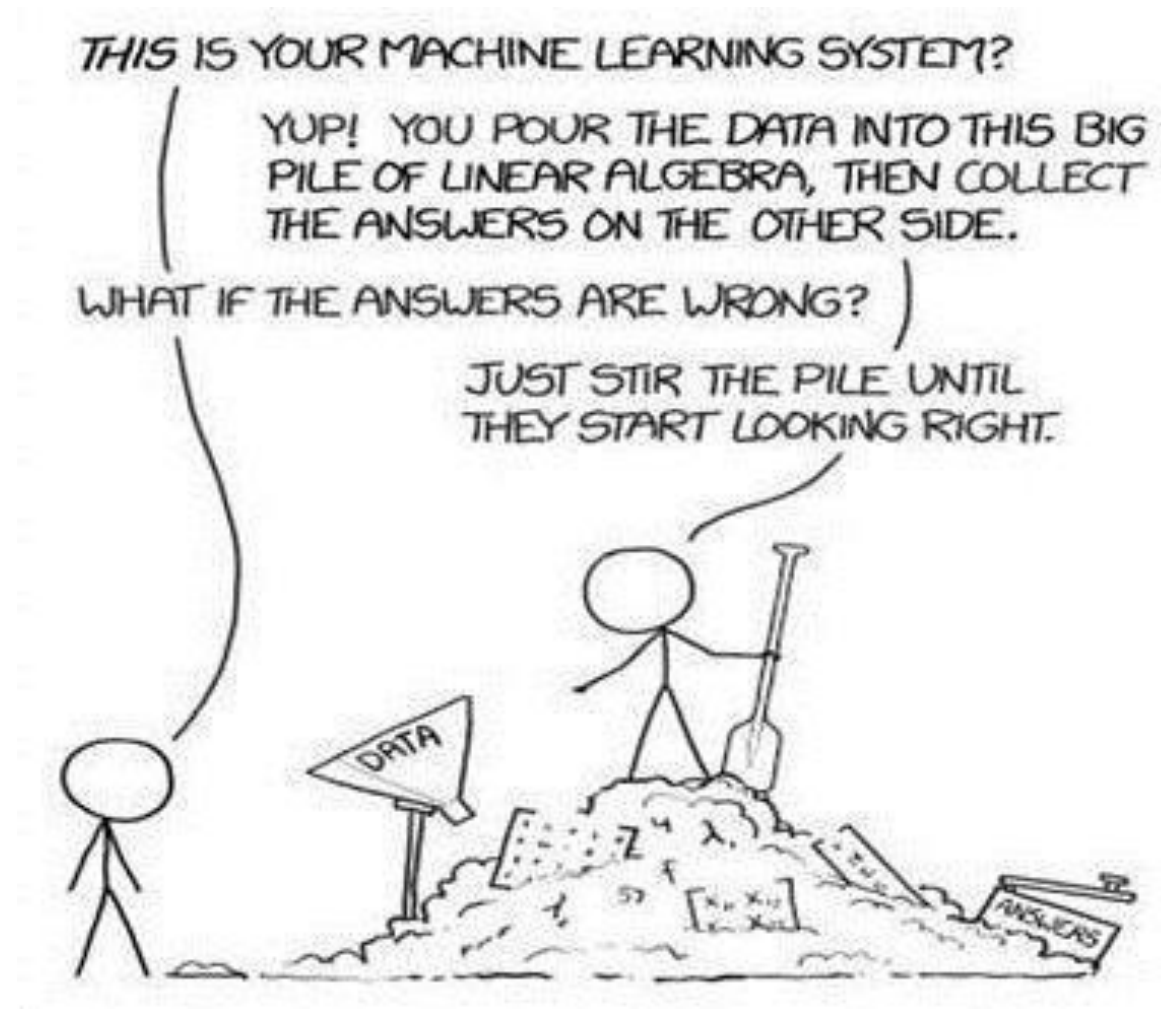
# Genetic Programming - Symbolic Regression

- Symbolic Regression with Genetic Programming:
  - represent formulas as tree data structures
  - Evaluation function: minimize  $\sum_{i=1}^n (y_i - f(x_i))^2$
  - Construct  $f(x)$  with Genetic Programming!



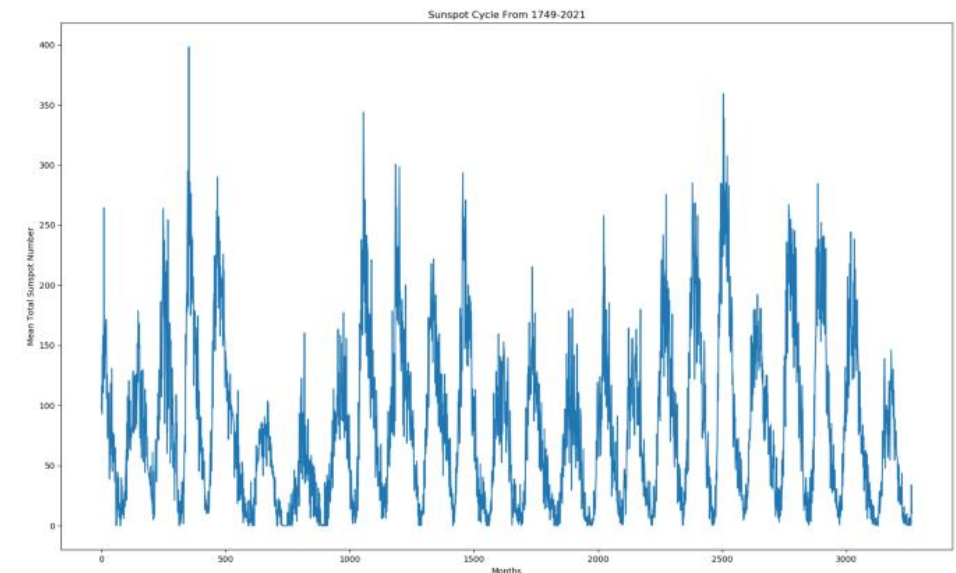
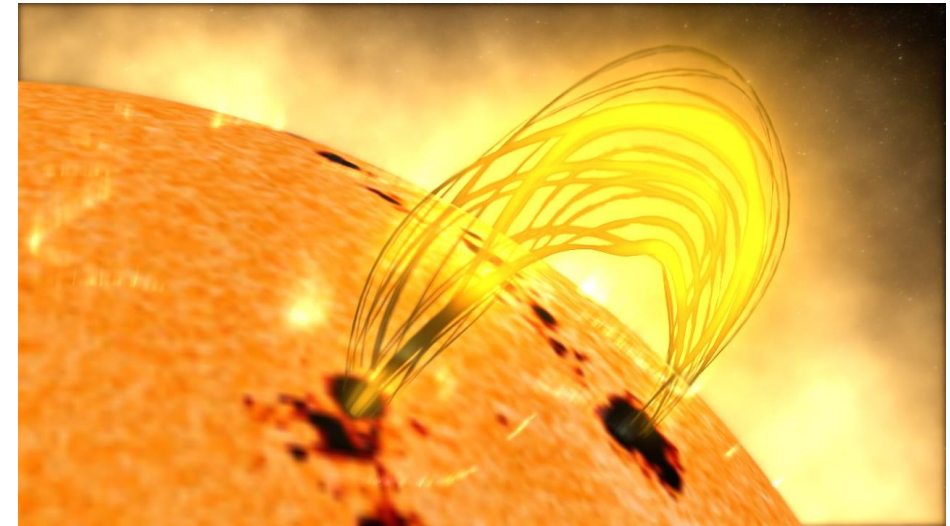


# Genetic Programming Algorithm



# Symbolic Regression in Python

- Sunspots have been observed for over four centuries, constituting the longest running, continuous time series of any natural phenomena in the Universe.
- Sunspots spawn severe space weather characterized by solar flares, coronal mass ejections, geomagnetic storms, enhanced radiative, and energetic particle flux
- Endangering satellites, global communication systems, air-traffic overpolar routes, and electric power grids
- Protection of planetary technologies and space situational awareness is therefore enabled by solar activity predictions.



Sunspots.csv



# Symbolic Regression in Python

- Programming a genetic programming model from scratch requires a lot of extracurricular preliminaries, such as automata theory, I will not be performing the algorithm from scratch.
- Instead, I will be using **gplearn**, a free python library designed specifically for genetic programming algorithms for both classification and regression.



Genetic Programming in Python,  
with a scikit-learn inspired API:

**gp**learn

<https://github.com/trevorstephens/gplearn>

# Symbolic Regression in Python

```
import numpy as np
from sklearn.metrics import mean_squared_error
import pandas as pd
import matplotlib.pyplot as plt
from gplearn.genetic import SymbolicRegressor
from sympy import *

#conda install python-graphviz

seed=2

np.random.seed(seed)

df = pd.read_csv("Sunspots.csv")
y = np.asarray(df['Monthly Mean Total Sunspot Number'])
size = len(y)
# 50% of data for training
train_ind = int(size * 0.50)
# 25% of data for validation and other 25% for testing
val_ind = int(size * 0.75)


# testing Genetic Programming algorithm using gplearn

max_window = 8
min_window = 3

best_models = [] # best model from each run of the algorithm per window size
best_fits = []
# randomly shuffle data through indices:
shuffled_indices = np.asarray(range(0, size-max_window))
np.random.shuffle(shuffled_indices)
# loop over each window size
```



Import gplearn package



We split our data into the **training, validation, and testing** sets. Our algorithms will be trained using the training dataset. The validation will be used to compare our models. Lastly, after we've chosen our final model, we will evaluate it's accuracy through the test dataset. See Slide 36 – Lecture 13

# Symbolic Regression in Python

```
# Loop over each window size
for vision in range(min_window, max_window + 1):
    input = []
    output = []
    # creates the window length size for each value
    # because the first couple values will not have
    # a full window we skip them, that's why start
    # at i and not 0
    for j in range(vision, size):
        input.append(y[(j - vision):j].tolist())
        output.append(y[j])

    input = np.asarray(input)
    output = np.asarray(output)

    temp = np.column_stack((output, input))

    # instead of shuffle each time here, we shuffle once outside loop
    # so that all window sizes have the same final array
    temp = temp[shuffled_indices]

    output = temp[:, 0]
    input = temp[:, 1:]

    y_train = output[0:train_ind]
    y_val = output[train_ind:val_ind]
    y_test = output[val_ind:size]
    x_train = input[0:train_ind]
    x_val = input[train_ind:val_ind]
    x_test = input[val_ind:size]
```

Loop throughout different window sizes (feature selection)

Prepare explanatory data

Split data into **training, validation, and testing** sets.

```
function_set = ['add', 'sub', 'mul', 'div']
temp_val = []
temp_models = []

for i in range(0, 3):
    gp = SymbolicRegressor(population_size=500, metric='mse',
                           generations=20, init_depth=(2, 6),
                           verbose=1, function_set=function_set, parsimony_coefficient=0.4)

    gp.fit(x_train, y_train)
    predictions = gp.predict(x_val)
    predictions = np.where(predictions < 0, 0, predictions)
    mse1 = mean_squared_error(y_val, predictions)
    print(" MSE Val: " + str(mse1))
    temp_val.append(mse1)
    temp_models.append(gp)

best_index = np.argmin(temp_val)
best_models.append(temp_models[best_index])
best_fits.append(temp_val[best_index])
```

Apply Genetic Programming (see next slide)

# Symbolic Regression in Python

```
function_set = ['add', 'sub', 'mul', 'div']
temp_val = []
temp_models = []
for i in range(0, 3):
    gp = SymbolicRegressor(population_size=500, metric='mse',
                           generations=20, init_depth=(2, 6),
                           verbose=1, function_set=function_set, parsimony_coefficient=0.4)

    gp.fit(x_train, y_train)
    predictions = gp.predict(x_val)
    predictions = np.where(predictions < 0, 0, predictions)
    mse1 = mean_squared_error(y_val, predictions)
    print(" MSE Val: " + str(mse1))
    temp_val.append(mse1)
    temp_models.append(gp)
best_index = np.argmin(temp_val)
best_models.append(temp_models[best_index])
best_fits.append(temp_val[best_index])
```

Set of Functions

Basic gplearn  
parameters  
(symbolic regression)

Compute MSE

We generate 3 programmes (i.e. we apply GP three times) to avoid getting stuck in local optima – multistart approach  
We then select the best programme from the three to test using the validation data

# Symbolic Regression in Python

Population Average			Best Individual			
Gen	Length	Fitness	Length	Fitness	OOB Fitness	Time Left
0	34.16	1.15449e+42	5	748.416	N/A	6.71s
1	11.32	4.13481e+10	5	748.416	N/A	4.76s
2	4.57	4.27409e+06	7	742.902	N/A	4.15s
3	4.27	4997.45	7	742.902	N/A	3.80s
4	5.90	5845.34	9	741.201	N/A	3.70s
5	6.44	6.13391e+07	13	707.885	N/A	3.45s
6	8.22	5.68907e+07	15	706.454	N/A	3.32s
7	11.74	7.37752e+07	27	668.907	N/A	3.13s
8	13.64	3.09287e+07	25	668.894	N/A	2.97s
9	18.06	5.05252e+07	21	663.364	N/A	2.80s
10	27.14	2.80803e+07	23	658.551	N/A	2.79s
11	25.20	2.37904e+07	23	658.551	N/A	2.41s
12	24.99	3.74852e+13	21	658.327	N/A	2.34s
13	22.15	1.08825e+07	21	658.327	N/A	1.76s
14	19.26	4.72406e+07	21	658.327	N/A	1.42s
15	18.78	2.39752e+07	21	658.327	N/A	1.12s
16	18.12	2.42965e+07	21	658.217	N/A	0.85s
17	18.49	5.56e+07	19	658.01	N/A	0.57s
18	16.93	3.17151e+07	19	658.01	N/A	0.28s
19	16.97	3.03626e+07	17	649.785	N/A	0.00s

MSE Val: 625.7448334070784

 Gplearn output

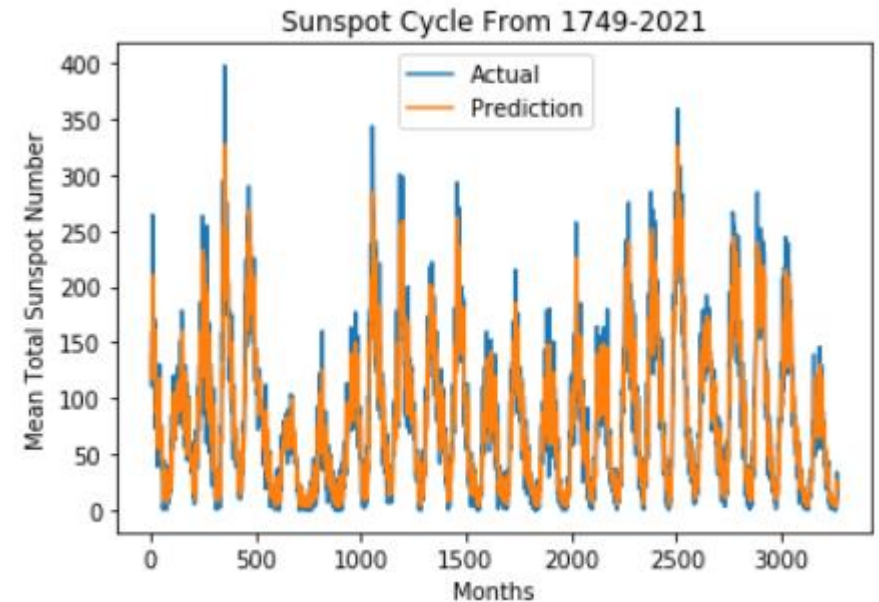
GP estimation of the time left  
to complete the calibration

 MSE for the  
validation data

The length of a program is the number of elements in  
the formula which is equal to the total number of  
nodes

# GP Application - Time Series Analysis

Best Validation Fitness Values Per Window Size:  
Window Size: 3 - Validation MSE: 707.5851095958388  
Window Size: 4 - Validation MSE: 625.7448334070784  
Window Size: 5 - Validation MSE: 603.3792897649598  
Window Size: 6 - Validation MSE: 695.9561339158797  
Window Size: 7 - Validation MSE: 705.650323050849  
Window Size: 8 - Validation MSE: 730.6631145004401  
Validation Error: Mean w/ std: 678.1631340391745+-46.60890223275122  
Best Model:  
Window Size : 5  
MSE for Test Data Set : 667.0923572895032



```
print(best_model._program)
```

```
add(div(sub(X2, X4), sub(sub(0.492, div(0.604, -0.420)), div(0.604, -0.420))), X4)
```

# Results

Genetic Algorithms + NN	Backpropagation + NN	Genetic Programming
Testing	Testing	Testing
613	615	608
674	661	667
659	601	642
700	734	645
626	636	615
705	657	751
670	693	694
677	689	704
607	662	787
576	574	559

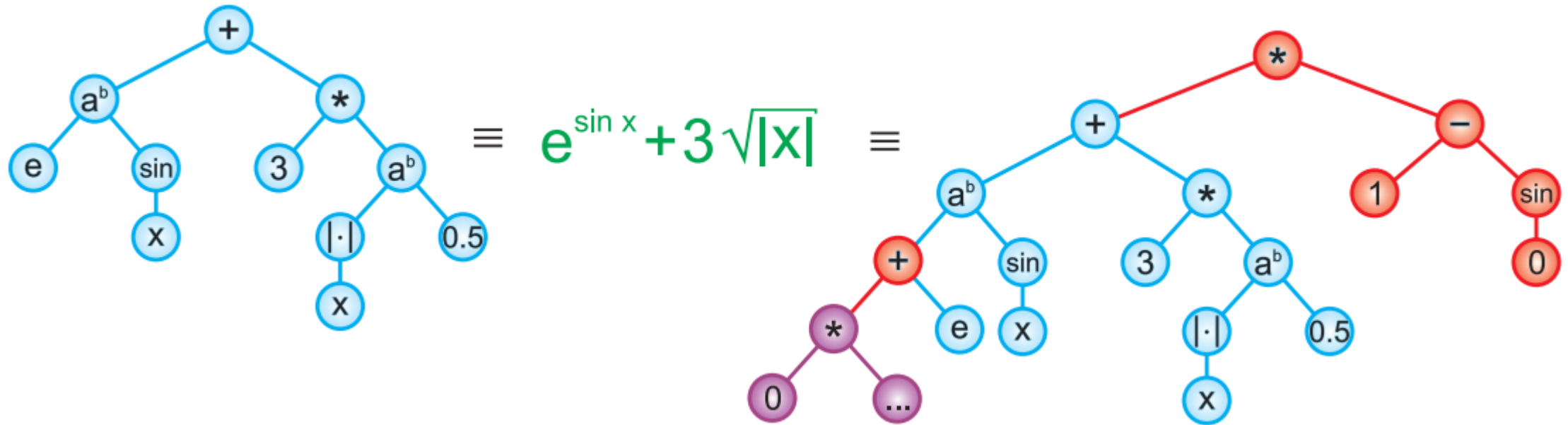
Symbolic regression is a great tool to be aware of.

It is not perfect for every kind of approach, but it gives you another ML option which can be really useful as the outcome is readily understandable.



# Bloat in Genetic Programming

- Bloat: uncontrolled growth of programs
- Intron: useless part of program, one type of bloat



# Bloat in Genetic Programming

- Bloat: uncontrolled growth of programs
- Why is it bad?
  - Elegant solutions are always simple and small
  - Larger programs = longer processing time for both, reproduction operations and evaluation
  - Larger programs = danger of overfitting
  - Larger programs occupy more memory
- What can we do against it?
  - Use multi-objective optimization: minimize also program size
  - Use **penalties in single-objective optimization**
  - Set a conservative upper bound for program size
  - Use specialized mutation and crossover operators which minimize bloat

# Bloat in Genetic Programming

- Bloat can be fought in gplearn in several ways. The principal weapon is using a penalized fitness measure during selection where the fitness of an individual is made worse the larger it is.
- In this way, should there be two programs with identical fitness competing in a tournament, the smaller program will be selected and the larger one discarded.
- The `parsimony_coefficient` parameter controls this penalty and may need to be experimented with to get good performance.



```
gp = SymbolicRegressor(population_size=500, metric='mse',  
                       generations=20, init_depth=(2, 6),  
                       verbose=1, function_set=function_set, parsimony_coefficient=0.4)
```



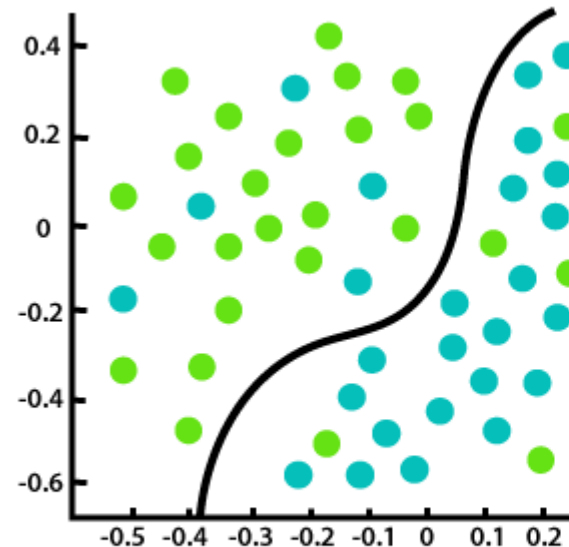
# Solving Classification Problems using Genetic Programming

Nuno Antunes Ribeiro

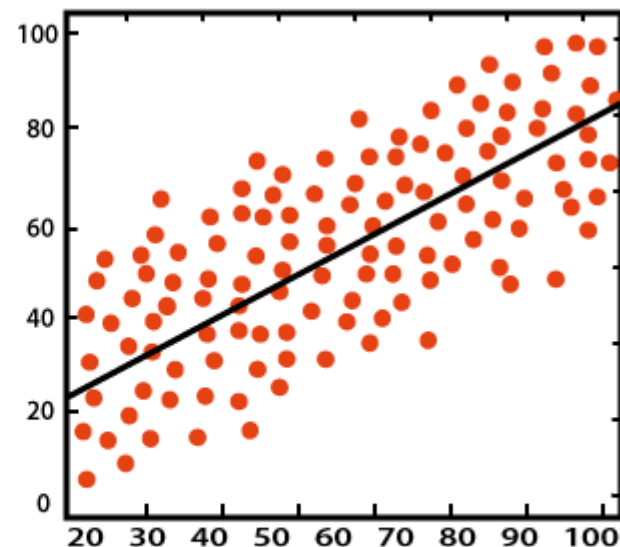
Assistant Professor

# Classification Problems

- Classification is a task that requires the use of machine learning algorithms that learn how to assign a class label to examples from the problem domain. An easy to understand example is classifying emails as “spam” or “not spam.”
- There are many different types of classification tasks that you may encounter in machine learning and specialized approaches to modeling that may be used for each.



Classification

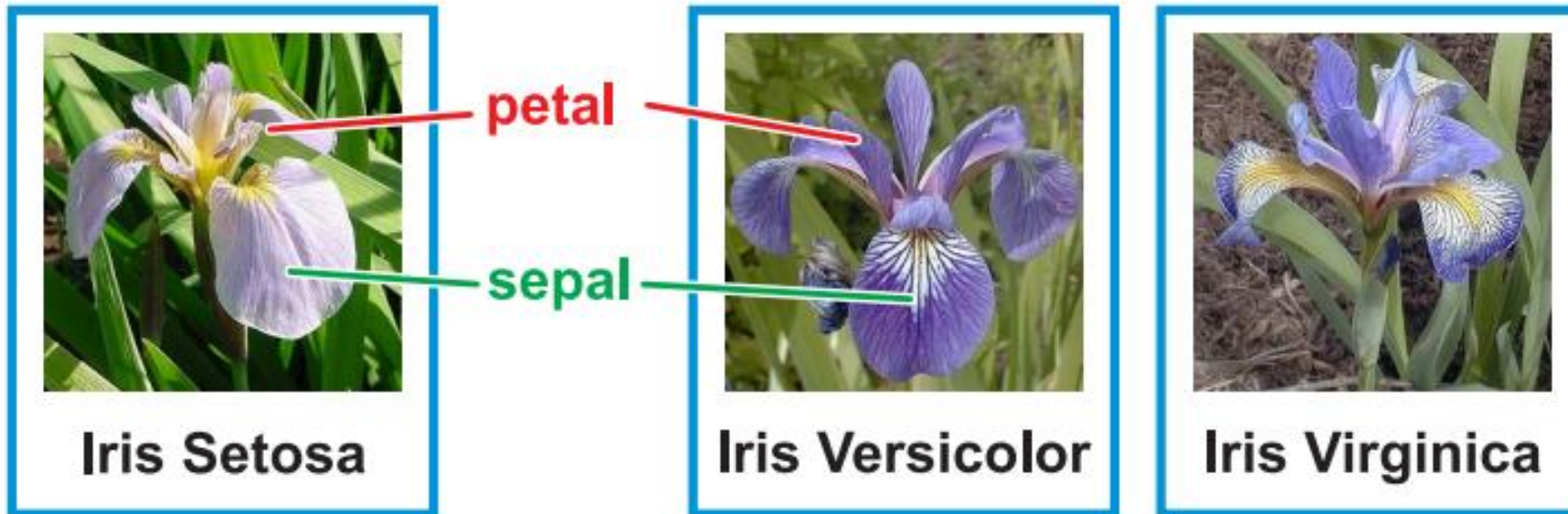


Regression



# Example: Classify Irises

- Most classical example of a classification problem
- The petals and sepals of different iris flowers have been measured
- Can we use this data to find a program which tells us to what type a flower belongs on basis of petal and sepal measurements?



# Example: Classify Irises

- Data samples  $t = (t_1, t_2, \dots, t_n)$ ;  $t_i \in \mathbb{R}$  belongs to classes  $k$  in  $K$
- Supervised learning: we use samples  $t \in A$  with known classes  $class(t) \in K$  to learn a function  $f(t_i)$

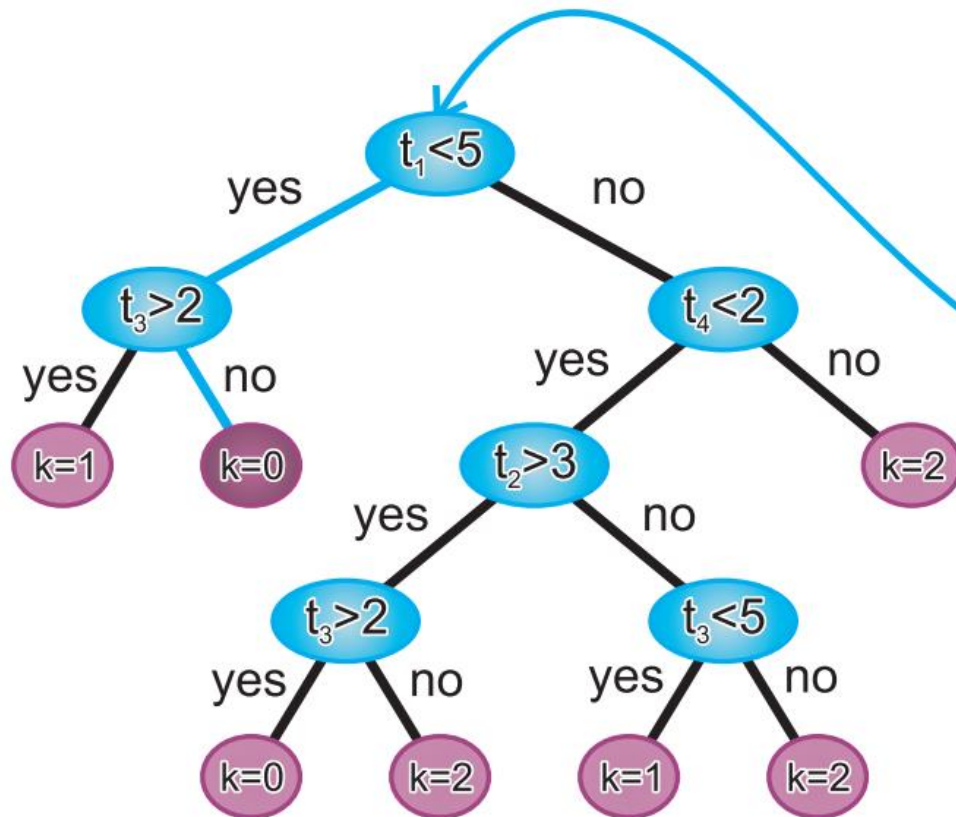
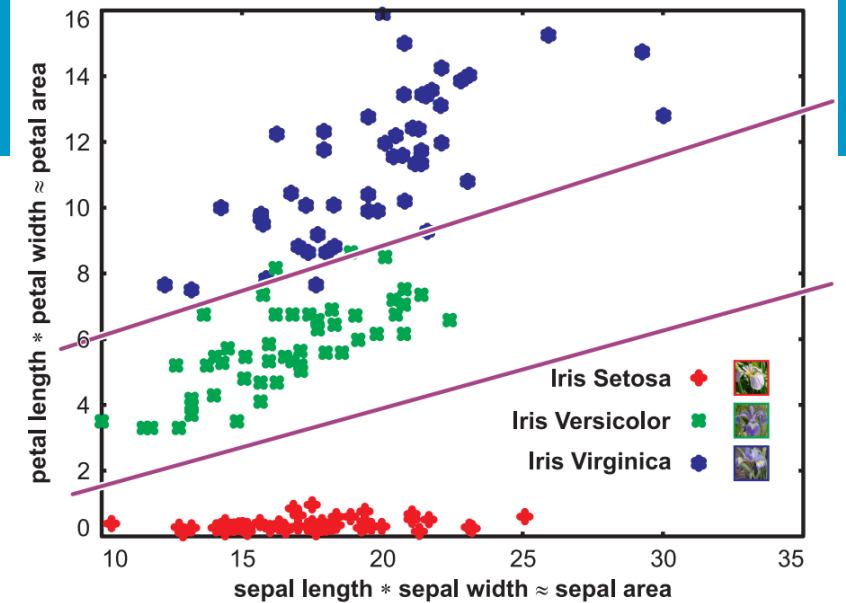
					Class	
				Sepal Length		
				Sepal Width	Class	
				Petal Length		
				Petal Width	Class	
A	$t \in A \{$	5.1	3.5	1.4	0.2	iris setosa
		4.9	3.0	1.4	0.2	iris setosa
		7.0	3.2	4.7	1.4	iris versicolor
		6.3	3.3	6.0	2.5	iris virginica
		...	...	...	...	...
		6.4	3.2	4.5	1.4	iris versicolor
		$t_1 \in \mathbb{R}$	$t_2 \in \mathbb{R}$	$t_3 \in \mathbb{R}$	$t_4 \in \mathbb{R}$	$K = \{setosa, versicolor, virginica\}$

$k = class(t)$



# Example: Classify Irises

- Common Approach: Decision Trees



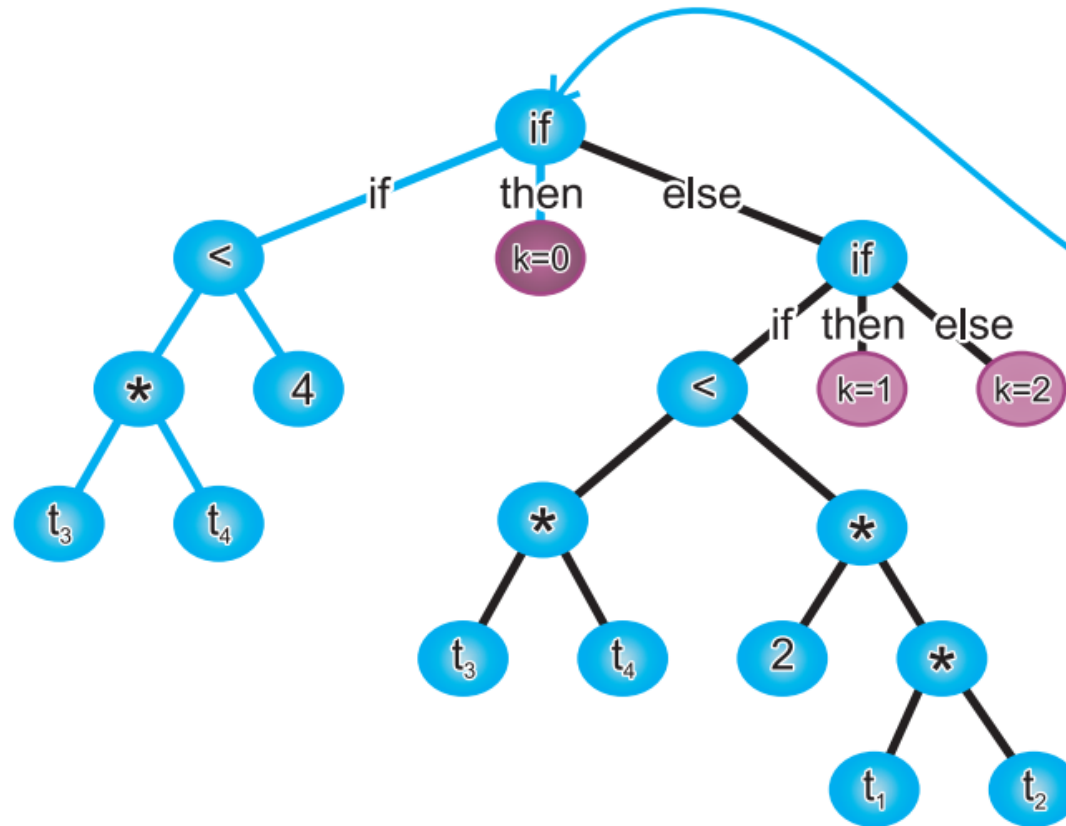
$t_1$	$t_2$	$t_3$	$t_4$	$k$
5.1	3.5	1.4	0.2	0
4.9	3.0	1.4	0.2	0
7.0	3.2	4.7	1.4	1
6.3	3.3	6.0	2.5	2
...	...	...	...	...
6.4	3.2	4.5	1.4	1

How decision trees work :

<https://www.youtube.com/watch?v=ZVR2Way4nwQ>

# Example: Classify Irises

- In Genetic Programming decisions and tree shapes are not limited to a certain shape



$t_1$	$t_2$	$t_3$	$t_4$	$k$
5.1	3.5	1.4	0.2	0
4.9	3.0	1.4	0.2	0
7.0	3.2	4.7	1.4	1
6.3	3.3	6.0	2.5	2
...	...	...	...	...
6.4	3.2	4.5	1.4	1

if  $t_3 * t_4 < 4 \Rightarrow$  class 0  
else if  $t_3 * t_4 < 2 * t_1 * t_2 \Rightarrow$  class 1  
else class 2