

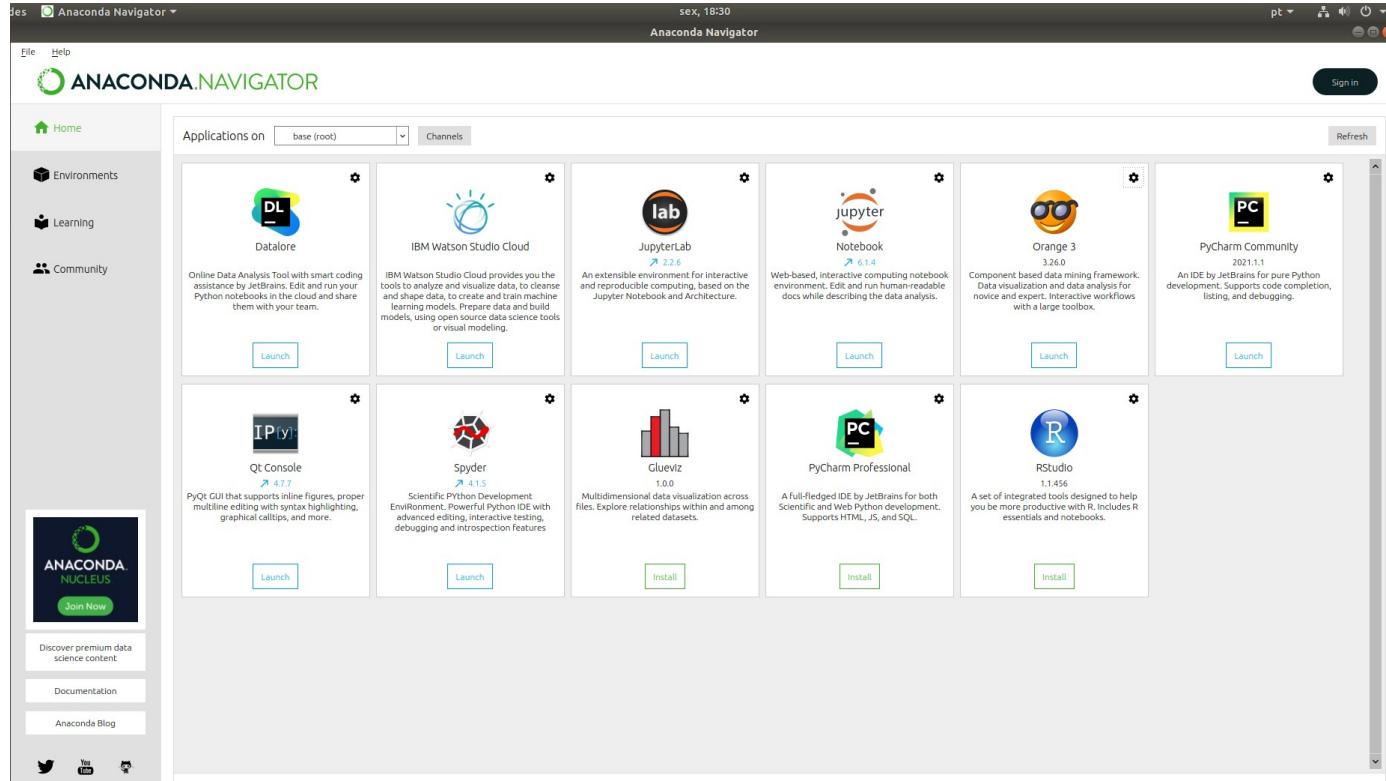
# Orange Data Mining Tool

Nádia Félix

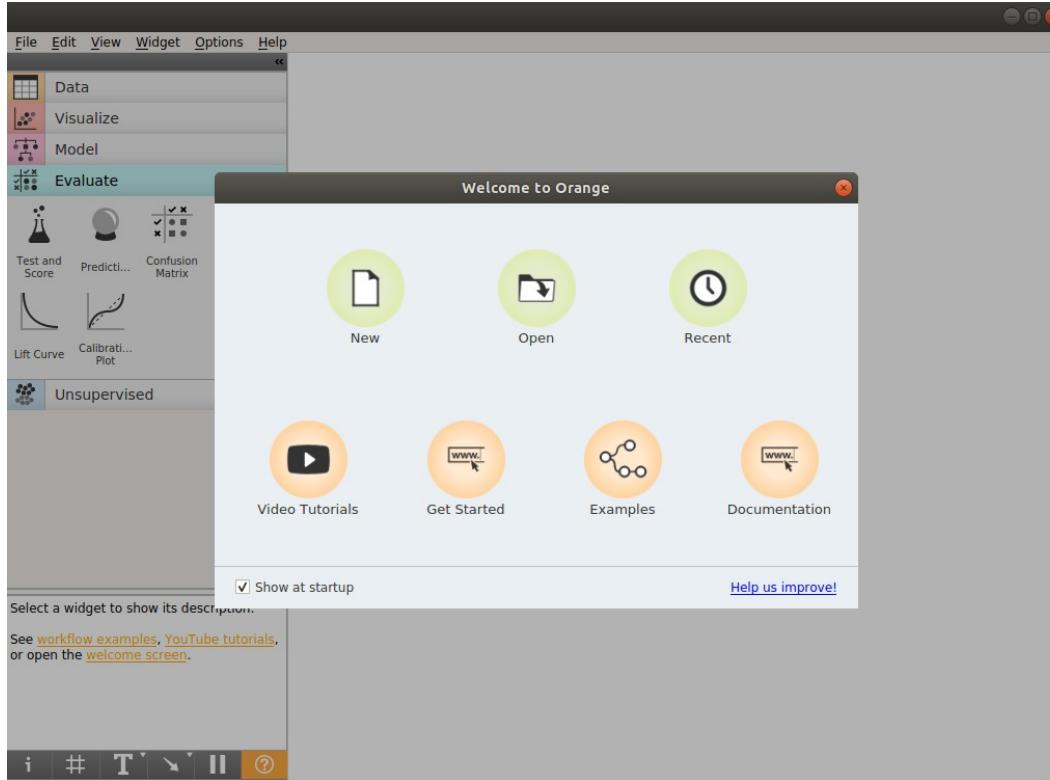
# Por que Orange?

- Open Source
- Component based
- No programming
- Data visualization
- Platform independent software
- Allows clustering and classification
- Data mining through visual programming
- and python scripting
- Por ser baixado em: <https://orangedatamining.com/>

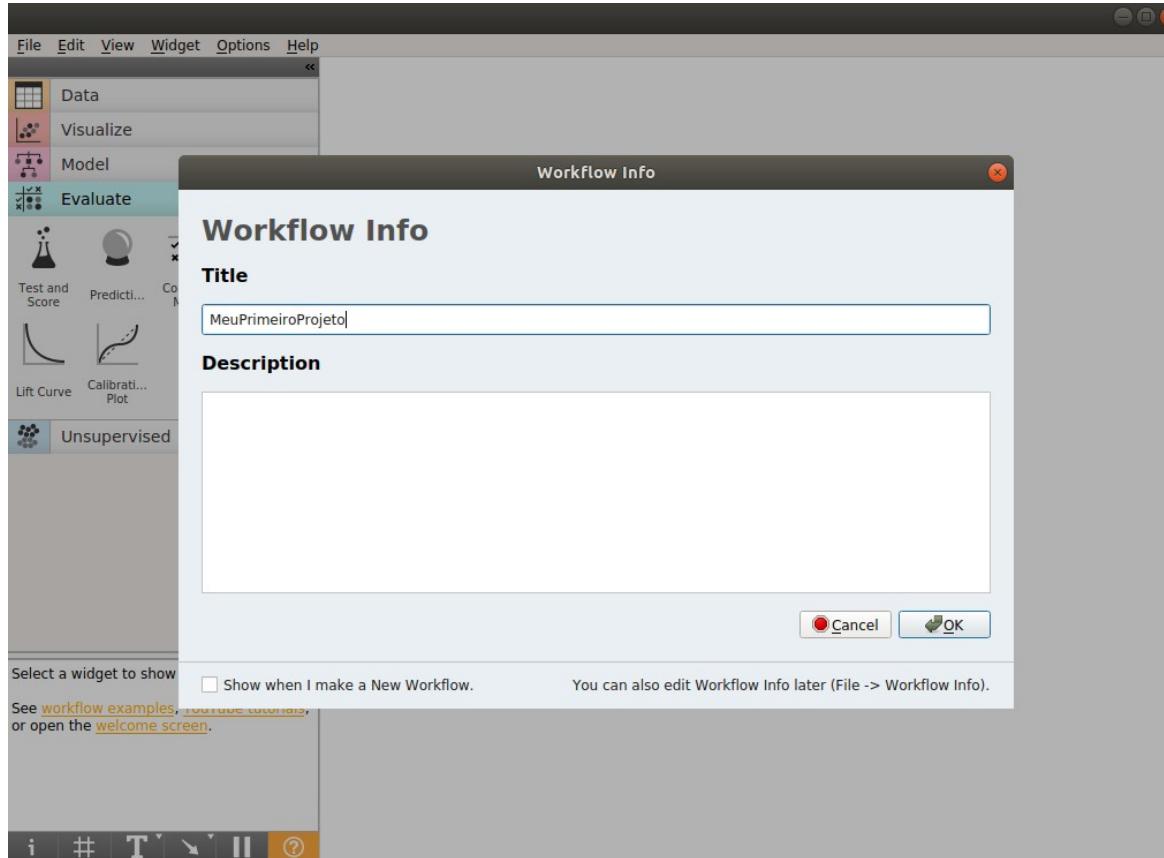
# Getting Started With ORANGE!!



# Getting Started With ORANGE!!



# Getting Started With ORANGE!!



The image displays three side-by-side screenshots of the Orange data mining software interface, each showing a different section of the application:

- Data Section:** Shows icons for Tree Viewer, Box Plot, Distributions, Scatter Plot, Sieve Diagram, Mosaic Display, FreeViz, Linear Projection, Radviz, Heat Map, Venn Diagram, Silhouette Plot, Pythagorean Tree, Pythagorean Forest, CN2 Rule Viewer, and Nomogram.
- Visualize Section:** Shows icons for Constant, CN2 Rule Induction, kNN, Tree, Random Forest, SVM, Linear Regression, Logistic Regression, Naive Bayes, AdaBoost, Neural Network, Stochastic Gradient Descent, Save Model, and Load Model.
- Evaluate Section:** Shows icons for Test & Score, Predictions, Confusion Matrix, ROC Analysis, Lift Curve, Calibration Plot, and Unsupervised.

# Análise exploratória dos dados – Exploratory DATA analysis

# Dataset: Heart Disease

## ATTRIBUTES

- Narrowing diameter
- Cholesterol
- Chest pain (dor no peito)
- Rest ECG
- Fasting blood sugar
- Max HR (Freq. Cardíaca)
- Age,gender and more

- Has 303 instances
- 13 attributes
- Categorical class with 2 values (0,1)
- In .csv format
- datasets of Orange.

# Verificar correlações entre os dados

- Idade: as doenças cardíacas aumentam com a idade acima de 65 anos
- Sexo: a doença cardíaca é a principal causa de morte tanto para homens como para mulheres

File

File: heart\_disease.tab

URL:

Info

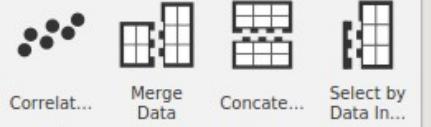
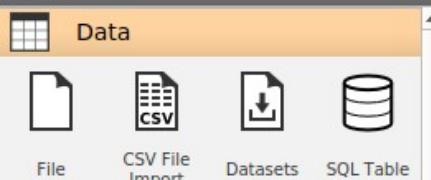
**Heart Disease dataset**  
Data on the presence of heart disease in patients.

303 instance(s)  
13 feature(s) (0.2% missing values)  
Classification; categorical class with 2 values (no missing values)  
0 meta attribute(s)

Columns (Double click to edit)

	Name	Type	Role	Values
9	exerc ind ang	C categorical	feature	0, 1
10	ST by exercise	N numeric	feature	
11	slope peak exc ST	C categorical	feature	upsloping, flat, downsloping
12	major vessels colored	N numeric	feature	
13	thal	C categorical	feature	normal, reversable defect, fixed defect
14	diameter narrowing	C categorical	target	0, 1

11/06/21



## Data Table

View the dataset in a spreadsheet.

[more...](#)



Data Table

### Variables

- Show variable labels (if present)
- Visualize numeric values
- Color by instance classes

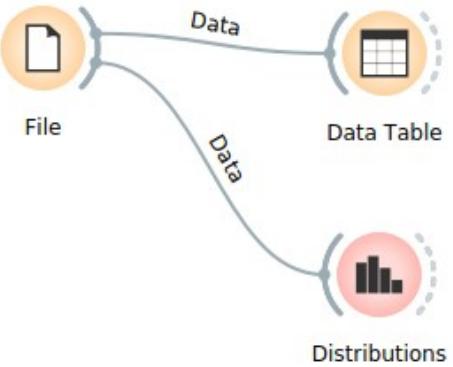
### Selection

- Select full rows

	meter narrowi	age	gender	chest pain	rest SBP
1	0	63	male	typical ang	145
2	1	67	male	asymptomatic	160
3	1	67	male	asymptomatic	120
4	0	37	male	non-anginal	130
5	0	41	female	atypical ang	130
6	0	56	male	atypical ang	120
7	1	62	female	asymptomatic	140
8	0	57	female	asymptomatic	120
9	1	63	male	asymptomatic	130
10	1	53	male	asymptomatic	140
11	0	57	male	asymptomatic	140
12	0	56	female	atypical ang	140
13	1	56	male	non-anginal	130
14	0	44	male	atypical ang	120
15	0	52	male	non-anginal	172
16	0	57	male	non-anginal	150
17	1	48	male	atypical ang	110
18	0	54	male	asymptomatic	140
19	0	48	female	non-anginal	130
20	0	49	male	atypical ang	130
21	0	64	male	typical ang	110
22	0	58	female	typical ang	150
23	1	58	male	atypical ang	120
24	1	58	male	non-anginal	132
25	1	60	male	asymptomatic	130
26	0	50	female	non-anginal	120

[Restore Original Order](#)

- Send Automatically



## Distributions

Display value distributions of a data feature in a graph.

[more...](#)

## Distributions

### Variable

- C diameter narrowing
- N age
- C gender
- C chest pain
- N rest SBP
- N cholesterol
- C fasting blood sugar > 120

### Distribution

Bin width

Fitted distribution

Hide bars

### Columns

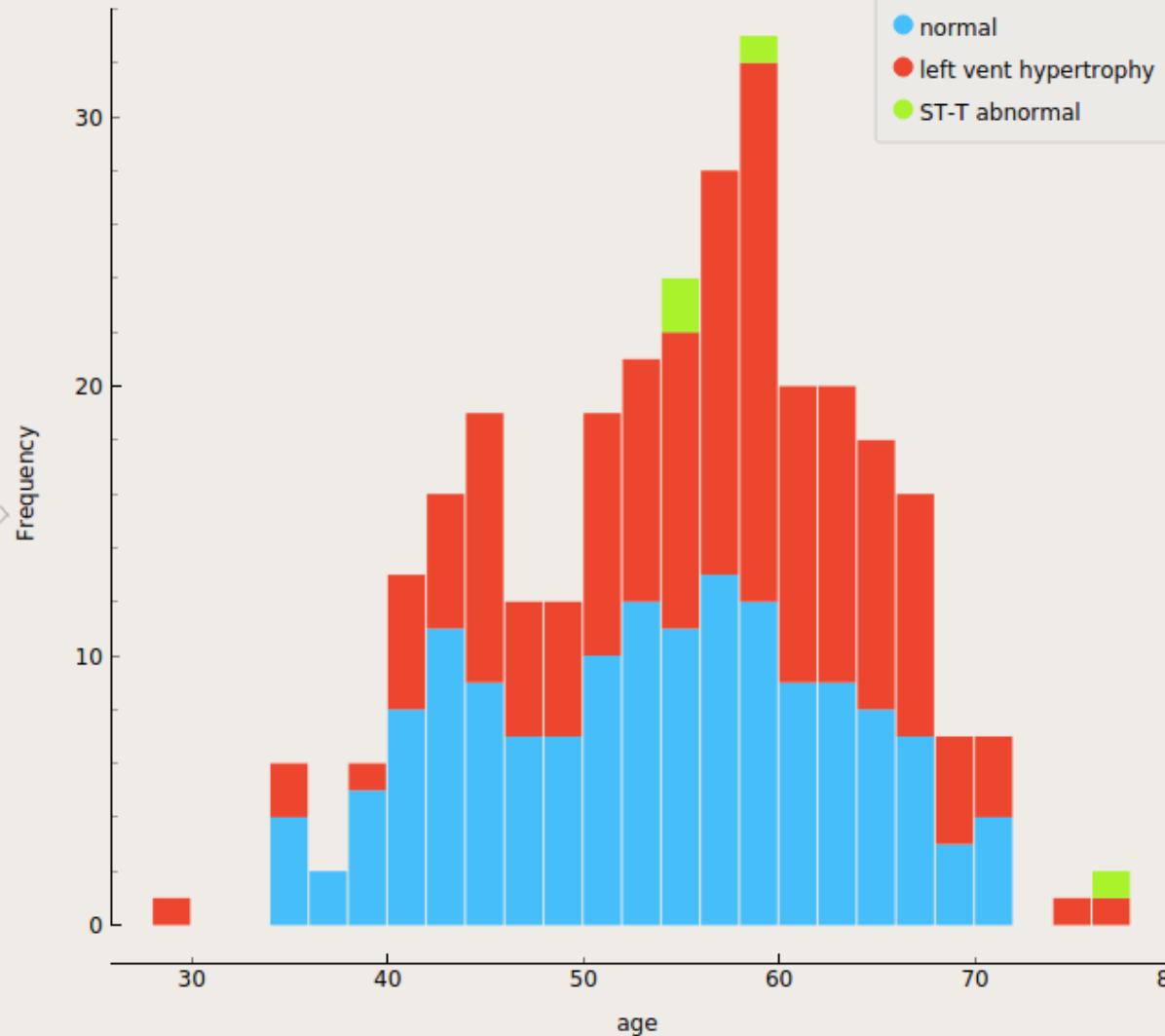
Split by

Stack columns

Show probabilities

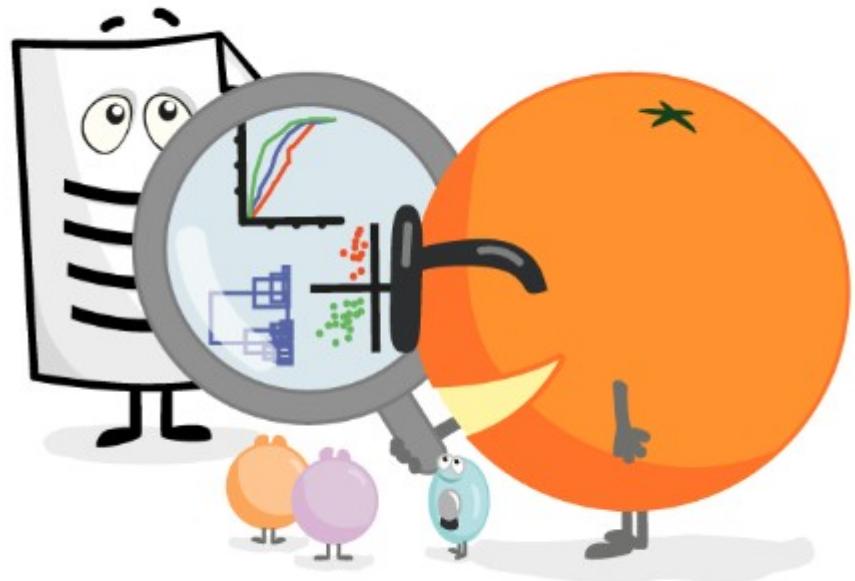
Show cumulative distribution

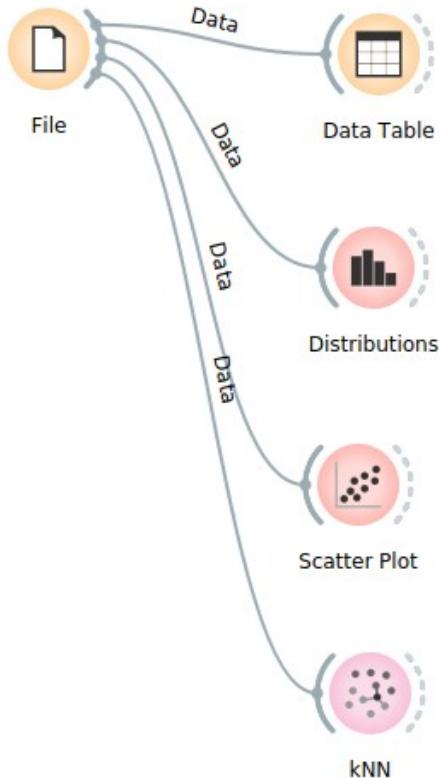
Apply Automatically



## Algorithms:

- KNN
- Naïve Bayes
- Decision Tree
- SVM





## kNN

Predict according to the nearest training instances.

[more...](#)

File Edit View Widget Options Help

«

Data

Visualize

Model

Constant CN2 Rule Induction Calibrated Learner kNN

Tree Random Forest SVM Linear Regress...

Logistic Regress... Naive Bayes AdaBoost Neural Network

Stochastic Gradien... Stacking Save Model Load Model

Evaluate

Unsupervised

kNN

Predict according to the nearest training instances.

more...

The diagram illustrates the data flow in a machine learning pipeline. It starts with a 'File' icon connected to a 'Data Table' icon, which is then connected to a 'kNN' model configuration window. The window shows the configuration for a 'kNN' model named 'kNN' with 5 neighbors, using Euclidean metric and Uniform weight. There are also 'Apply Automatically' and help buttons at the bottom.

File Data Table kNN

Name: kNN

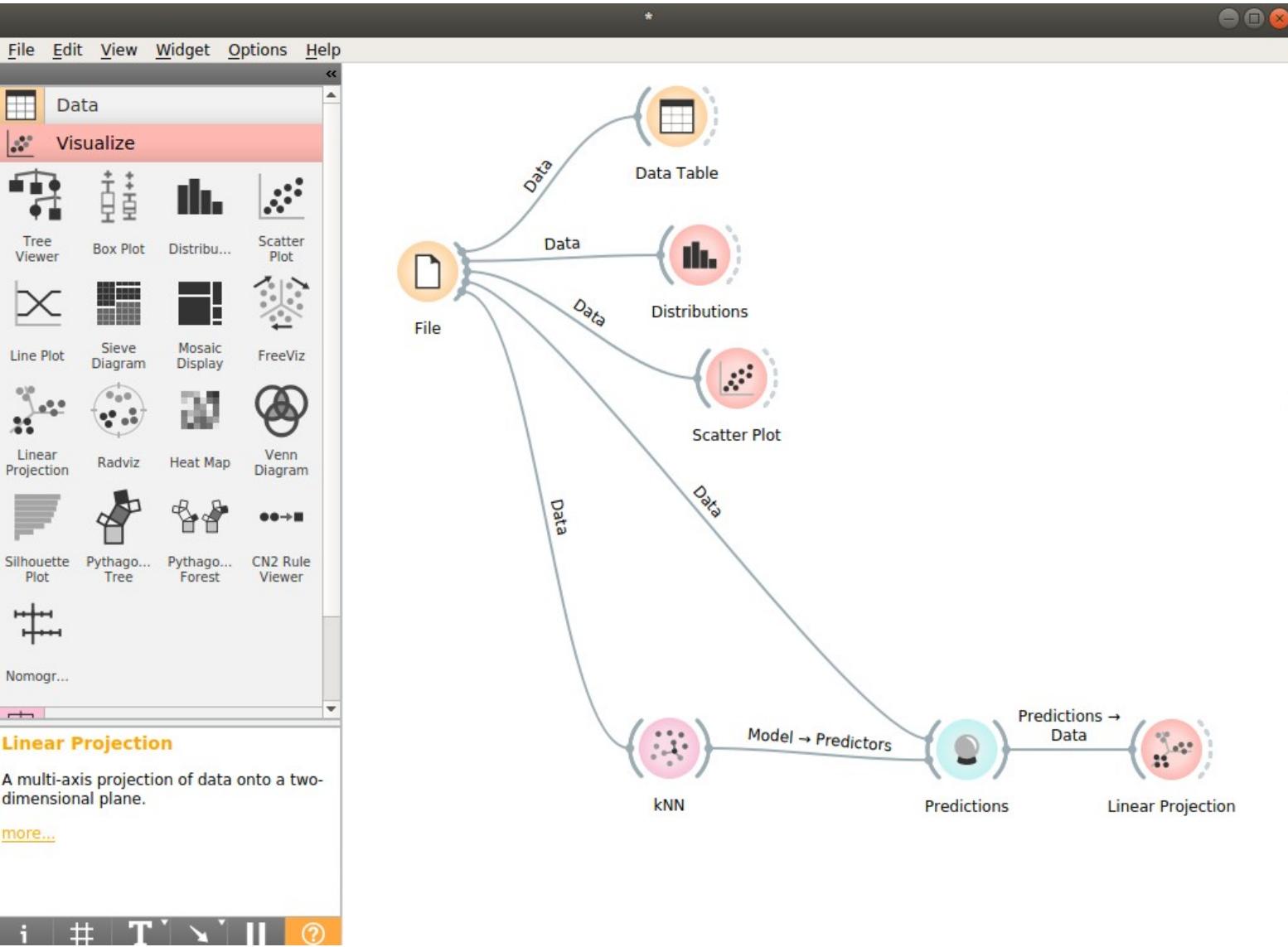
Neighbors:

- Number of neighbors: 5
- Metric: Euclidean
- Weight: Uniform

Apply Automatically

?

303

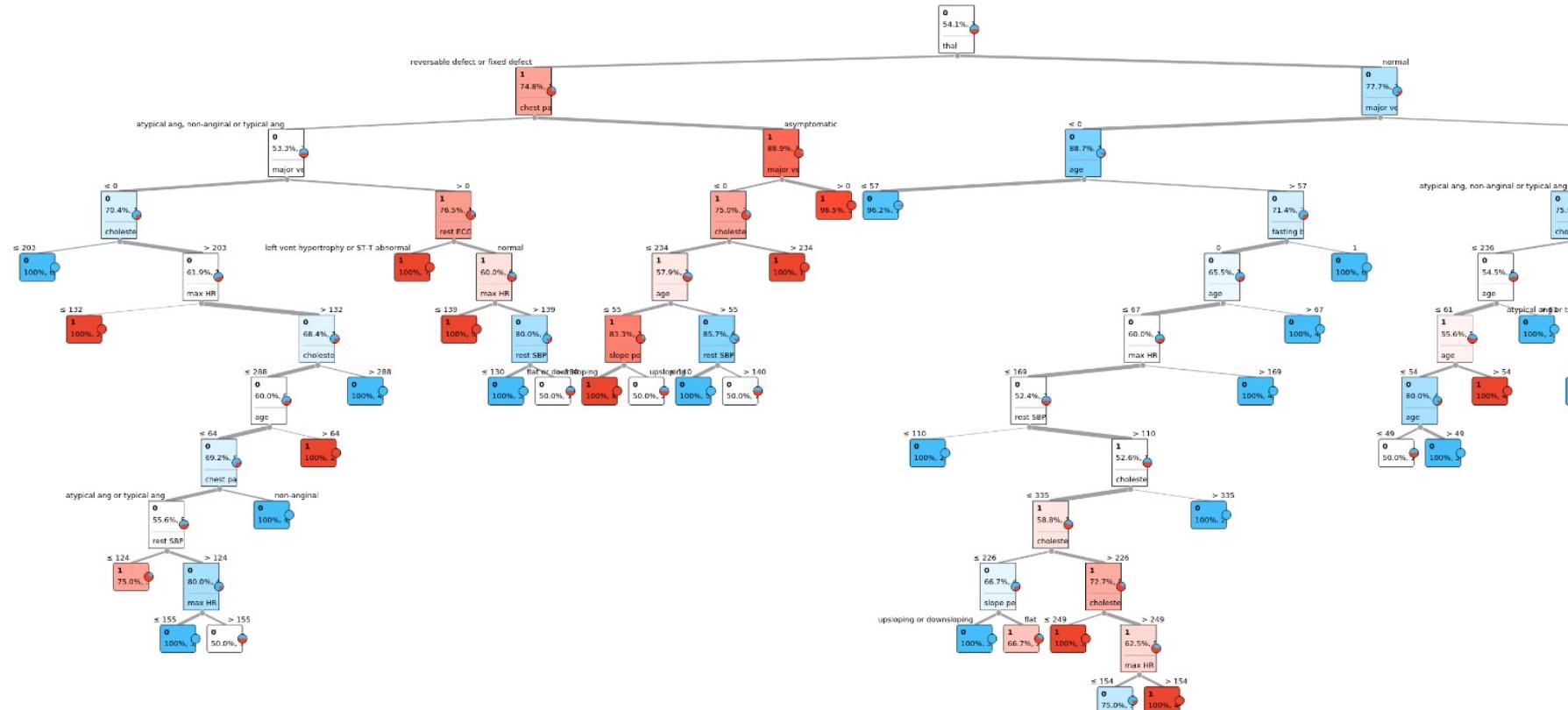


## Predictions

Show probabilities for	kNN	ameter narrowi	age	gender	chest pain	rest SBP	cholesterol	ig blood sugar >	rest ECG	max HR	exerc ind ang	ST by exercise	lope peak exc	Major vessels color	thal	
0	1	0.60 : 0.40 → 0	0	63	male	typical ang	145	233	1	left vent hyp...	150	0	2.3	downsloping	0	fixed defect
1	2	0.00 : 1.00 → 1	1	67	male	asymptomatic	160	286	0	left vent hyp...	108	1	1.5	flat	3	normal
2	3	0.00 : 1.00 → 1	1	67	male	asymptomatic	120	229	0	left vent hyp...	129	1	2.6	flat	2	reversible d...
3	4	1.00 : 0.00 → 0	0	37	male	non-anginal	130	250	0	normal	187	0	3.5	downsloping	0	normal
4	5	1.00 : 0.00 → 0	0	41	female	atypical ang	130	204	0	left vent hyp...	172	0	1.4	upsloping	0	normal
5	6	0.80 : 0.20 → 0	0	56	male	atypical ang	120	236	0	normal	178	0	0.8	upsloping	0	normal
6	7	0.60 : 0.40 → 0	1	62	female	asymptomatic	140	268	0	left vent hyp...	160	0	3.6	downsloping	2	normal
7	8	0.60 : 0.40 → 0	0	57	female	asymptomatic	120	354	0	normal	163	1	0.6	upsloping	0	normal
8	9	0.00 : 1.00 → 1	1	63	male	asymptomatic	130	254	0	left vent hyp...	147	0	1.4	flat	1	reversible d...
9	10	0.40 : 0.60 → 1	1	53	male	asymptomatic	140	203	1	left vent hyp...	155	1	3.1	downsloping	0	reversible d...
10	11	0.40 : 0.60 → 1	0	57	male	asymptomatic	140	192	0	normal	148	0	0.4	flat	0	fixed defect
11	12	0.80 : 0.20 → 0	0	56	female	atypical ang	140	294	0	left vent hyp...	153	0	1.3	flat	0	normal
12	13	0.00 : 1.00 → 1	1	56	male	non-anginal	130	256	1	left vent hyp...	142	1	0.6	flat	1	fixed defect
13	14	1.00 : 0.00 → 0	0	44	male	atypical ang	120	263	0	normal	173	0	0.0	upsloping	0	reversible d...
14	15	0.60 : 0.40 → 0	0	52	male	non-anginal	172	199	1	normal	162	0	0.5	upsloping	0	reversible d...
15	16	0.60 : 0.40 → 0	0	57	male	non-anginal	150	168	0	normal	174	0	1.6	upsloping	0	normal
16	17	0.40 : 0.60 → 1	1	48	male	atypical ang	110	229	0	normal	168	0	1.0	downsloping	0	reversible d...
17	18	0.80 : 0.20 → 0	0	54	male	asymptomatic	140	239	0	normal	160	0	1.2	upsloping	0	normal
18	19	0.60 : 0.40 → 0	0	48	female	non-anginal	130	275	0	normal	139	0	0.2	upsloping	0	normal
19	20	0.80 : 0.20 → 0	0	49	male	atypical ang	130	266	0	normal	171	0	0.6	upsloping	0	normal
20	21	0.80 : 0.20 → 0	0	64	male	typical ang	110	211	0	left vent hyp...	144	1	1.8	flat	0	normal
21	22	0.80 : 0.20 → 0	0	58	female	typical ang	150	283	1	left vent hyp...	162	0	1.0	upsloping	0	normal
22	23	0.80 : 0.20 → 0	1	58	male	atypical ang	120	284	0	left vent hyp...	160	0	1.8	flat	0	normal
23	24	0.60 : 0.40 → 0	1	58	male	non-anginal	132	224	0	left vent hyp...	173	0	3.2	upsloping	2	reversible d...
24	25	0.40 : 0.60 → 1	1	60	male	asymptomatic	130	206	0	left vent hyp...	132	1	2.4	flat	2	reversible d...
25	26	1.00 : 0.00 → 0	0	50	female	non-anginal	120	219	0	normal	158	0	1.6	flat	0	normal
26	27	0.80 : 0.20 → 0	0	58	female	non-anginal	120	340	0	normal	172	0	0.0	upsloping	0	normal
27	28	0.40 : 0.60 → 1	0	66	female	typical ang	150	226	0	normal	114	0	2.6	downsloping	0	normal
28	29	1.00 : 0.00 → 0	0	43	male	asymptomatic	150	247	0	normal	171	0	1.5	upsloping	0	normal
29	30	0.40 : 0.60 → 1	1	40	male	asymptomatic	110	167	0	left vent hyp...	114	1	2.0	flat	0	reversible d...
30	31	0.60 : 0.40 → 0	0	69	female	typical ang	140	239	0	normal	151	0	1.8	upsloping	2	normal

Tree Viewer

**Tree size:** 77 nodes, 39 leaves  
**Edge widths:** Relative to parent  
**Target class:** None



**Write a comment...**

# Validação Cruzada

The screenshot shows the Weka interface with a focus on the "Test and Score" option in the "Widget" menu. The main workspace displays a flow diagram for cross-validation:

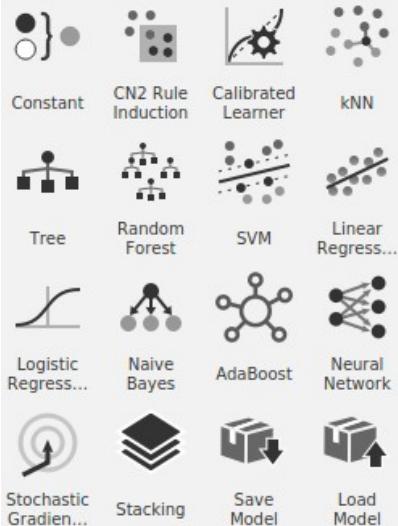
```
graph LR; File -- Data --> DataSampler[Data Sampler]; DataSampler -- "Data Sample → Data" --> Learner[Tree]; Learner -- Learner --> TestScore[Test and Score];
```

The "Model" panel on the left lists various machine learning algorithms, including Tree, Random Forest, SVM, Linear Regression, Logistic Regression, Naive Bayes, AdaBoost, Neural Network, Stochastic Gradient Descent, Stacking, Save Model, and Load Model.

**Widget menu (Test and Score):**

- Cross-validation accuracy estimation.
- [more...](#)

Data  
Visualize  
Model

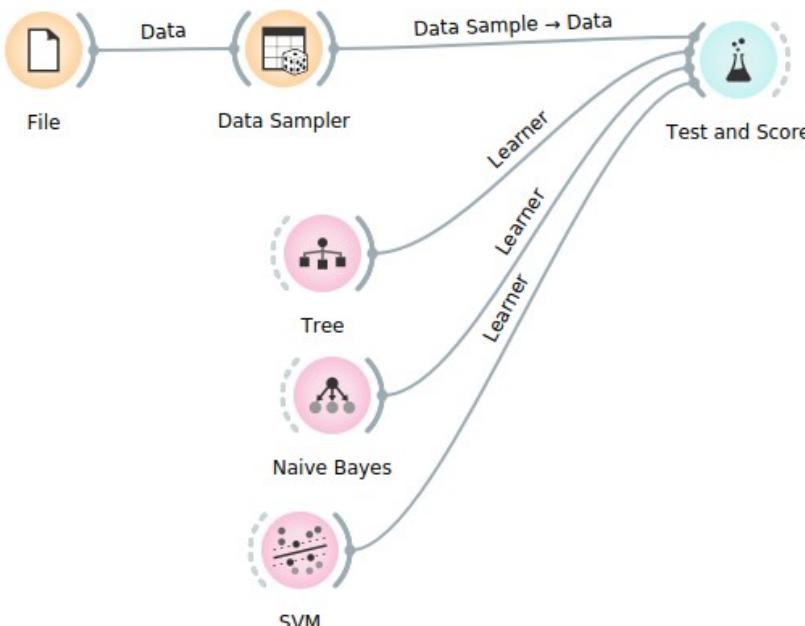


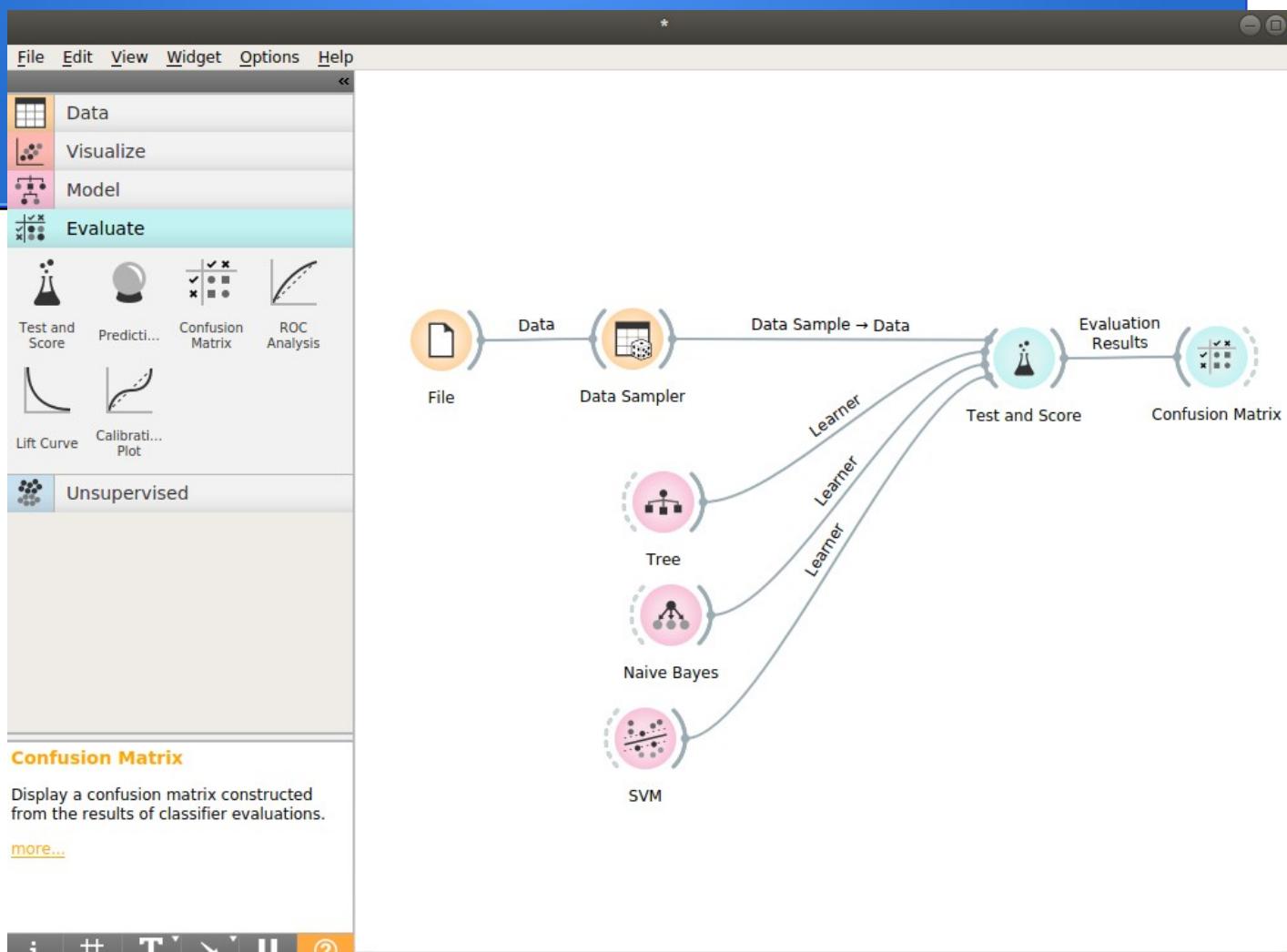
Evaluate  
Unsupervised

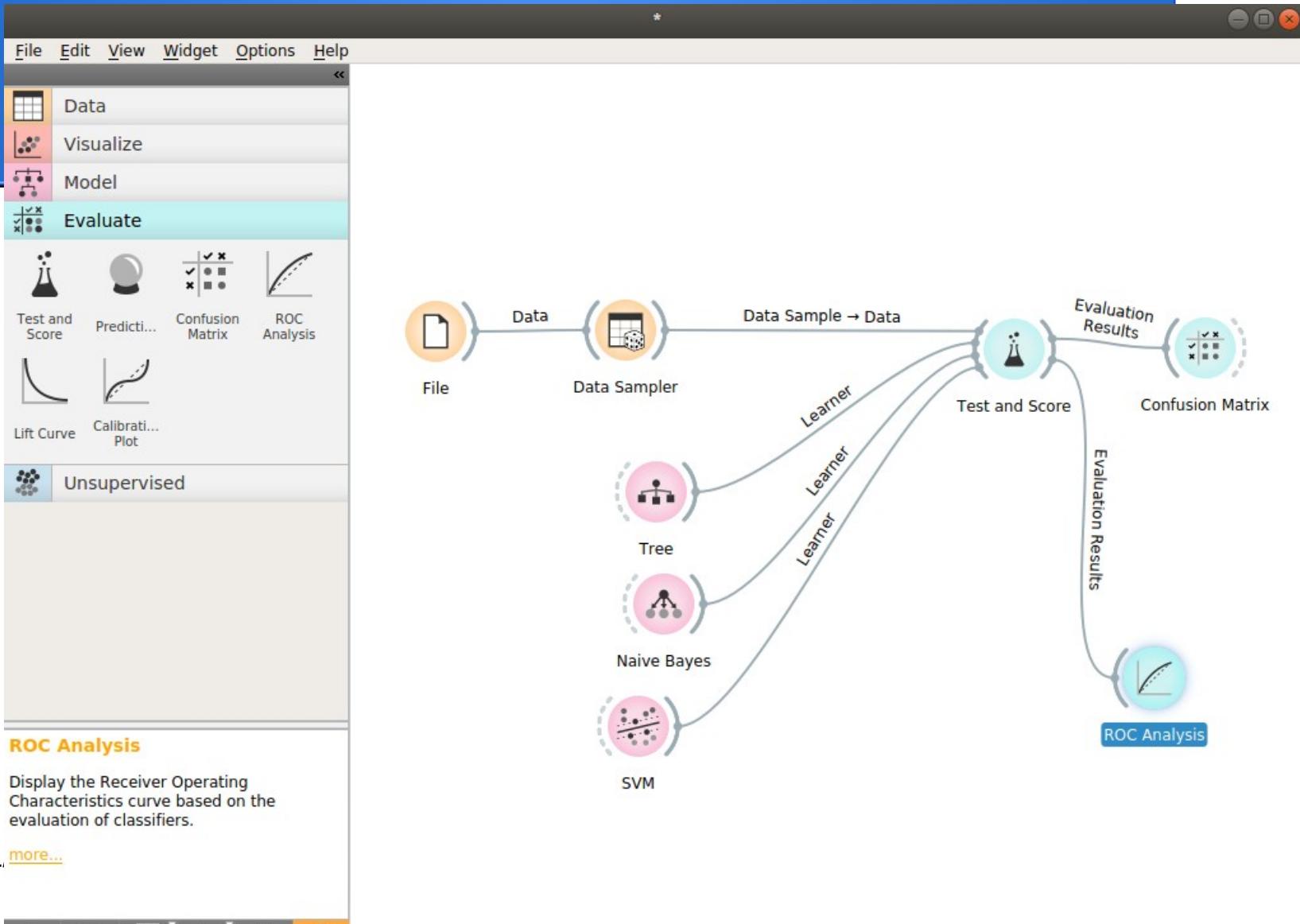
**Test and Score**

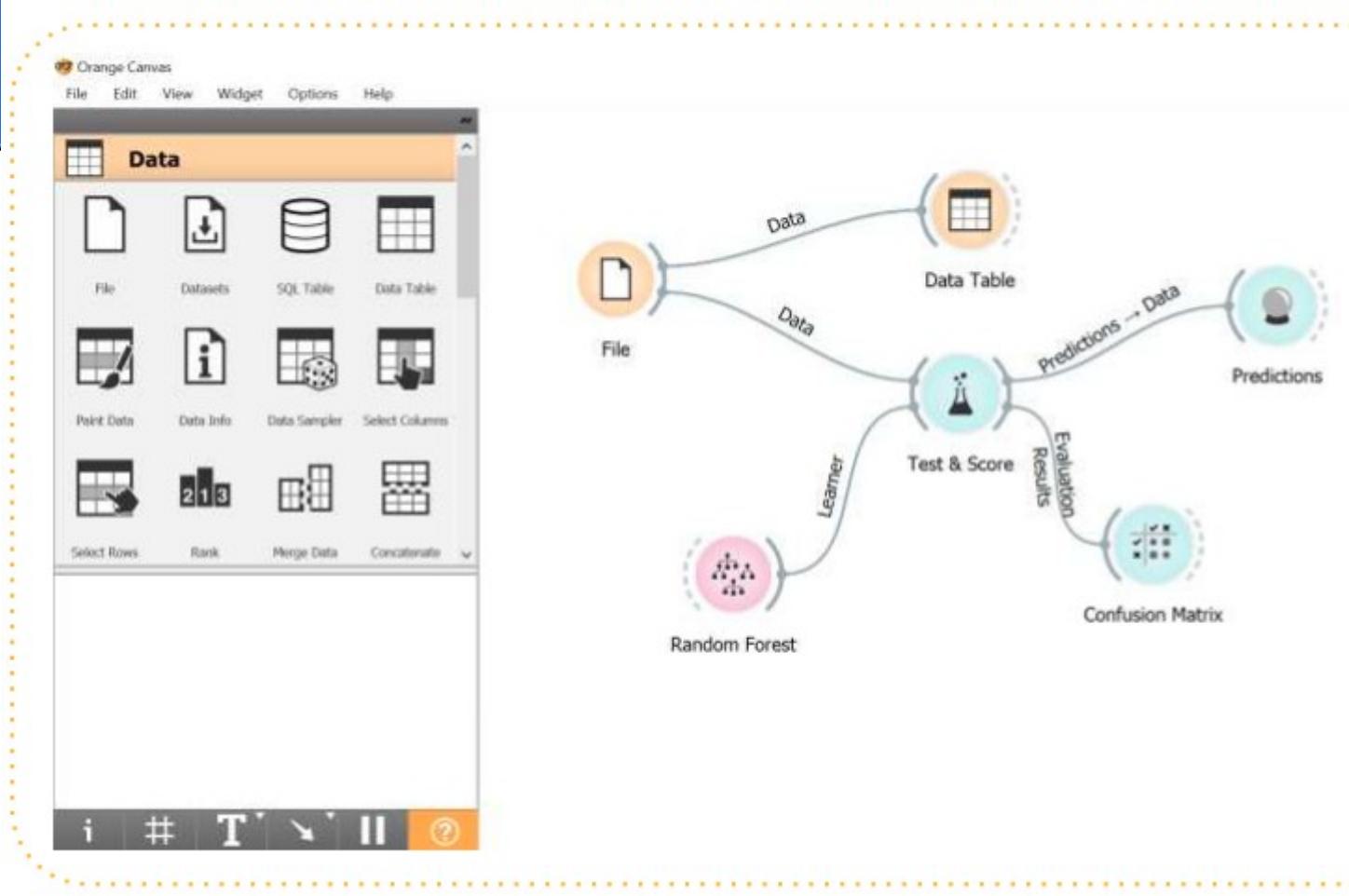
Cross-validation accuracy estimation.

[more...](#)









# Referências

<https://orange.biolab.si/getting-started/>  
<https://orangedatamining.com/>