



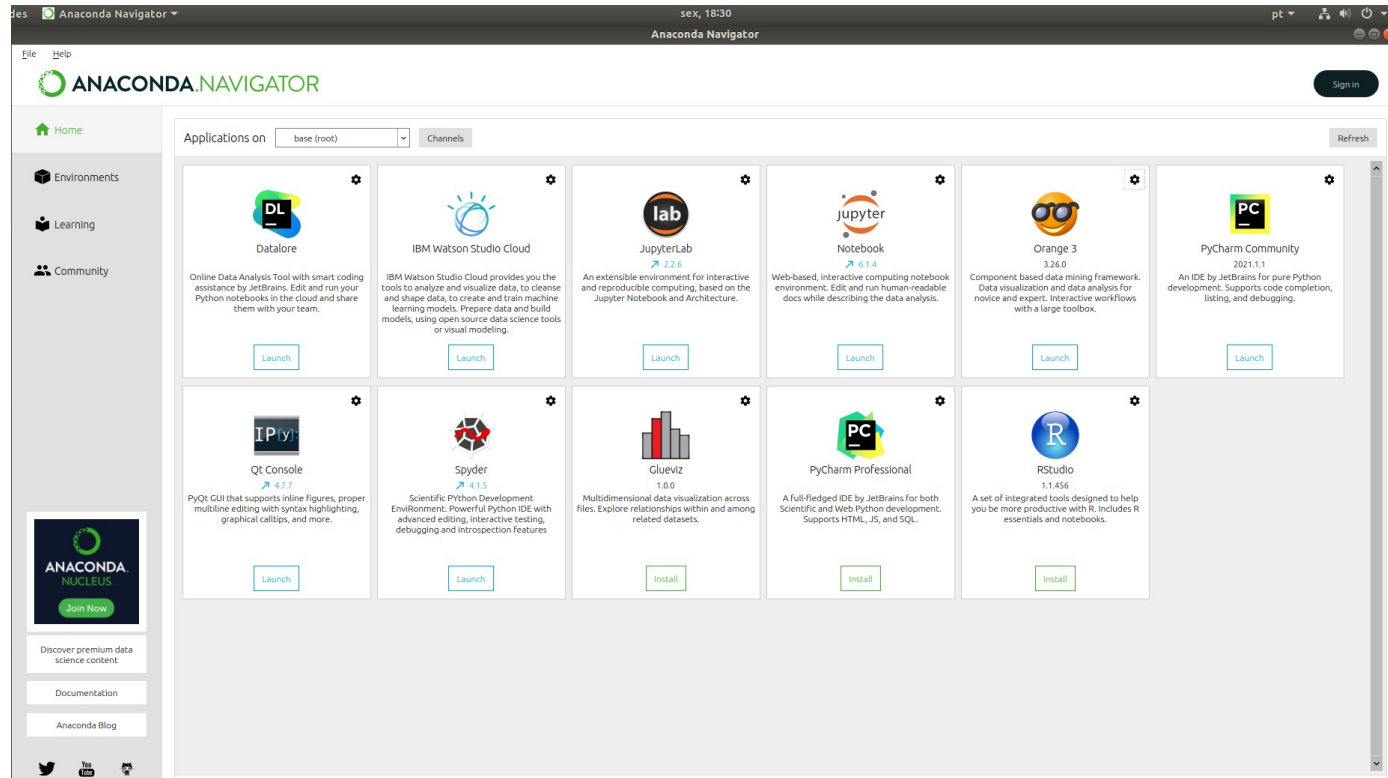
Orange Data Mining Tool

Nádia Félix

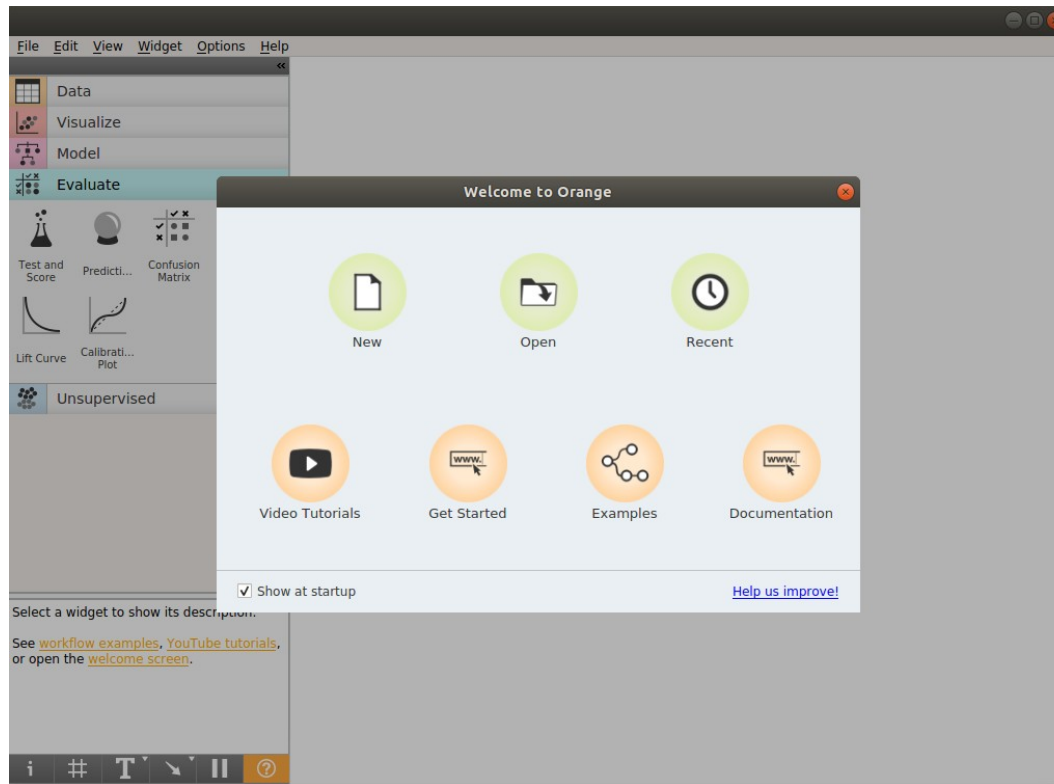
Por que Orange?

- 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/>

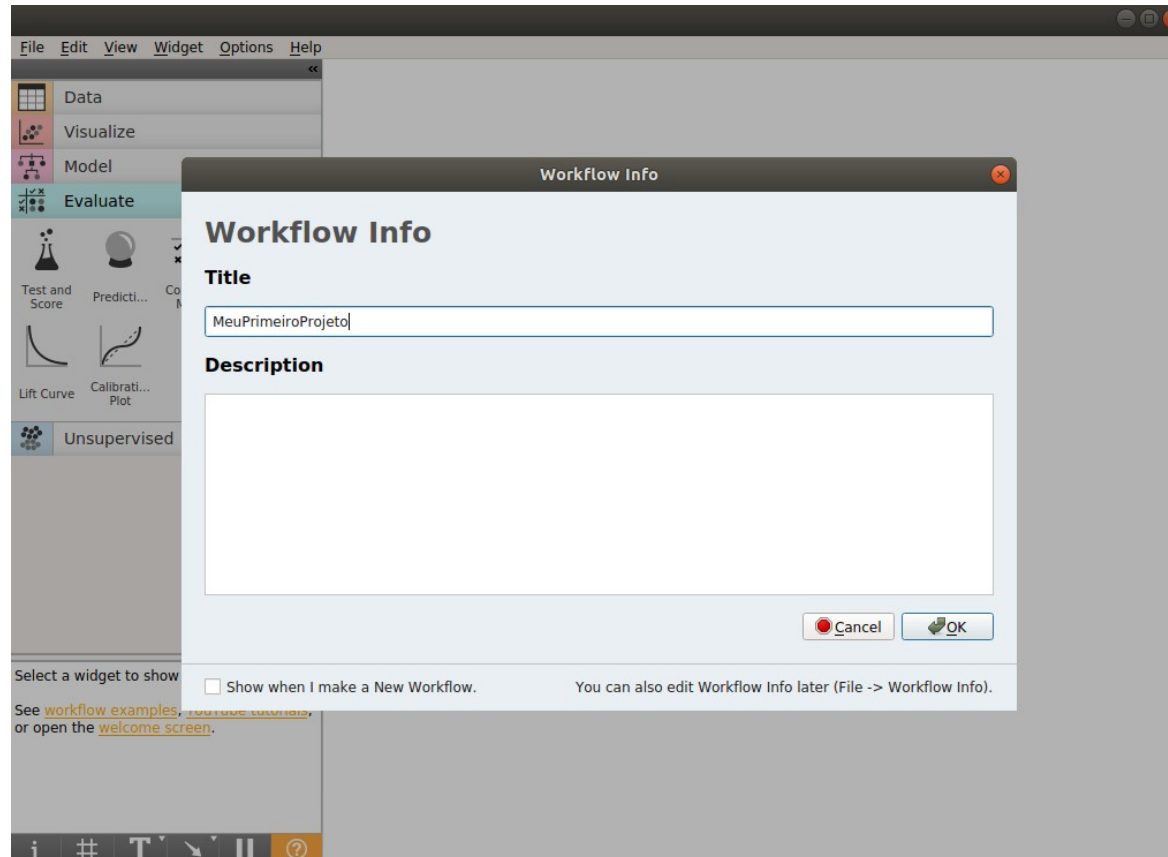
Getting Started With ORANGE!!

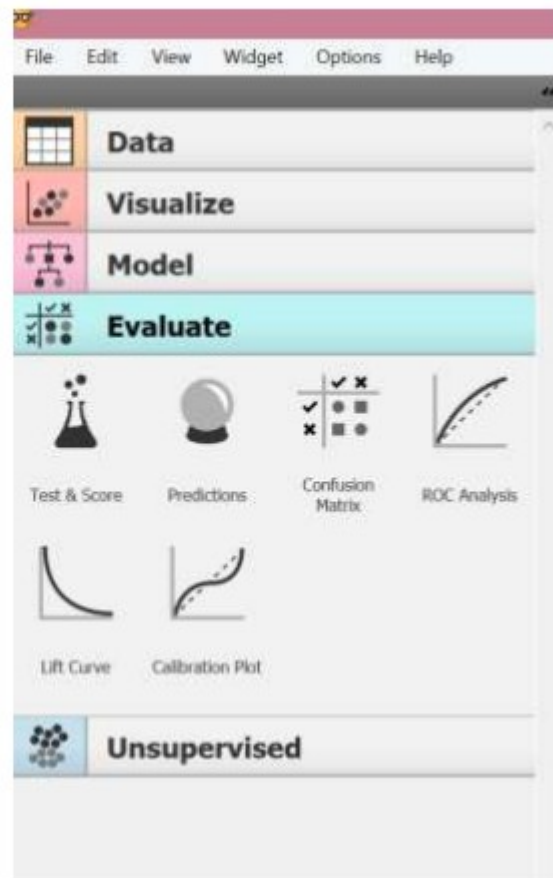
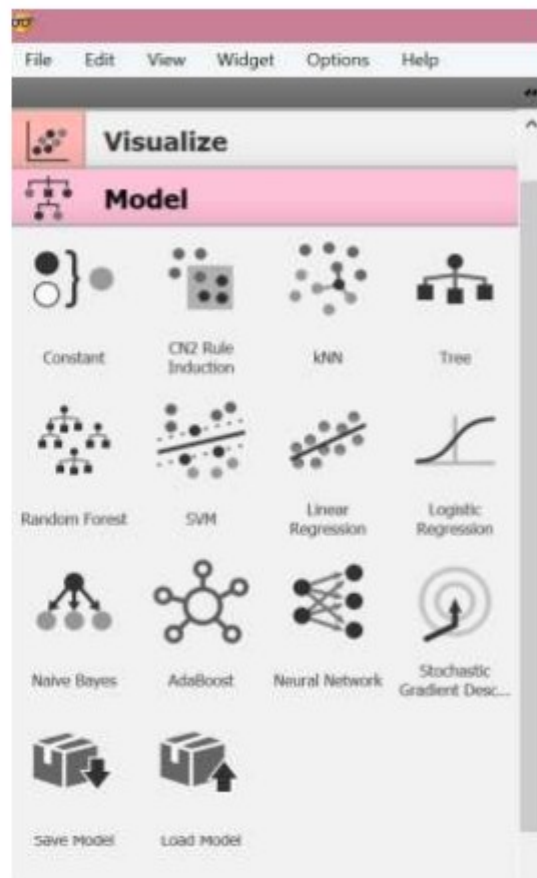


Getting Started With ORANGE!!



Getting Started With ORANGE!!





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

...

Reload

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

Browse documentation datasets

Reset

Apply

File Edit View Widget Options Help

Data

File CSV File Import Datasets SQL Table

Data Table Paint Data Data Info Data Sampler

Select Columns Select Rows Pivot Table Rank

Correlat... Merge Data Concatenate... Select by Data In...

Transpose Random... Preproc... Apply Domain

Data Table

View the dataset in a spreadsheet.
[more...](#)

File Data Data Table

Data Table

Variables

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

Selection

- ☒ Select full rows

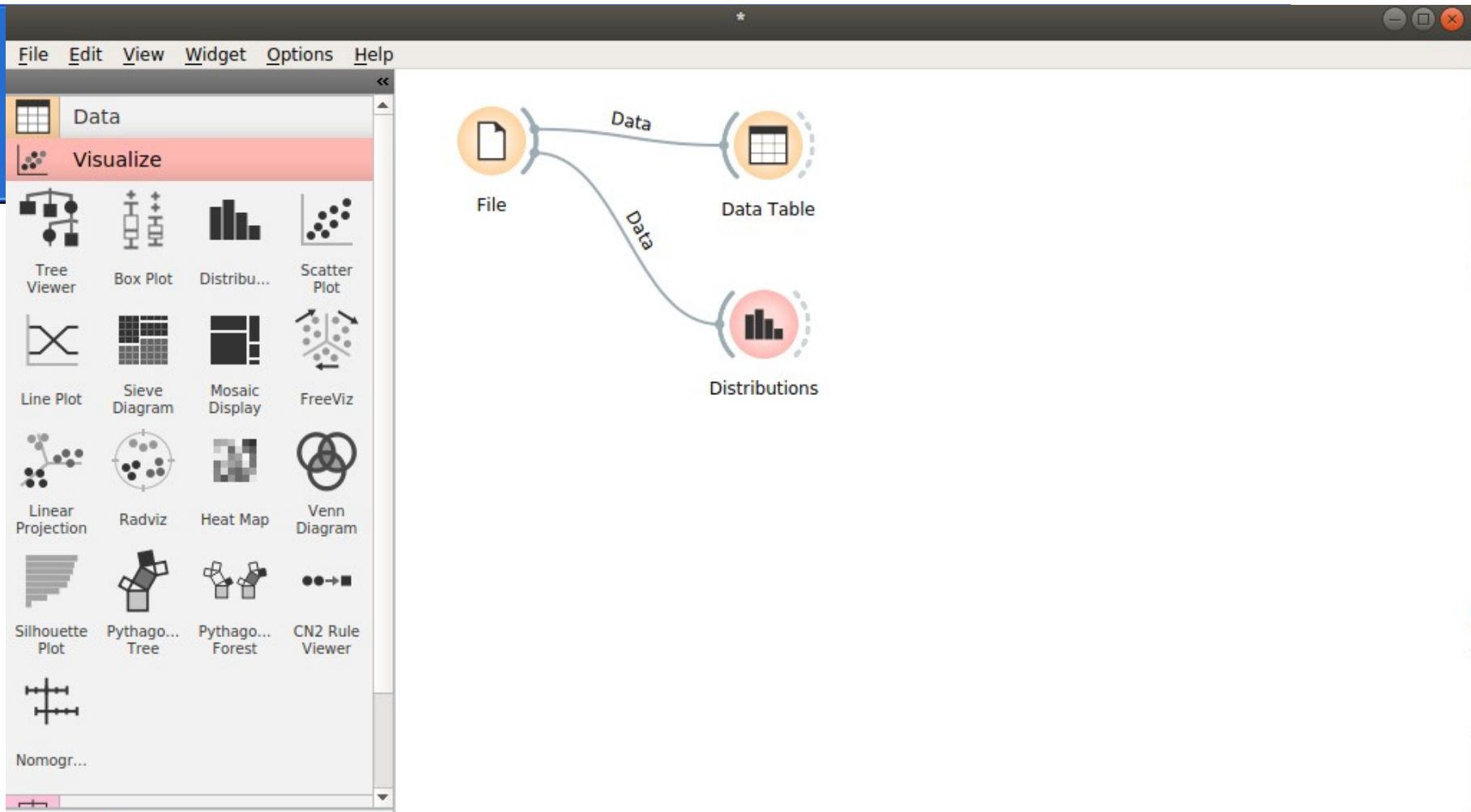
meter narrowi age gender chest pain rest SBP

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

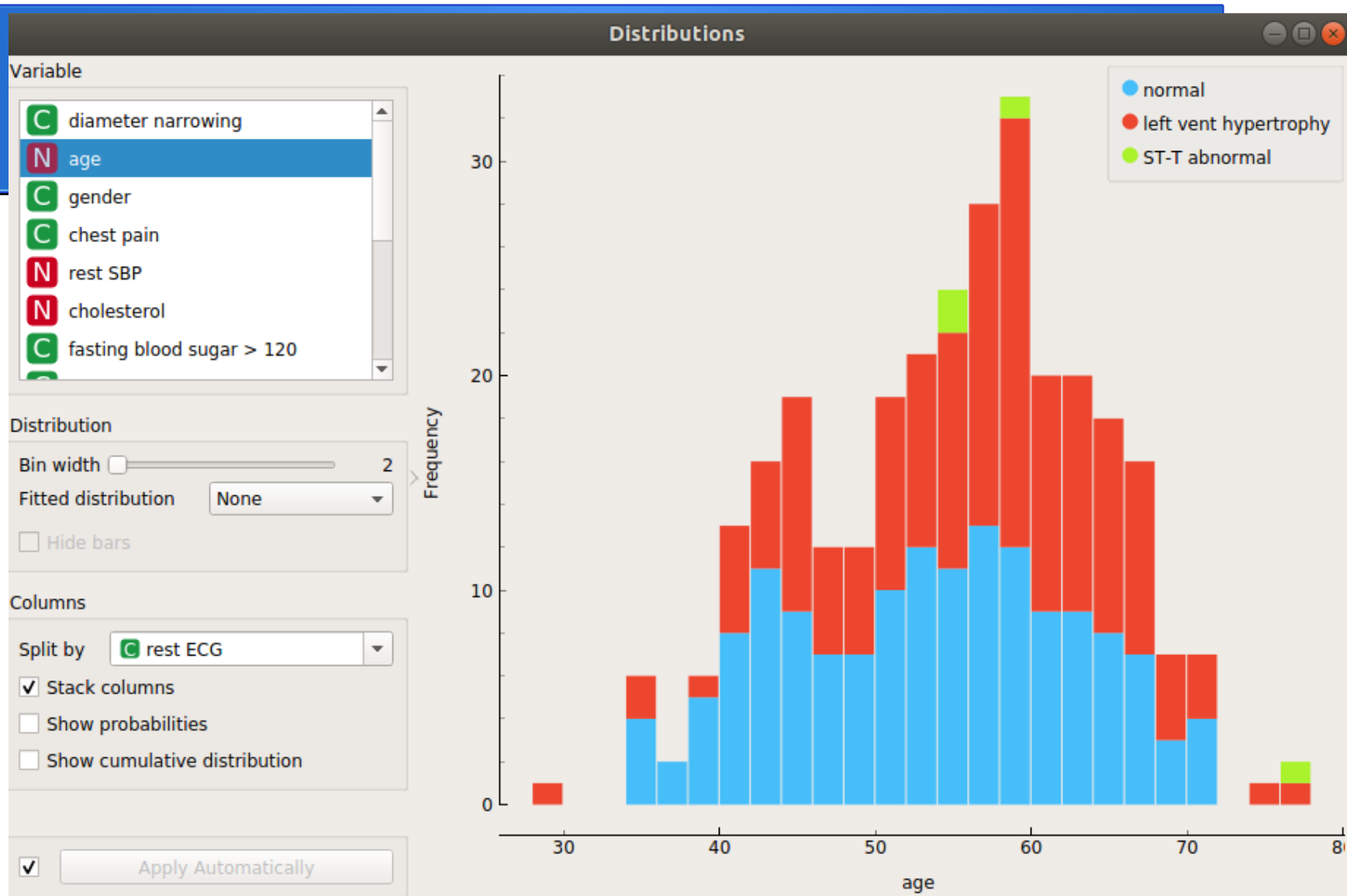
? | 303 | 1



Distributions

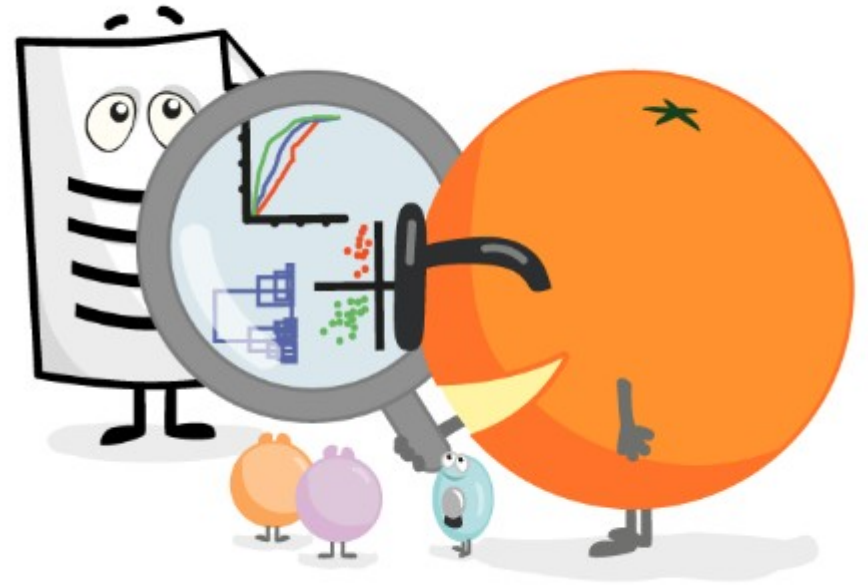
Display value distributions of a data feature in a graph.

[more...](#)



Algorithms:

- KNN
- Naïve Bayes
- Decision Tree
- SVM



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

File

Data

Data

Data

Data

Data Table

Distributions

Scatter Plot

kNN

kNN

Predict according to the nearest training instances.

[more...](#)

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The screenshot displays the Orange3 data mining software interface. On the left, a widget palette is visible with categories: Data, Visualize, and Model. The 'Model' category is selected, showing various machine learning widgets. A workflow is built on the canvas, starting with a 'File' widget connected to three 'Data' widgets, which are then connected to a 'Data Table' widget and a 'kNN' widget. The 'kNN' widget's configuration window is open, showing the following settings:

- Name: kNN
- Neighbors: Number of neighbors: 5
- Metric: Euclidean
- Weight: Uniform
- Apply Automatically: ☒

At the bottom of the widget palette, the 'kNN' widget is highlighted, with a description: 'Predict according to the nearest training instances.' and a link to 'more...'.

File Edit View Widget Options Help

Data

Visualize

Tree Viewer

Box Plot

Distribu...

Scatter Plot

Line Plot

Sieve Diagram

Mosaic Display

FreeViz

Linear Projection

Radviz

Heat Map

Venn Diagram

Silhouette Plot

Pythago... Tree

Pythago... Forest

CN2 Rule Viewer

Nomogr...

Linear Projection

A multi-axis projection of data onto a two-dimensional plane.

[more...](#)

File

Data

Data

Data

Data

Data

kNN

Predictions

Linear Projection

Data

Data

Data

Data

Model → Predictors

Predictions → Data

17

Predictions

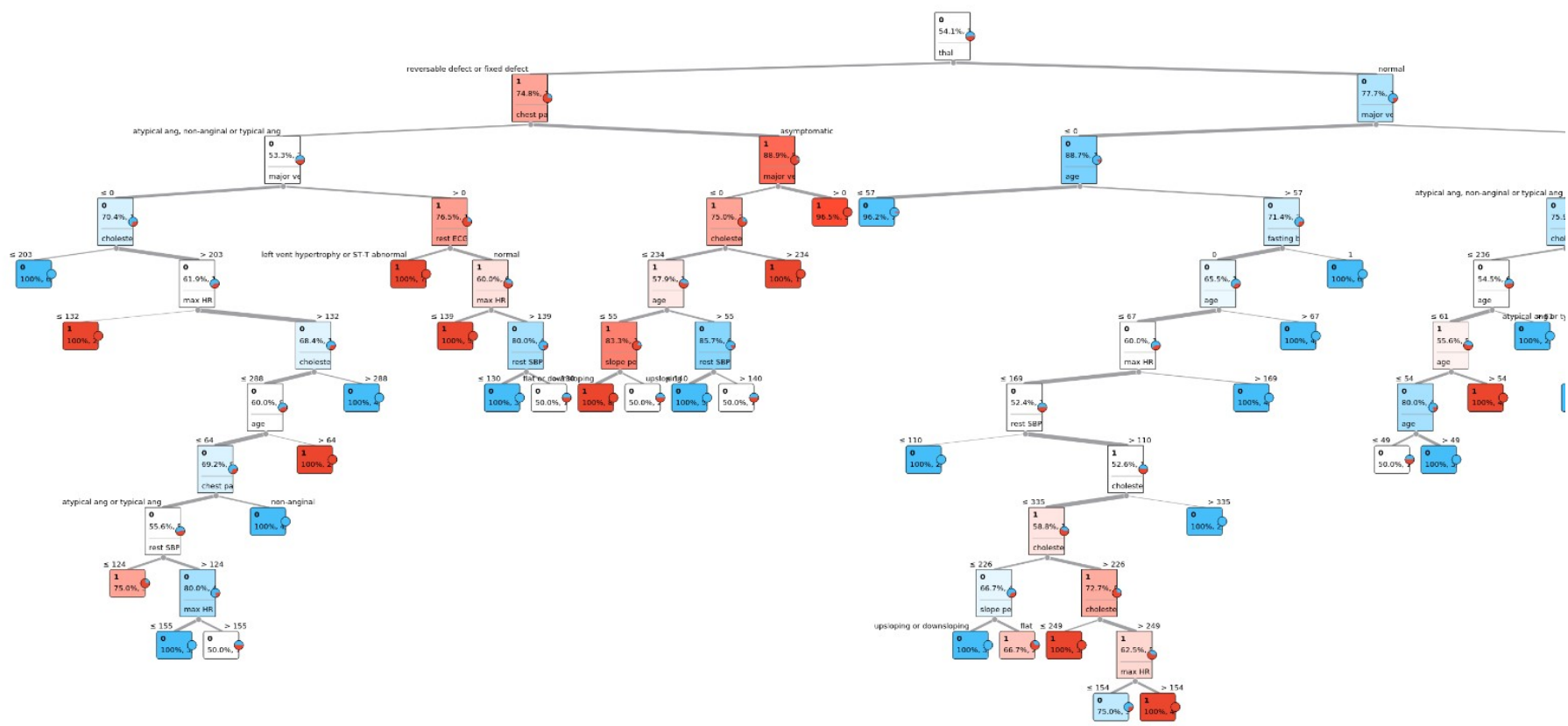
Show probabilities for

0

1

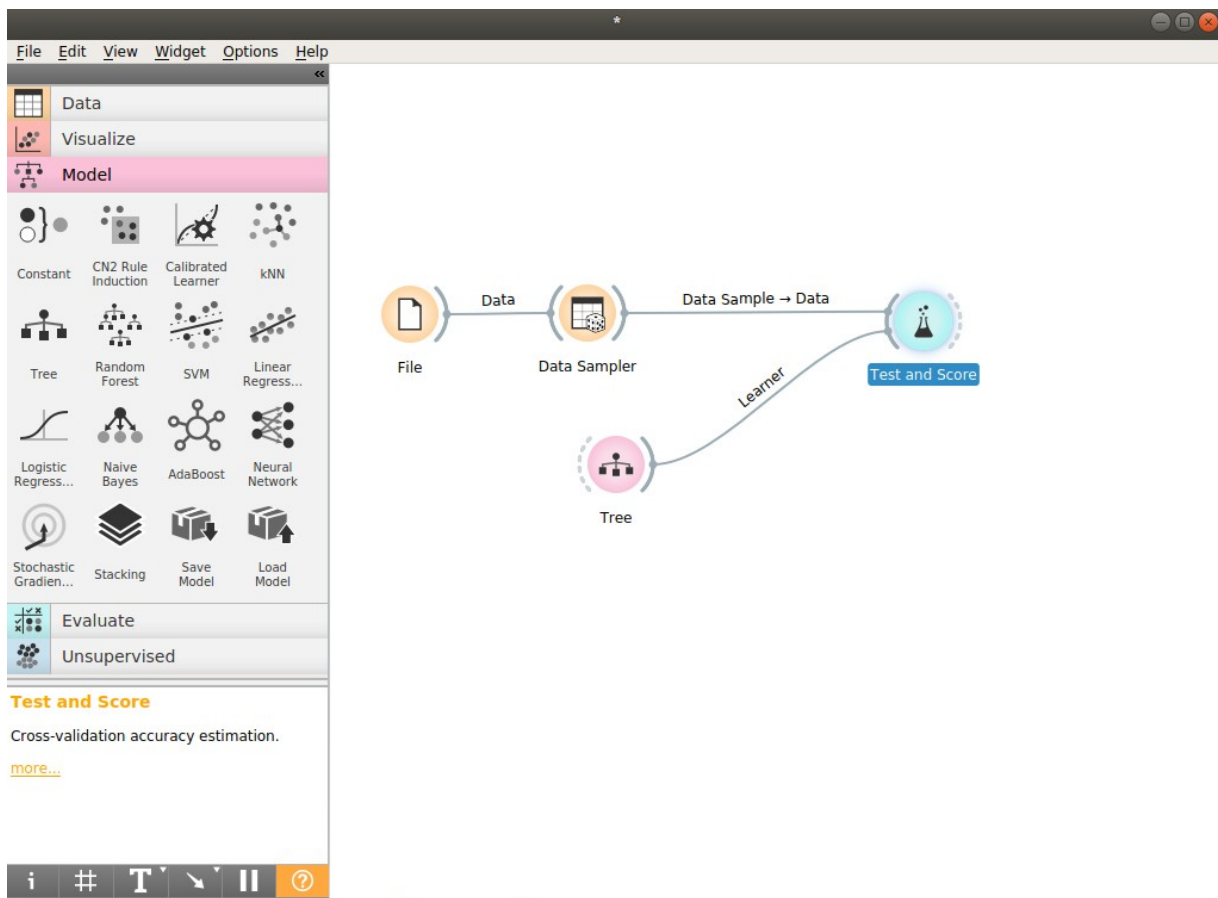
	kNN	ameter narrowir	age	gender	chest pain	rest SBP	cholesterol	ig blood sugar >	rest ECG	max HR	exerc ind ang	ST by exercise	lope peak exc S	lor vessels color	thal
1	0.60 : 0.40 → 0	0	63	male	typical ang	145	233	1	left vent hyp...	150	0	2.3	downsloping	0	fixed defect
2	0.00 : 1.00 → 1	1	67	male	asymptomatic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

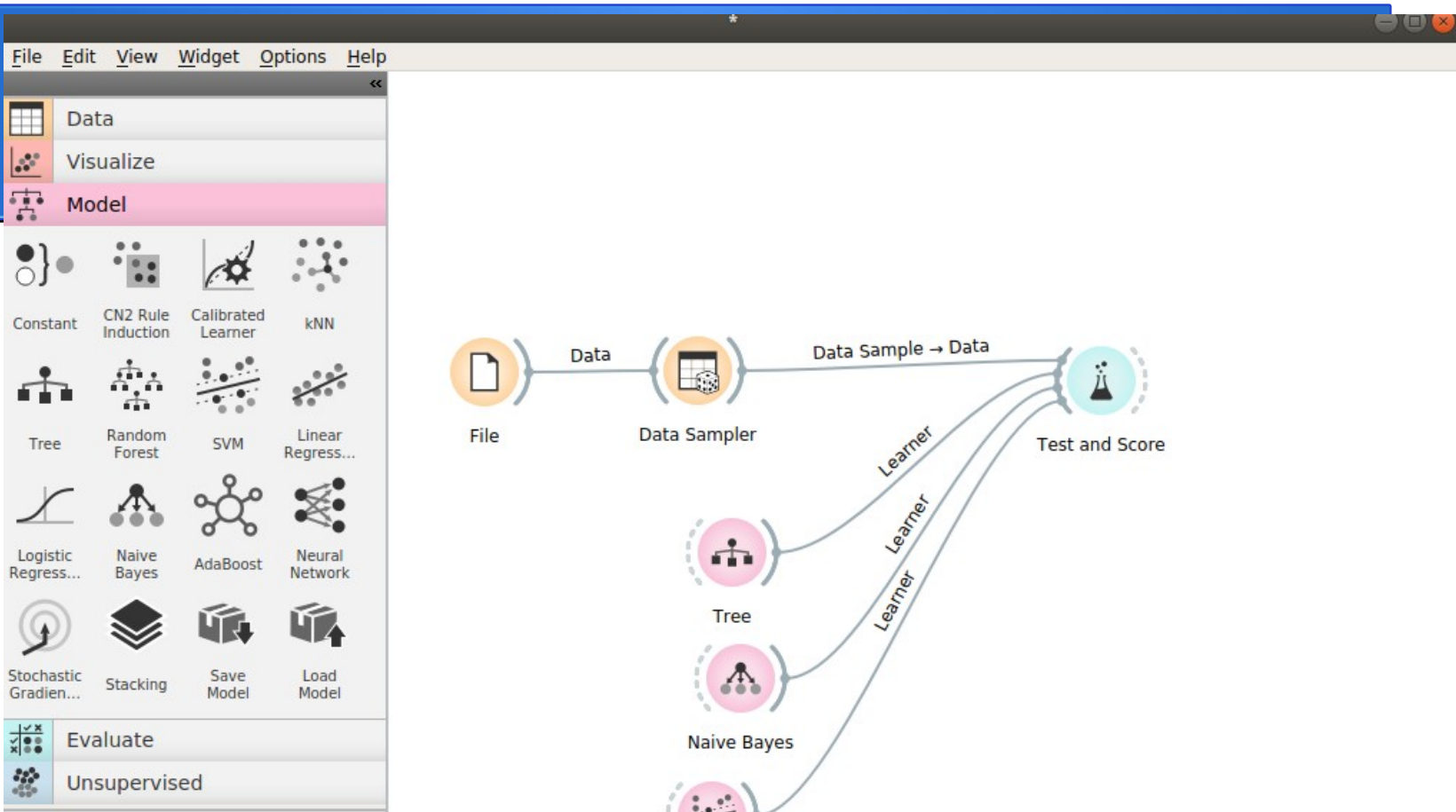
Tree size: 77 nodes, 39 leaves
Edge widths: Relative to parent
Target class: None



Write a comment...

Validação Cruzada

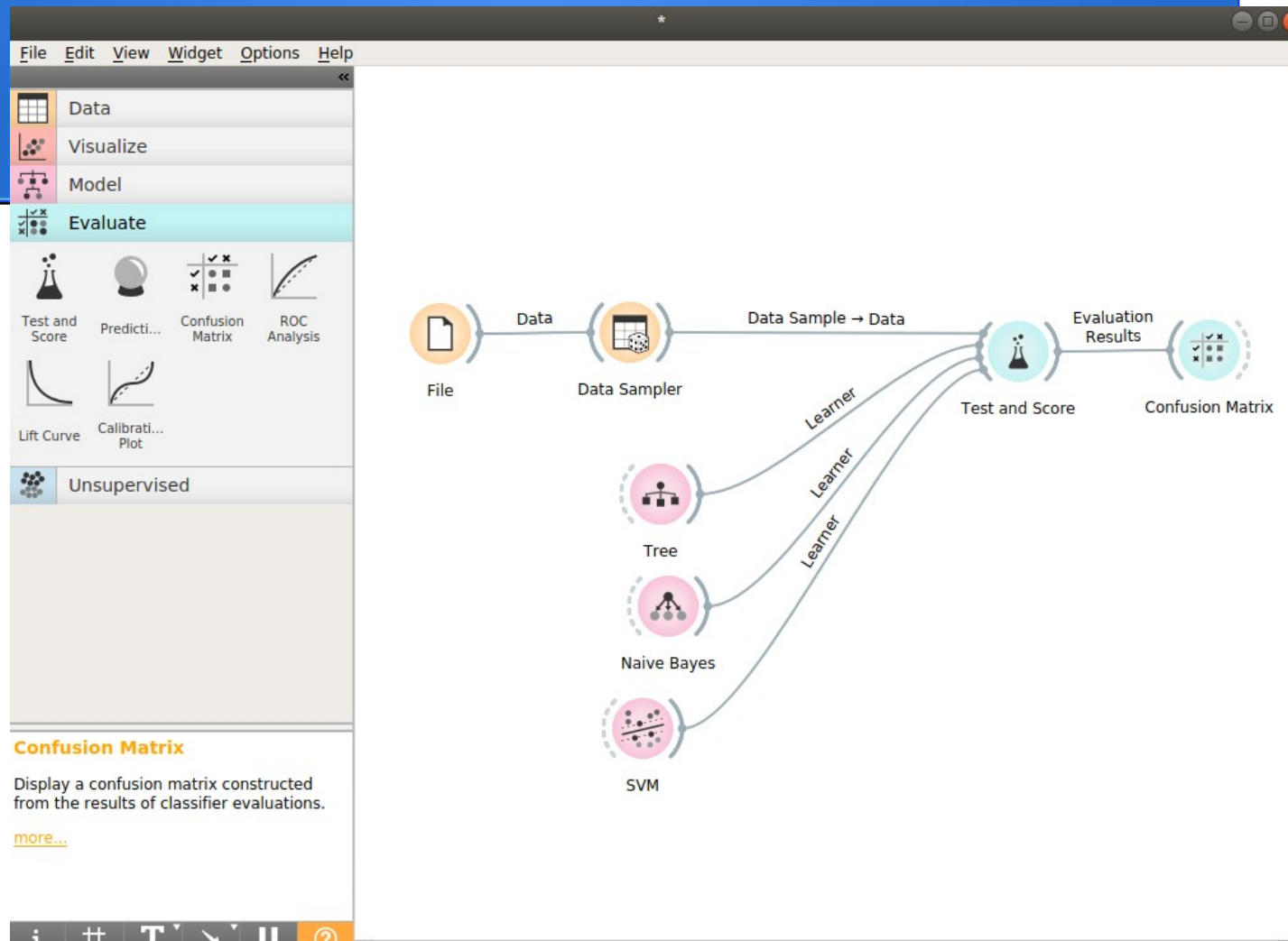




Test and Score

Cross-validation accuracy estimation.

[more...](#)



FileEditViewWidgetOptionsHelp

Data

Visualize

Model

Evaluate

Test and Score

Predicti...

Confusion Matrix

ROC Analysis

Lift Curve

Calibrati... Plot

Unsupervised

ROC Analysis

Display the Receiver Operating Characteristics curve based on the evaluation of classifiers.

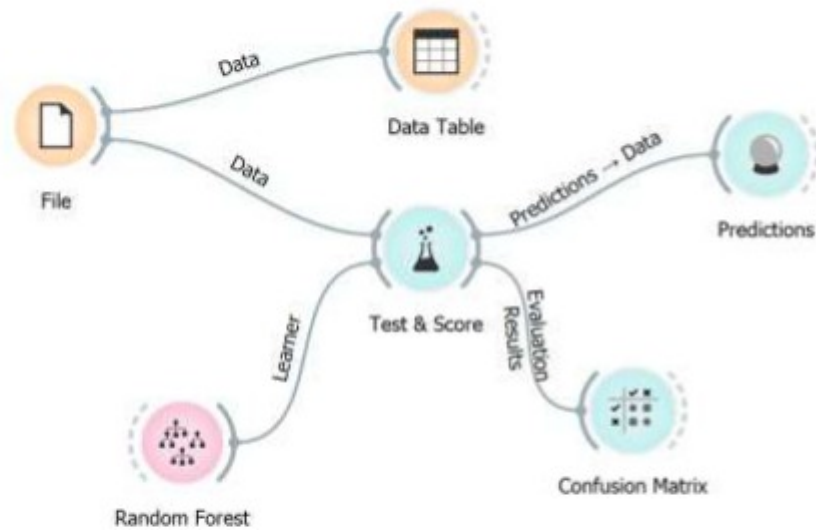
[more...](#)

```
graph LR; File[File] -- Data --> DS[Data Sampler]; DS -- "Data Sample -> Data" --> TS[Test and Score]; TS -- "Learner" --> Tree[Tree]; TS -- "Learner" --> NB[Naive Bayes]; TS -- "Learner" --> SVM[SVM]; TS -- "Evaluation Results" --> CM[Confusion Matrix]; TS -- "Evaluation Results" --> ROC[ROC Analysis];
```

The diagram illustrates a machine learning workflow. It begins with a 'File' icon, which leads to a 'Data' label. This data is then processed by a 'Data Sampler' icon, resulting in a 'Data Sample -> Data' label. This sampled data is then fed into a 'Test and Score' icon. From 'Test and Score', three paths emerge, each labeled 'Learner', leading to 'Tree', 'Naive Bayes', and 'SVM' icons. Additionally, two paths emerge from 'Test and Score', both labeled 'Evaluation Results', leading to a 'Confusion Matrix' icon and an 'ROC Analysis' icon (highlighted with a blue box).

11 [more...](#)

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Referências

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