

4086/1 – APRENDIZAGEM DE MÁQUINA

01/2021

Prof. Dra. Valéria Delisandra Feltrim

Doutorado Acadêmicos em Ciência da Computação

Aluno: Jéfer Benedett Dorr

pg54802@uem.br

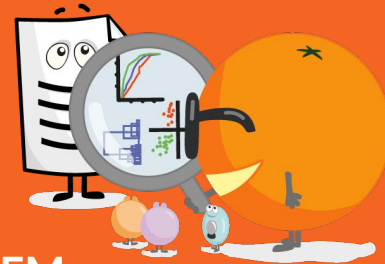




Ferramenta no-code, produtiva e divertida para ensino lúdico dos conceitos de Aprendizagem de Máquina

Prof. Me. Jéfer Benedett Dörr

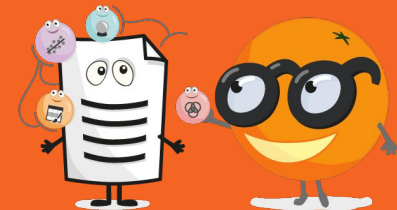
- Graduado em Informática pela UNIOESTE
- Especialista em Software Livre pela Unipan
- Mestre em Informática pela UFPR
- Doutorando em Ciência da Computação - UEM



ORANGE
is the
new
BLACK



Docente do Curso de
Licenciatura em Computação
da UFPR/Palotina



<https://orangedatamining.com/>

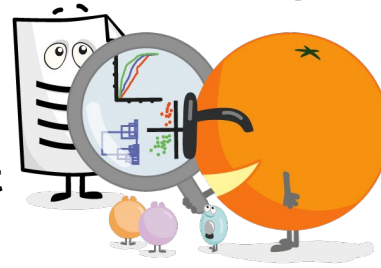


Orange Data Mining - Fruitful & Fun



Atividade prática, mão da massa, de Machine Learning para quem não sabe nada de Inteligência Artificial. Uso de uma ferramenta gráfica, no coding, onde serão apresentados conceitos e realizados exemplos utilizando alguns dos principais algoritmos de aprendizagem de máquina, como:

- Naive Bayes
- Classificador de Máxima Entropia
- Redes neurais
- KNN
- k-means
- SVM
- Random Forest



links para acompanhar

[colab - fofo](#)

[dataset principal](#)

[pasta](#)

[modelo teachable machine](#)

[colab - teachable machine](#)

[colab - cat vs dog](#)

Machine Learning é uma técnica de Inteligência Artificial que permite que a máquina aprenda através de exemplos

Exemplo: PROBLEMA identificar o que é um gato e por eliminação, se o bichinho não for um gato, vamos identificá-lo como cachorro.

características do nosso gato referência:

- É fofo?
- Tem orelhinha pequena?
- Faz miau?



cat	fofo?	orelha pequena?	mia?
cat	1	1	1
cat	1	0	1
cat	0	1	1



dog	fofo?	orelha pequena?	mia?
dog	1	1	0
dog	0	1	0
dog	0	1	0

#Dataset:

bichinho1 = [1, 1, 1]

bichinho2 = [1, 0, 1]

bichinho3 = [0, 1, 1]

bichinho4 = [1, 1, 0]

bichinho5 = [0, 1, 0]

bichinho6 = [0, 1, 0]

#label

#1 = Gato

#-1 = Cachorro

marcacoes = [1, 1, 1, -1, -1, -1]

dados = [bichinho1, bichinho2, bichinho3,
bichinho4, bichinho5, bichinho6]

Create model - Naive Bayes

```
!pip install scikit-learn
from sklearn.naive_bayes import MultinomialNB

modelo = MultinomialNB()
modelo.fit(dados, marcacoes)
```

Naive-Bayes: calcular a probabilidade que uma amostra desconhecida pertença a cada uma das classes possíveis, ou seja, prever (ou adivinhar) a classe mais provável.

Predict

bicho_misterioso1 = [1, 1, 1]

bicho_misterioso2 = [1, 0, 0]

bicho_misterioso3 = [0, 0, 1]

-	fofo?	orelha pequena?	mia?
?	1	1	1
?	1	0	0
?	0	0	1

Resultados

```
teste = [bicho_misterioso1, bicho_misterioso2,  
bicho_misterioso3]  
resultado = modelo.predict(teste)  
print(resultado)
```

Correto?

```
marcacoes_teste = [1, -1, 1]
print('Resultado: ')
print(resultado)
print ('Marcacoes: ')
print(marcacoes_teste)
```

Resultado[Gato, Gato, Gato]

e a resposta correta deveria ser

Marcacoes [Gato,Cachorro,Gato]

algoritmo tem a acurácia de 66,66%, ou seja,

ele acertou 2 bichos misteriosos do nosso total de 3.



curiosidade

O algoritmo “**Naive Bayes**” é um classificador probabilístico baseado no “**Teorema de Bayes**”, o qual foi criado por **Thomas Bayes** (1701 - 1761) para tentar provar a existência de Deus.

Hoje, usado para identificar se determinado e-mail é um **SPAM!**

```
# Criar um classificador
nbytes = NBytes::Base.new
# Treinar o classificador com exemplos - as palavras da String são
#divididas em um array
nbytes.train( "You need to buy some Viagra".split(/\s+/), 'SPAM' )
nbytes.train( "This is not spam, just a letter to Bob.".split(/\s+/), 'HAM' )
nbytes.train( "Hey Oasis, Do you offer consulting?".split(/\s+/), 'HAM' )
nbytes.train( "You should buy this stock".split(/\s+/), 'SPAM' )

# Dividir mensagem que precisa ser classificada
tokens = "Now is the time to buy Viagra cheaply and
discreetly".split(/\s+/)
result = @nbytes.classify(tokens)
# Imprime a classe em que o texto foi classificado. (SPAM ou HAM)
p result.max_class
# Imprime a probabilidade da mensagem ser SPAM
p result['SPAM']
# Imprime a probabilidade da mensagem ser HAM
p result['HAM']
```

exemplo

- 100 pessoas realizaram o teste.
- 20% das pessoas que realizaram o teste possuíam a doença.
- 90% das pessoas que possuíam a doença, receberam positivo no teste.
- 30% das pessoas que não possuíam a doença, receberam positivo no teste.

Se uma nova pessoa realizar o teste e receber um resultado positivo, qual a probabilidade de ela possuir a doença?

O algoritmo de **Naive Bayes** consiste em encontrar uma probabilidade a posteriori (parte do efeito para a causa, ou seja, possuir a doença, dado que recebeu um resultado positivo), multiplicando a probabilidade a priori (da causa para o efeito, ou, possuir a doença) pela probabilidade de “receber um resultado positivo, dado que tem a doença”.

Devemos também computar a probabilidade a posteriori da negação (Não possuir a doença, dado que recebeu um resultado Positivo).

$$P(\text{doença}|\text{positivo}) = 20\% * 90\% = 0,2 * 0,9 = 0,18$$

$$P(\text{doença}|\text{negativo}) = 80\% * 30\% = 0,8 * 0,3 = 0,24$$

normalizando:

$$P(\text{doença}|\text{positivo}) = 0,18 / (0,18 + 0,24) = 0,4285$$

$$P(\text{não doença}|\text{positivo}) = 0,24 / (0,18 + 0,24) = 0,5714$$

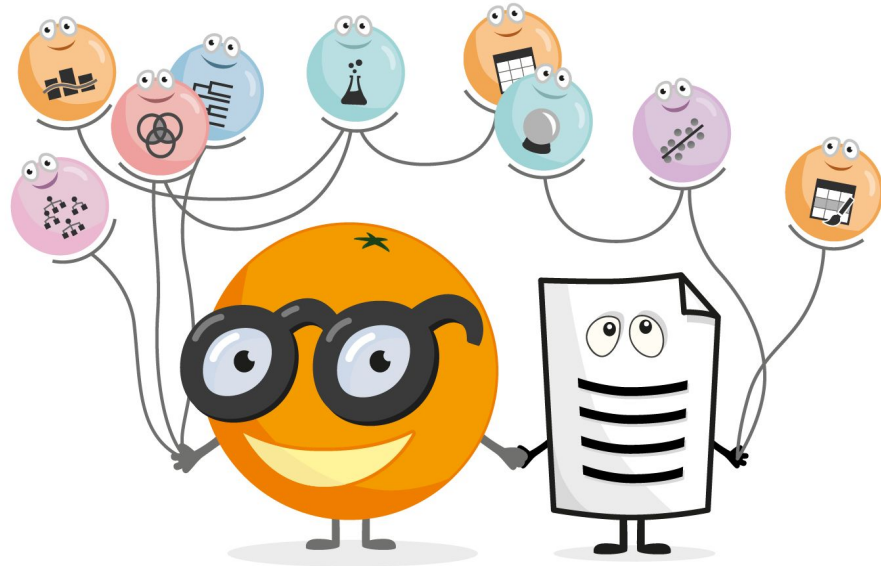
se o resultado do teste da nova pessoa for positivo, ela possui aproximadamente 43% (0,4285) de chance de estar doente.



- Orange is a component-based visual programming software package for data visualization, machine learning, data mining, and data analysis.
 - open-source software GPL
 - Python
 - scientific computing: numpy, scipy and scikit-learn
 - interface: Qt framework
 - *Slovenia*
 - Initial release: 1996
 - Last release: 2020
 - drag and drop
-

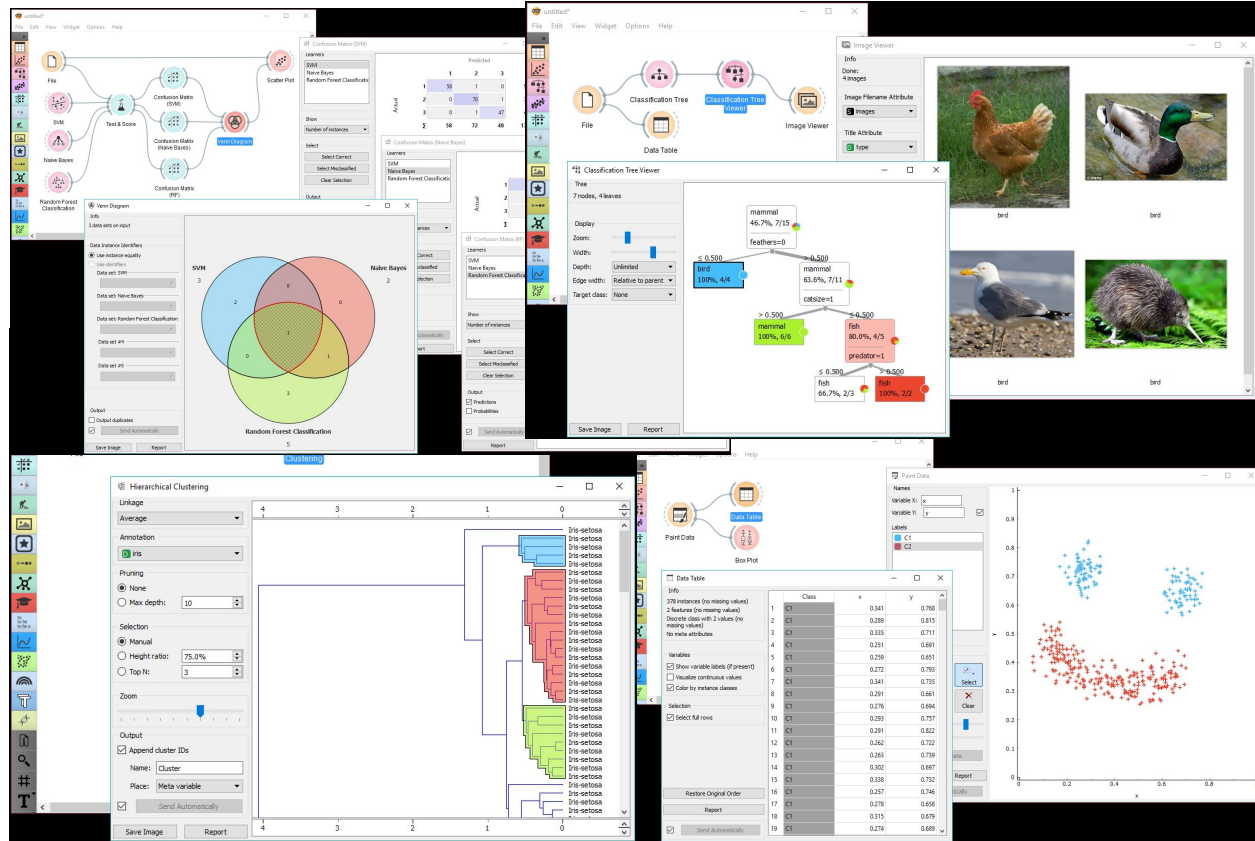
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images



blog

It's Sailing Time (Again)

By: BLAZ, Aug 11, 2017

Every fall I teach a course on Introduction to Data Mining. And while the course is really on statistical learning and its applications, I also venture into [classification trees](#). For several reasons. First, I can introduce [information gain](#) and with it feature scoring and ranking. Second, classification trees are one of the first machine learning approaches co-invented by engineers ([Ross Quinlan](#)) and statisticians ([Leo Breiman](#), [Jerome Friedman](#), [Charles J. Stone](#), [Richard A. Olshen](#)). And finally, because they make the base of random forests, one of the most accurate machine learning models for smaller and mid-size data sets.

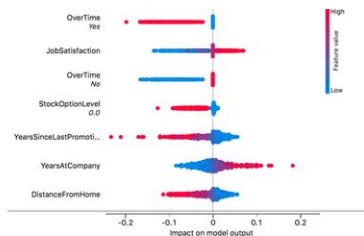
Related: [Introduction to Data Mining Course in Houston](#)

Lecture on classification trees has to start with the data. Years back I have crafted a data set on sailing. Every data set has to have a story. Here is one:

Sara likes weekend sailing. Though, not under any condition. Past twenty Wednesdays I have asked her if she will have any company, what kind of boat she can rent, and I have checked the weather forecast. Then, on Saturday, I wrote down if she actually went to the Sea.

Data on Sara's sailing contains three attributes (Outlook, Company, Sailboat) and a class (Sail).

Data Table				
	Sail	Outlook	Company	Sailboat
1	yes	rainy	big	big
2	yes	rainy	big	small
3	no	rainy	med	big
4	no	rainy	med	small
5	yes	sunny	big	big
6	yes	sunny	big	small
7	yes	sunny	med	big



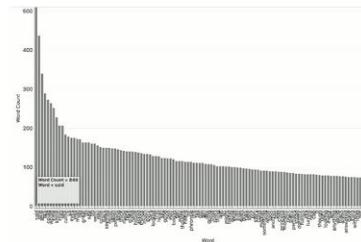
By: Ajda Pretner, Jan 27, 2021

Observing Word Distribution

How to inspect word distribution in a corpus with a clever combination of widgets in Orange.

Categories: [text mining](#) [word distribution](#) [bar plot](#) [word cloud](#)

[Read More](#)



By: Ajda Pretner, Jan 11, 2021

Orange in Classroom

Orange is actively used in classrooms at over two hundred universities from around the world.

Categories: [Orange](#) [education](#) [teaching](#) [university](#)

download



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Workflows

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Blog

Docs

Workshops



Windows



macOS



Linux / Source

Download the latest version for Windows

Download Orange 3.29.1

<https://orangedatamining.com/download/>

Install

Windows: double click - next - next - ok

Linux:

```
sudo apt install virtualenv build-essential python3-dev
```

```
virtualenv --python=python3 --system-site-packages orange3venv
```

```
pip install PyQt5 PyQtWebEngine
```

```
source bin/activate
```

```
pip install orange3
```

anaconda

 ANACONDA NAVIGATOR

Sign in

Home

Environments

Projects (beta)

Learning

Community








Documentation

Developer Blog

Feedback

Twitter YouTube GitHub

Applications on Channels

 <p>glueviz</p> <p>0.10.4</p> <p>Multidimensional data visualization across files. Explore relationships within and among related datasets.</p> <p>Launch</p>	 <p>jupyter notebook</p> <p>5.0.0</p> <p>Web-based, interactive computing notebook environment. Edit and run human-readable code while describing the data analysis process.</p> <p>Launch</p>	 <p>orange3</p> <p>3.4.1</p> <p>Component based data mining framework. Data visualization and data analysis for novice and expert. Interactive workflows with a large toolbox.</p> <p>Launch</p>	 <p>IPyQt</p> <p>qtconsole</p> <p>4.3.0</p> <p>IPyQt GUI that supports inline figures, proper multiline editing with syntax highlighting, graphical calltips, and more.</p> <p>Launch</p>
 <p>rstudio</p> <p>1.1.383</p> <p>A set of integrated tools designed to help you be more productive with R. Includes R essentials and notebooks.</p> <p>Launch</p>	 <p>spyder</p> <p>3.1.4</p> <p>Scientific Python Development Environment. Powerful Python IDE with advanced editing, interactive testing, debugging and introspection features</p> <p>Launch</p>	 <p>jupyterlab</p> <p>0.31.8</p> <p>Install</p>	

examples

Info

164 instances
10 features (7.0 % missing data)
Target with 3 values
5 meta attributes (7.1 % missing data)

Variables

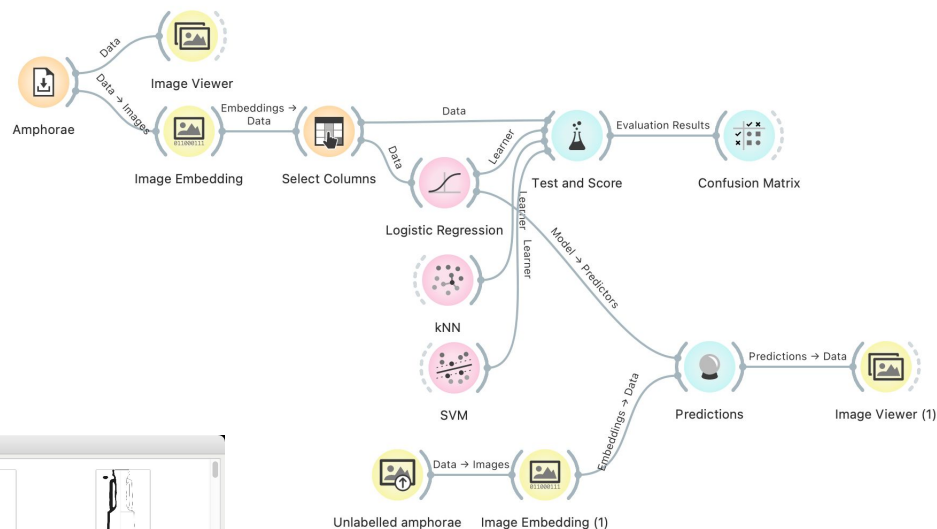
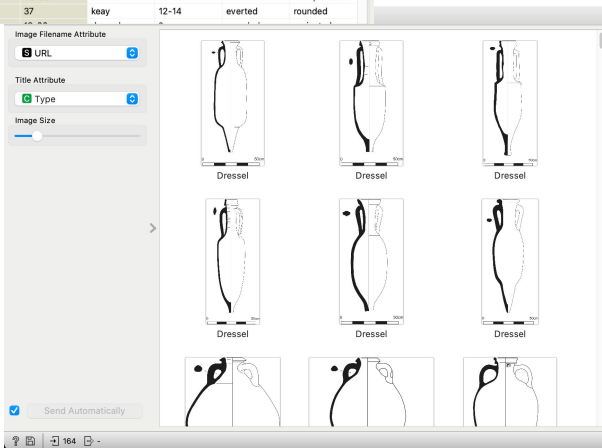
- Show variable labels (if present)
- Visualize numeric values
- Color by instance classes

Selection

- Select full rows

Restore Original Order

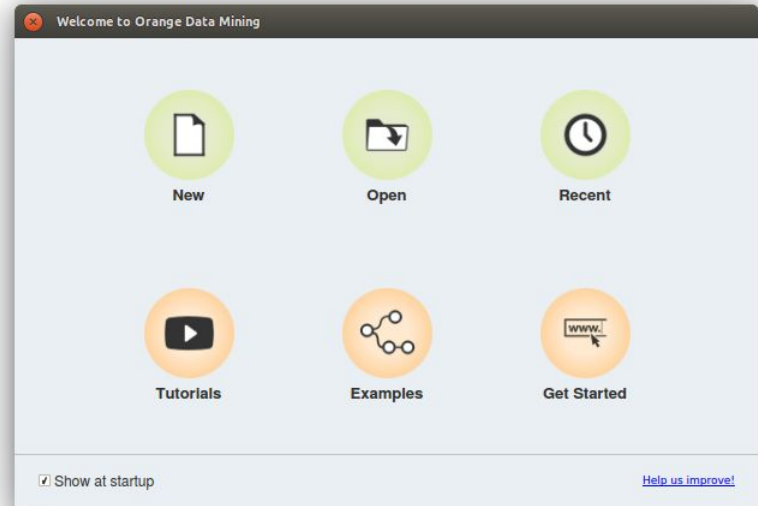
	Type	URL	Name	Subtype	Height (cm)	Width (cm)	Fabric	Rim diameter	Rim type	Shoulder type
1	Dressel	file.biola.../...	DR101	Dressel 14	90-110	27-33	dressel	16-22	beaded	rounded
2	Dressel	file.biola.../...	DR104	Dressel 1	97-117	28-29	campanian	17-18	collar	carinated
3	Dressel	file.biola.../...	DR105	Dressel 1	97-117	28-29	campanian	17-18	collar	carinated
4	Dressel	file.biola.../...	DR106	Dressel 1	97-117	28-29	campanian	17-18	collar	carinated
5	Dressel	file.biola.../...	DR109	Dressel 12	103-105	25-30	dressel	16-18	everted	rounded
6	Dressel	file.biola.../...	DR110	Dressel 14	90-110	27-33	dressel	16-22	beaded	rounded
7	Dressel	file.biola.../...	DR111	Dressel 20	70-97	56-66	dressel	18-20	rounded	none/smooth
8	Dressel	file.biola.../...	DR113	Dressel 23	53-56	24-36	dressel	10-14	triangular	none/smooth
9	Dressel	file.biola.../...	DR114	Dressel 28	?	?	dressel	?	pulley wheel	rounded
10	Dressel	file.biola.../...	DR127	Dressel 6B	85-95	34-37	dressel	15-16	rounded	none/smooth
11	Dressel	file.biola.../...	DR135	Dressel 21-22	88-90	24-26	dressel	18-20	beaded	rounded
12	Dressel	file.biola.../...	DR137	Dressel 6A	95-96	34-35	?	16-17	rounded	carinated
13	Dressel	file.biola.../...	DR138	Dressel 6B	85-95	34-37	dressel	15-16	rounded	none/smooth
14	Dressel	file.biola.../...	DR139	Dressel 2-4	95-108	26-29	campanian	14-16	beaded	carinated
15	Dressel	file.biola.../...	DR140	Dressel 20	70-97	56-66	dressel	18-20	rounded	none/smooth
16	Dressel	file.biola.../...	DR141	Dressel 20	70-97	56-66	dressel	18-20	rounded	none/smooth
17	Dressel	file.biola.../...	DR149	Dressel 30	67-69	37	keay	12-14	everted	rounded



Predictions

	image name	image	size	width
Regression	dressel	dressel.png	16893	167
l	gauloise	gauloise.png	21614	268
se	keay	keay.png	25827	217

use



menus

File Edit View Widget Options H

Data

File	Datasets	SQL Table	Data Table
Paint Data	Data Info	Data Sampler	Select Columns
Select Rows	Rank	Correl...	Merge Data
Conca...	Select by Dat...	Trans...	Rando...

Visualize

Tree Viewer	Box Plot	Distrib...	Scatter Plot
Line Plot	Sieve Diagram	Mosaic Display	FreeViz
Linear Projec...	Radviz	Heat Map	Venn Diagram
Silhou... Plot	Pythag... Tree	Pythag... Forest	CN2 Rule V...
Nomo...			

Model

Constant	CN2 Rule I...	kNN	Tree
Random Forest	SVM	Linear Regre...	Logistic Regre...
Naive Bayes	AdaBo...	Neural Network	Stoch... Gradie...
Stacking	Save Model	Load Model	

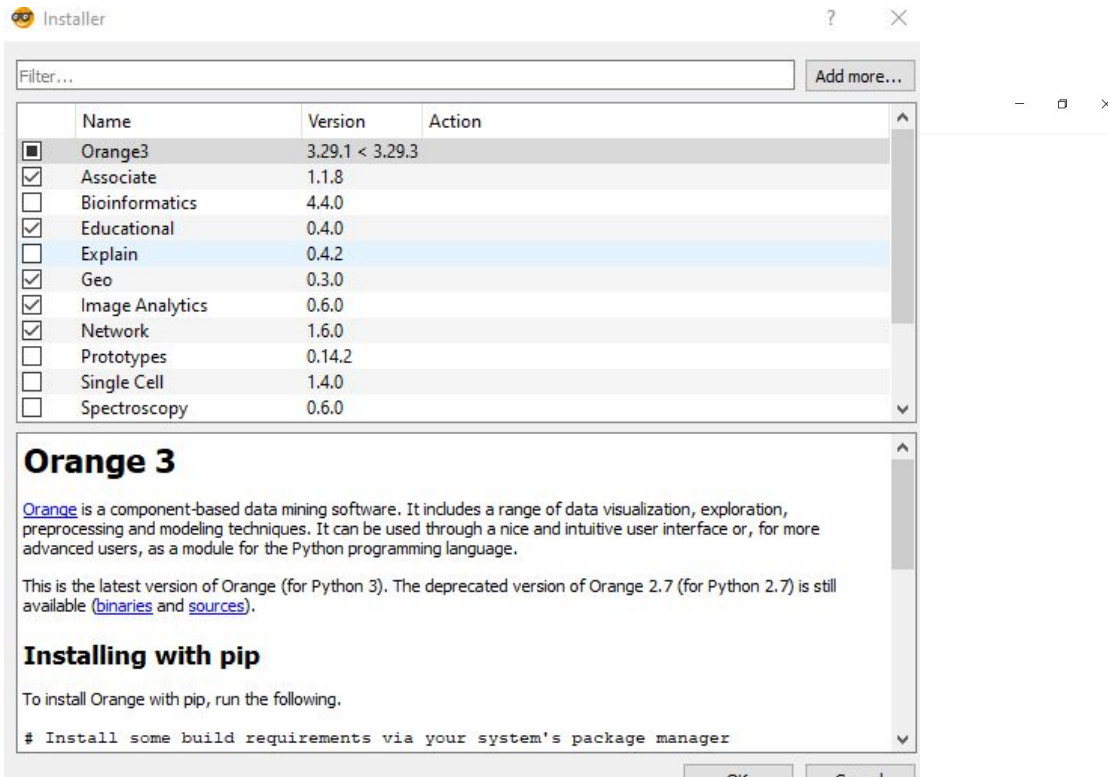
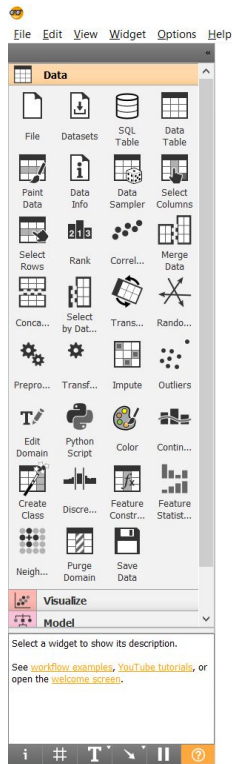
Evaluate

Test & Score	Predic...	Confu... Matrix	ROC Analysis
Lift Curve	Calibr... Plot		

Unsupervised

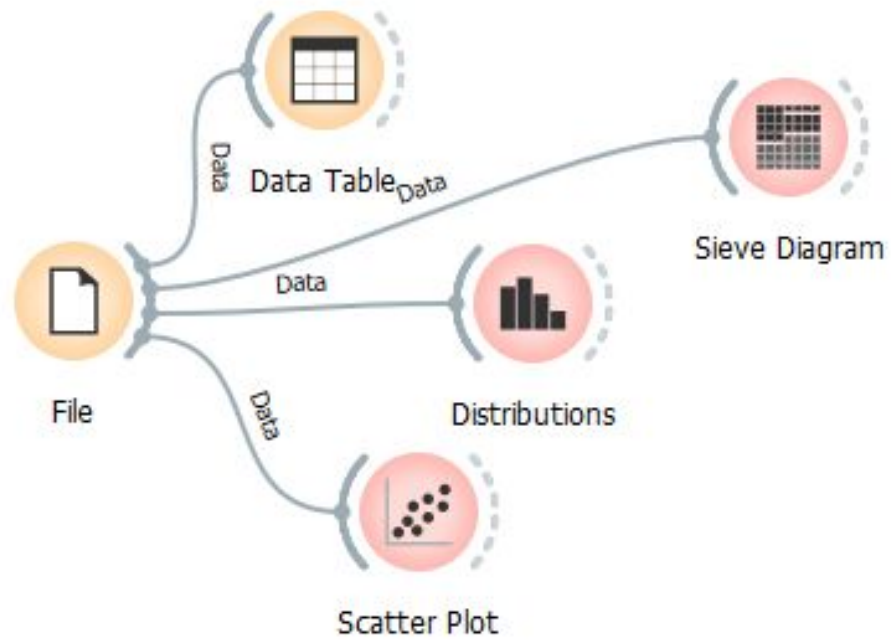
Distance File	Distance Matrix	t-SNE	Distance Map
Hierar... Cluste...	k-Means	Louvain Cluste...	Manifold Learning
PCA	Corres... Analysis	Distan...	Distance Transf...
MDS	Save Distan...		

addons

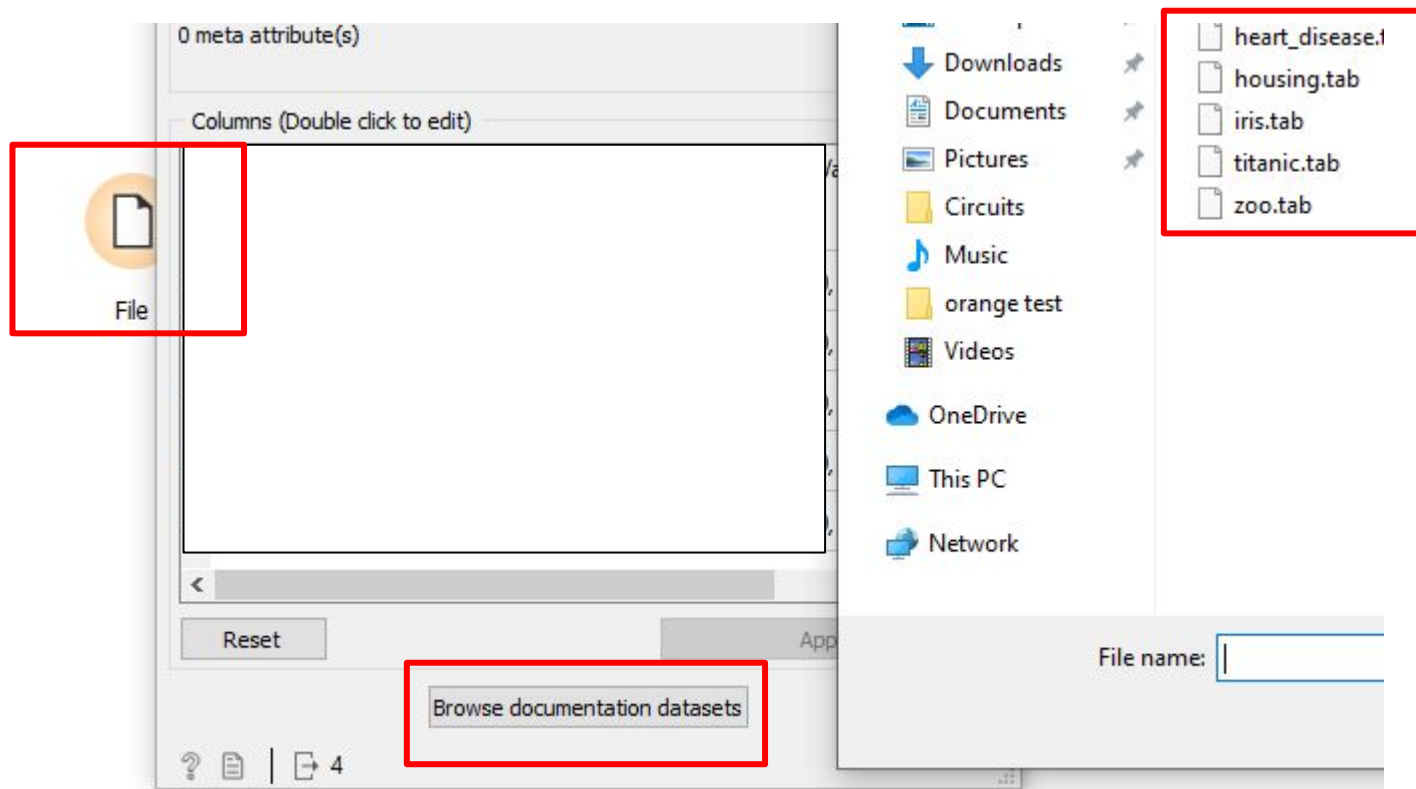


—

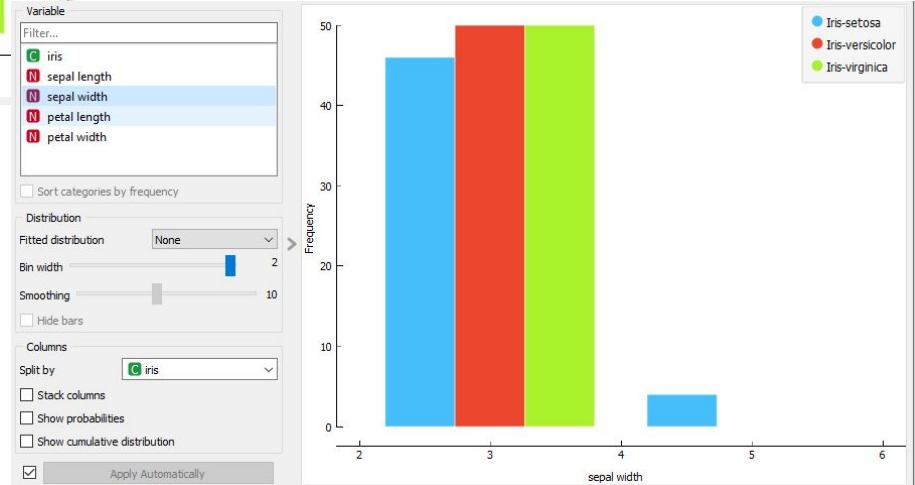
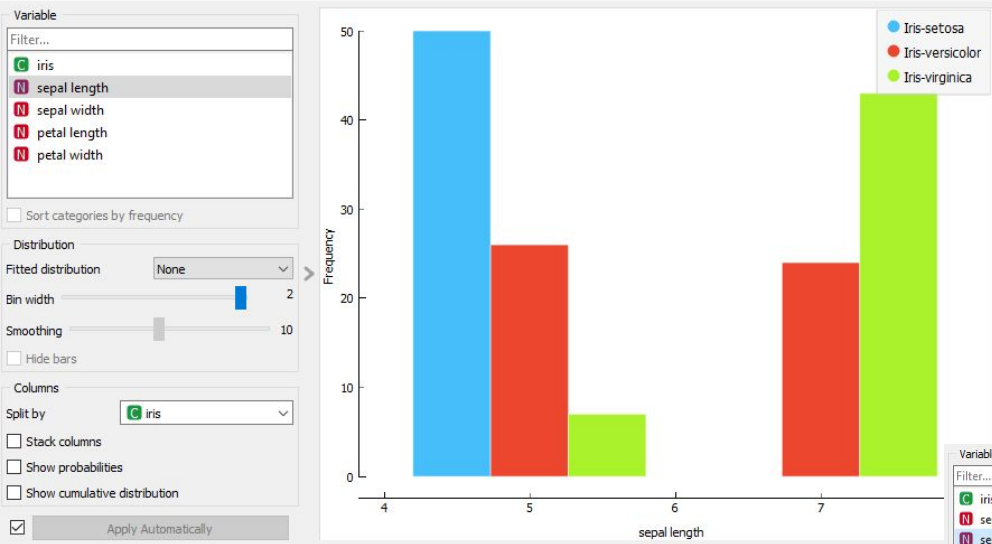
basic use



documentation datasets



Distributions

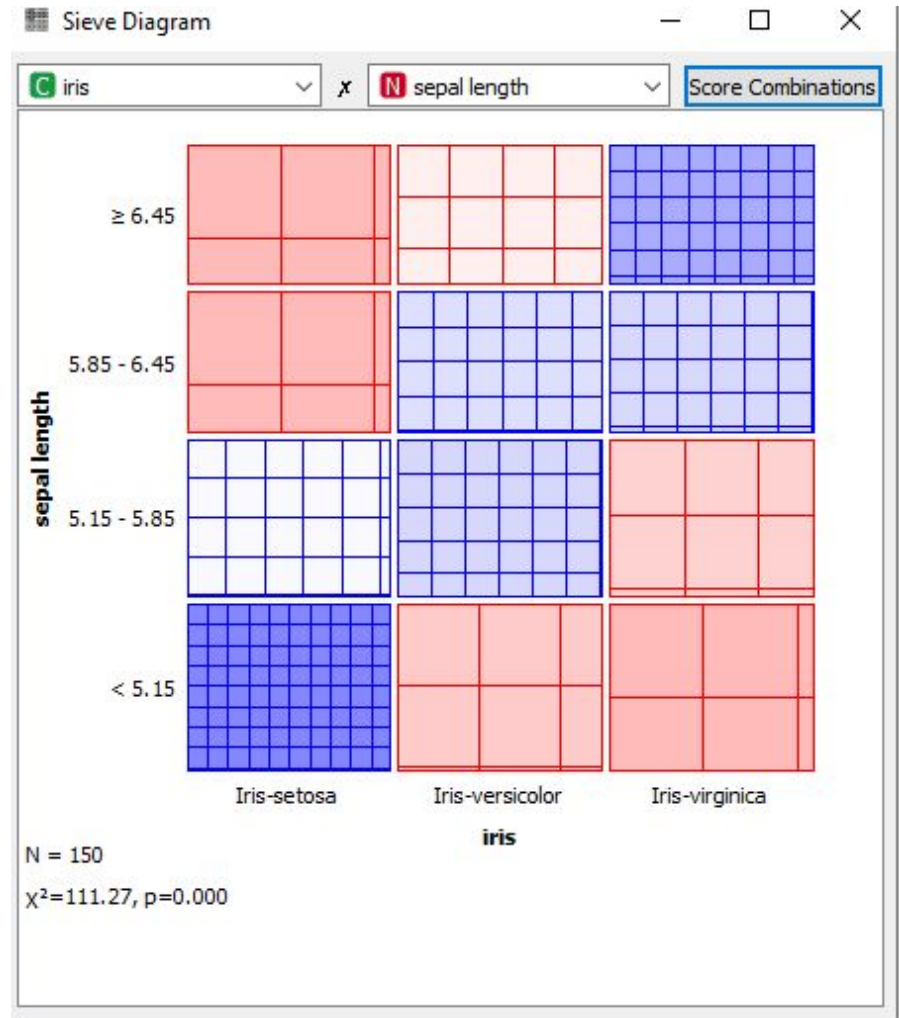


Sieve Diagram

[Sieve Diagram](#)

summarizes the relationship between the categorical variables using frequencies

It was proposed by Riedwyl and Schüpbach in a technical report in 1983 and later called a parquet diagram (Riedwyl and Schüpbach 1994).



—

end basic

atividade 1

Atividade 1

[monet or manet - inceptionv3 \(google\)](#)

[the lost monet - model pre trained painters](#)

—

end atividade 1

Machine Learning: cat or dog?



dataset

<https://www.kaggle.com/tongpython/cat-and-dog>

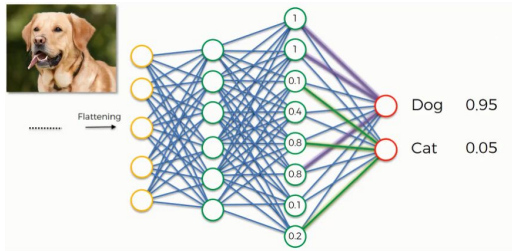


kaggle

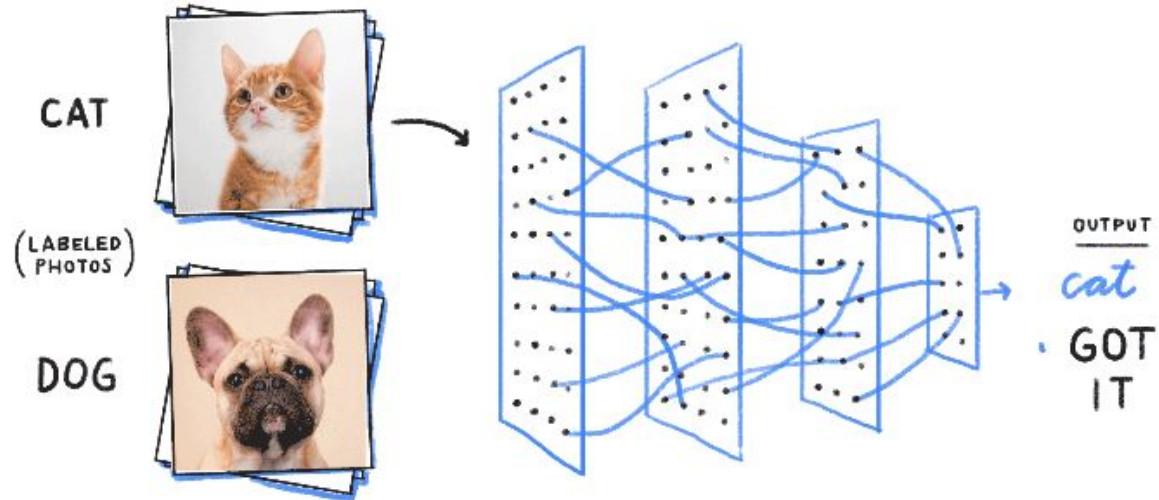
This dataset is for running the code from this site:

<https://becominghuman.ai/building-an-image-classifier-using-deep-learning-in-python-totally-from-a-beginners-perspective-be8dbaf22dd8>.

solve an image classification problem



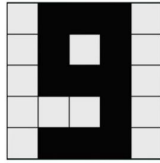
como uma rede neural aprende



training an artificial neural network on few thousand images of cats and dogs and make the Neural Network learn to predict which class the image belongs to, next time it sees an image having a cat or dog in it.

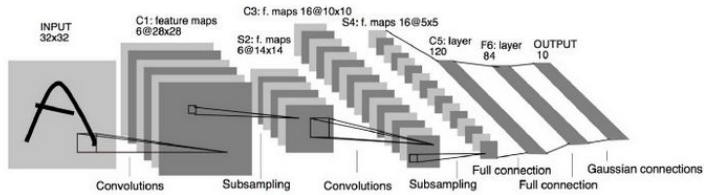
rede neural

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

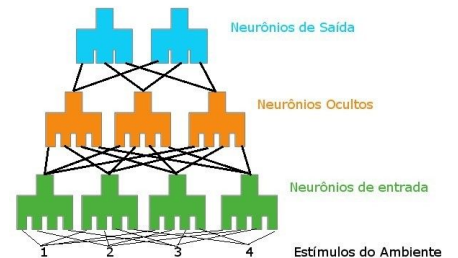
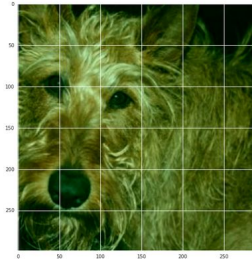
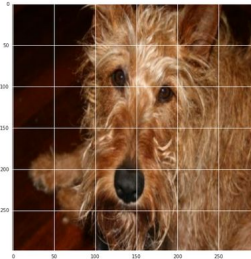


primeira aplicação com sucesso de uma CNN foi desenvolvida por [Yann LeCun em 1998](#)

0,1,1,1,0,0,1,0,1,0,0,1,1,1,0,0,0,0,1,0,0,1,1,1,0,9

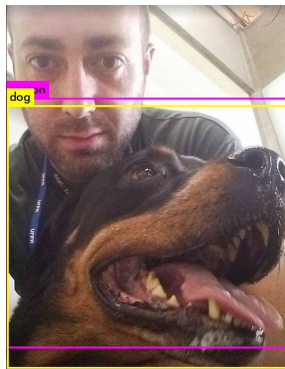
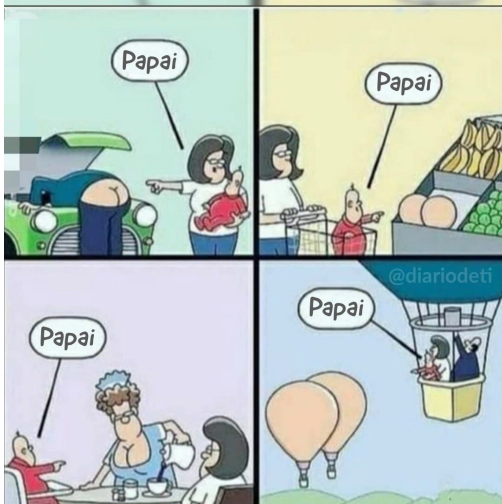


Arquitetura da rede LeNet5

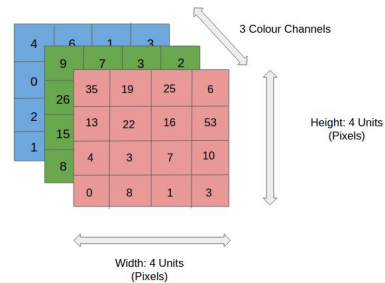


colored images - rgb

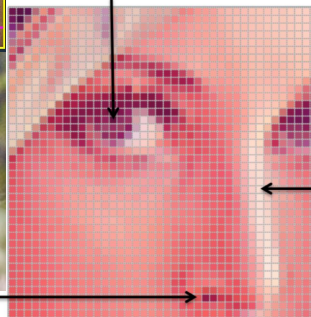
COMO FUNCIONA O MACHINE LEARNING



[213, 60, 67]



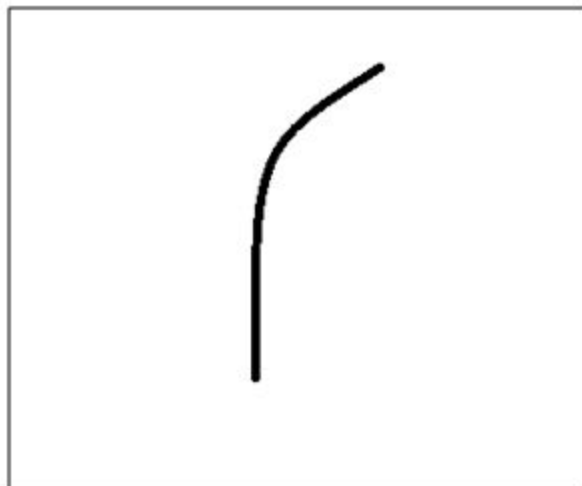
[90, 0, 53]



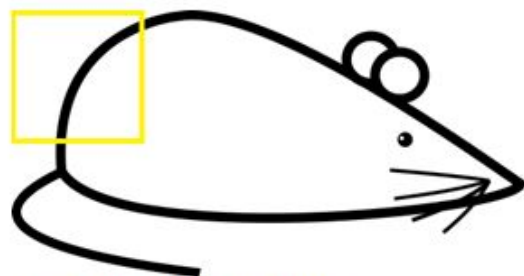
[249, 215, 203]

0	0	0	0	0	30	0
0	0	0	0	30	0	0
0	0	0	30	0	0	0
0	0	0	30	0	0	0
0	0	0	30	0	0	0
0	0	0	30	0	0	0
0	0	0	0	0	0	0

Pixel representation of filter



Visualization of a curve detector filter



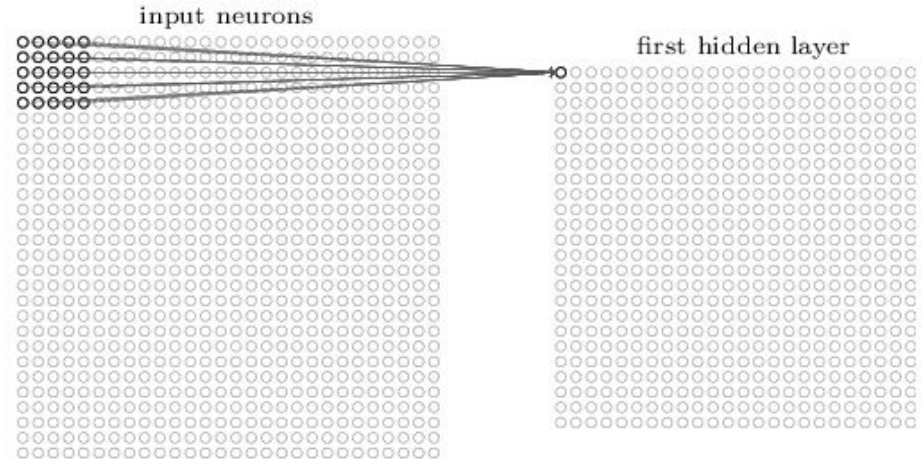
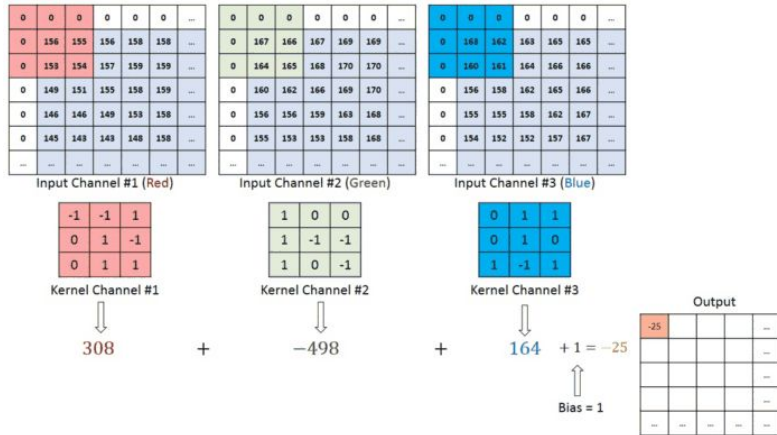
Visualization of the filter on the image



Convoluções

<https://medium.com/neuronio-br/entendendo-redes-convolucionais-cnn-s-d10359f21184>

filtros = kernel formados por pesos e atualizados por backpropagation



Entrada de 28×28 dimensões com receptive field de área 5×5.

1 - The input image is the image being detected. The feature detector is a matrix, usually 3x3. A feature detector is also referred to as a kernel or a filter.

CNN (Convolutional Neural Network)

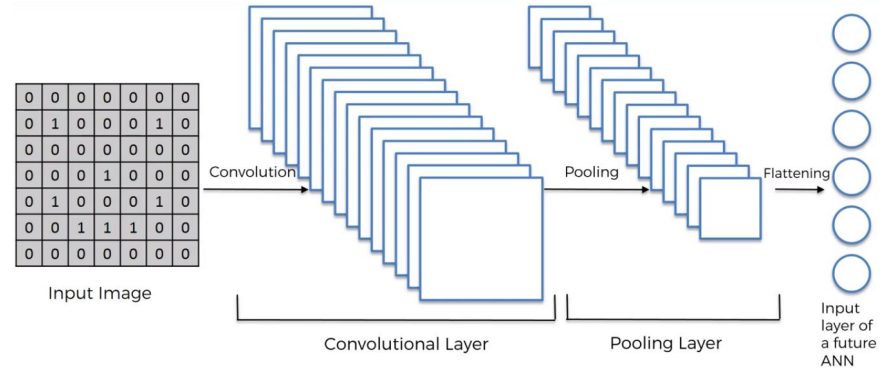
The process of building a Convolutional Neural Network always involves four major steps.

Step - 1 : Convolution

Step - 2 : Pooling

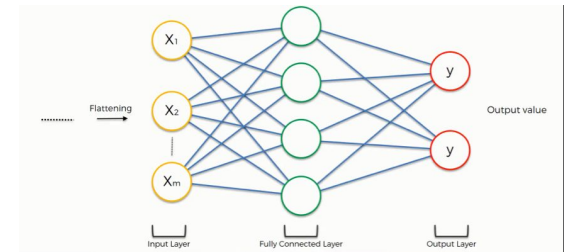
Step - 3 : Flattening

Step - 4 : Full connection

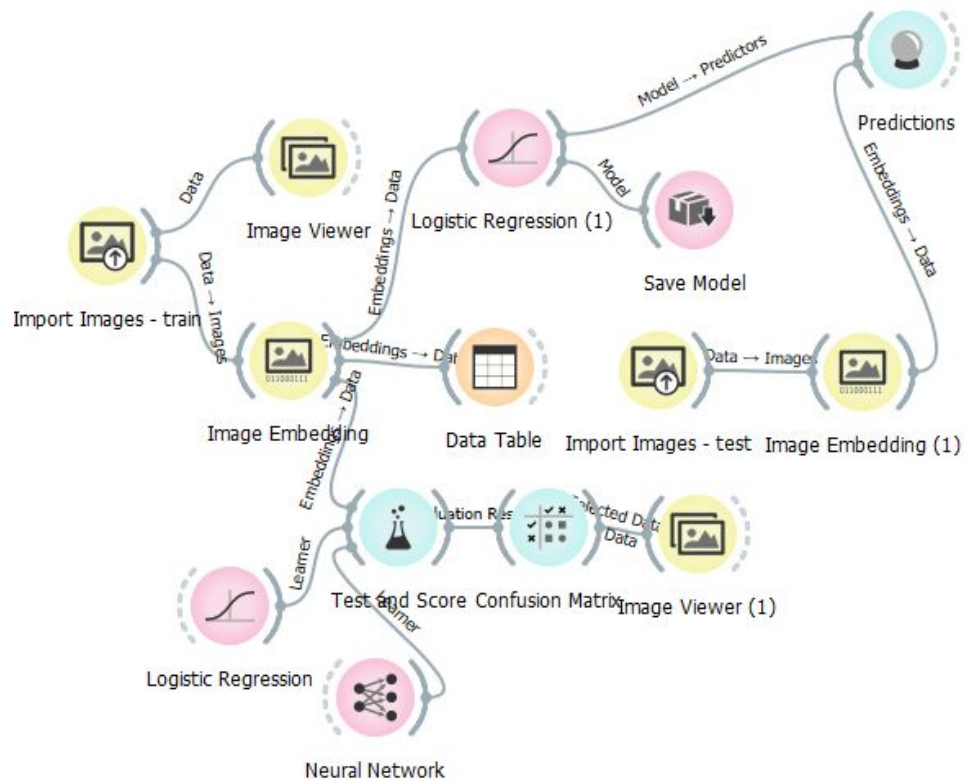
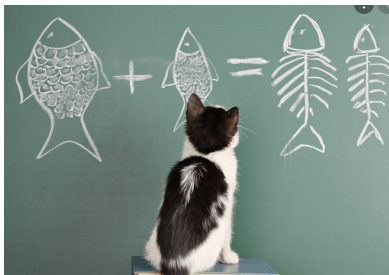


compile and Train model

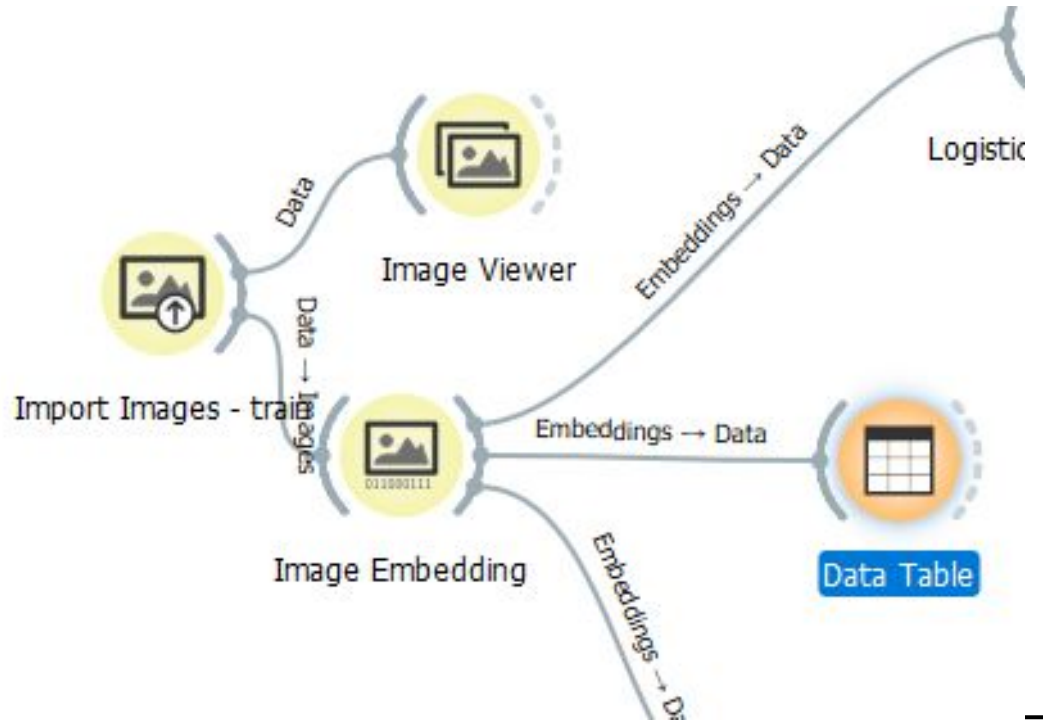
new predictions from trained model



this



import - view - infos



folders named “training_set” and “test_set”

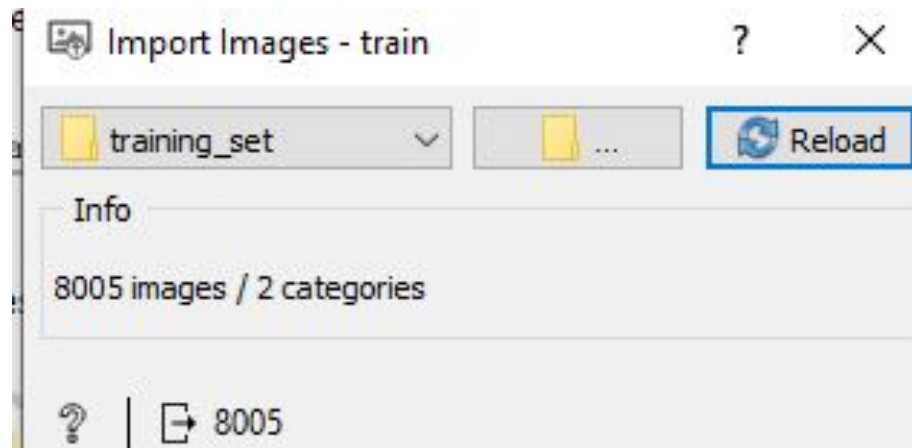




Image Viewer

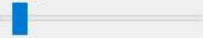
Image Filename Attribute

S image

Title Attribute

C category

Image Size



cats



cats



cats



cats



cats



cats



cats



cats



cats



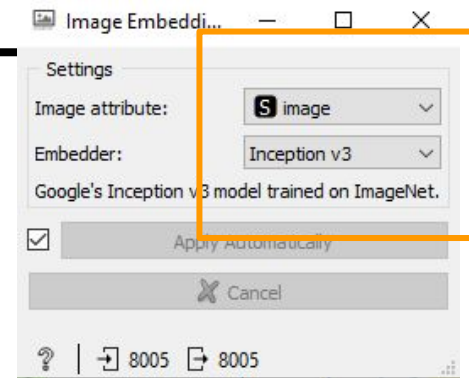
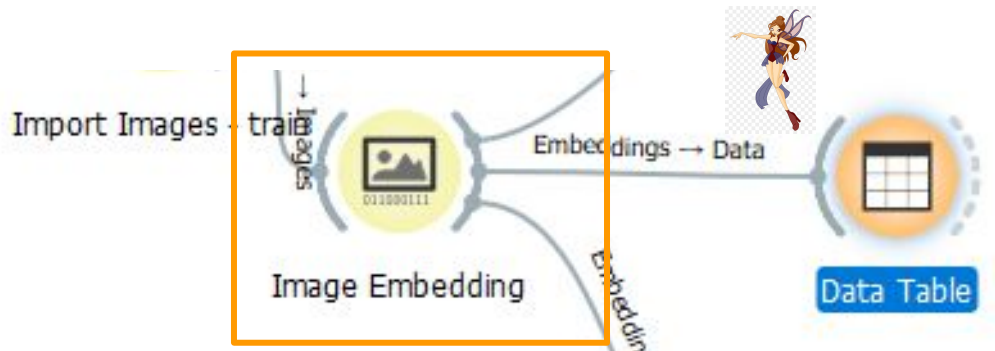
cats



cats



cats



Inception v3: [Google's Inception v3](<https://arxiv.org/abs/1512.00567>)

Image embedding through deep neural networks.

Inputs: Images: List of images.

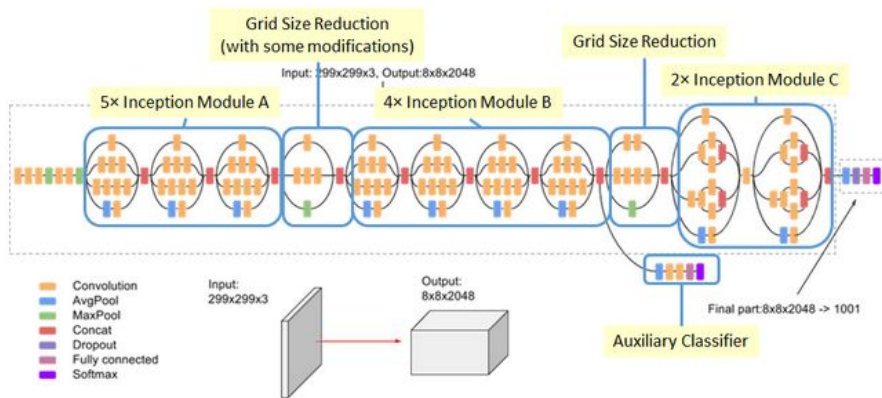
Outputs: Embeddings: Images represented with a vector of numbers.

Skipped Images: List of images where embeddings were not calculated.

Image Embedding reads images and uploads them to a remote server or evaluate them locally. Deep learning models are used to calculate a feature vector for each image. It returns an enhanced data table with additional columns (image descriptors).



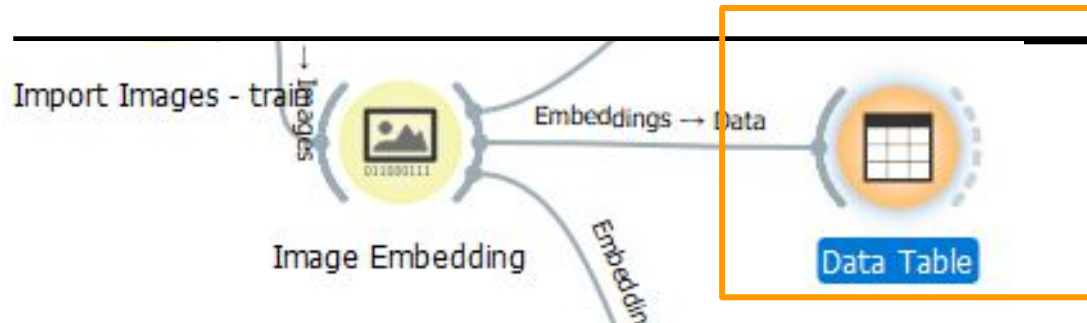
Inception v3 (GoogLeNet)



Inception-v3 Architecture (Batch Norm and ReLU are used after Conv)

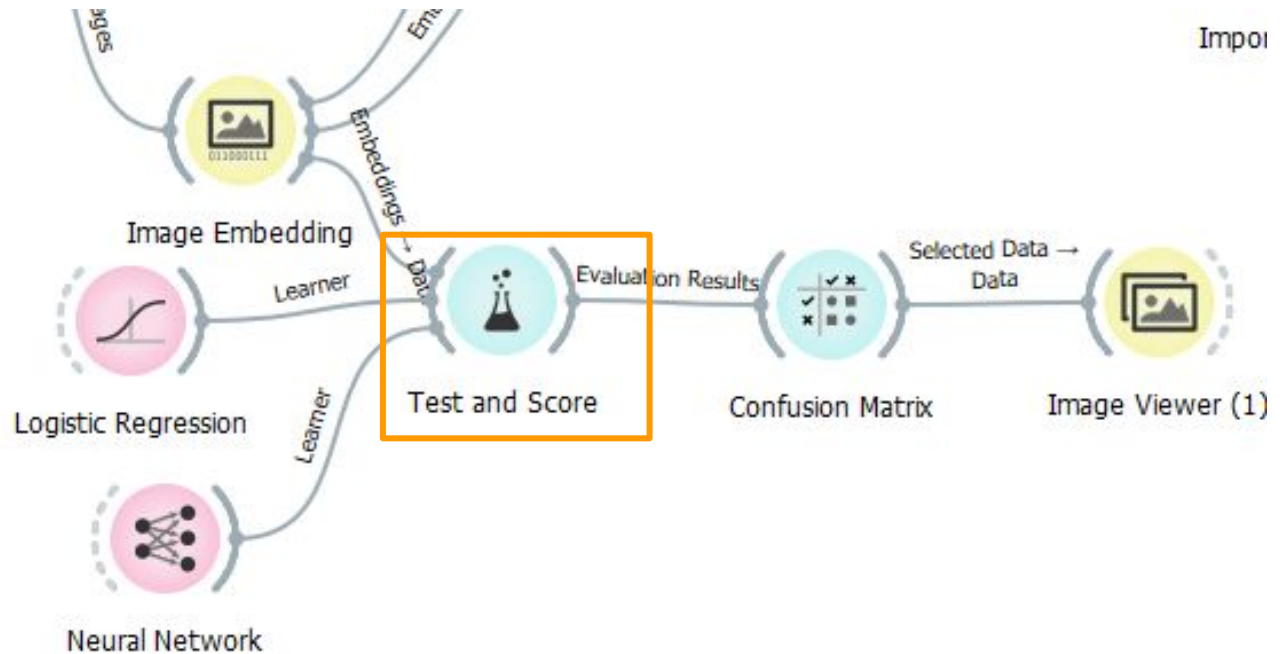
[inception v3] C. Szegedy, W. Liu, Y. Jia, P. Sermanet, S. Reed, D. Anguelov, D. Erhan, V. Vanhoucke, and A. Rabinovich. Going deeper with convolutions. In Proceedings of the IEEE Conference on Computer Vision and Pattern Recognition, pages 1–9, 2015

<https://cloud.google.com/tpu/docs/inception-v3-advanced>



Data Table

hidden origin	category	image name	image	size	width	height	n0 True	n1 True	n2 True			
1	cats	cat.1	cats\cat.1.jpg	16880	300	280	0.475694	0.432614	0.69527			
2	cats	cat.10	cats\cat.10.jpg	34315	489	499	0.31333	0.241181	0.432035			
3	cats	cat.100	cats\cat.100.jpg	28277	402	499	0.629442	0.265338	0.541589			
4	cats	cat.1000	cats\cat.1000.jpg	n2043 True	n2044 True	n2045 True	n2046 True	n2047 True	149	0.663272	0.333273	1.13996
5	cats	cat.1001	cats\cat.1001.jpg	0.0000036	0.171371	0.390427	0.000073291	0.399337	499	0.345729	0.111364	0.275736
6	cats	cat.1002	cats\cat.1002.jpg	0.371487	0.147471	0.492729	0.140827	0.976613	407	0.576959	0.631882	0.716025
7	cats	cat.1003	cats\cat.1003.jpg	0.147725	0.00344998	0.0549949	0.0806171	0.0727724	269	0.366972	0.44163	0.867934
8	cats	cat.1004	cats\cat.1004.jpg	0.427751	0.181229	0.813569	0.211402	0.264475	375	0.432891	0.0436033	0.38925
			cats\cat.1004.jpg	0.640509	0	0.239425	0.195675	0.174547				
			cats\cat.1004.jpg	0.243388	0.356838	0.0456228	0.112791	0.293549				
			cats\cat.1004.jpg	0.072368	0.0131649	0.373584	0.0435946	0.241524				
			cats\cat.1004.jpg	0.0247444	0.0243987	0.082787	0.0150555	0.595918				
			cats\cat.1004.jpg	0.329187	0.0473317	0.243795	0.254472	0.0699269				



Test and Score

Tests learning algorithms on data.

Inputs

- Data: input dataset
- Test Data: separate data for testing
- Learner: learning algorithm(s)

Outputs

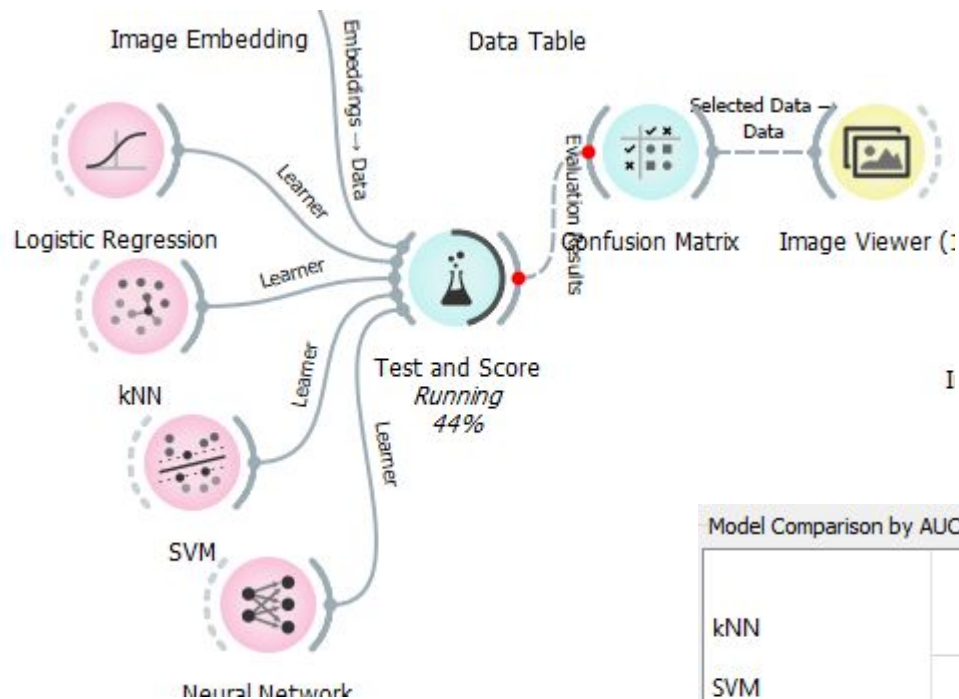
- Evaluation Results: results of testing classification algorithms

The widget tests learning algorithms.
Different sampling schemes are available,
including using separate test data.

The image shows the 'Test and Score' widget interface and its workflow diagram. The interface is divided into a configuration panel on the left and a results table on the right. The configuration panel includes options for 'Sampling' (Cross validation, Random sampling), 'Number of folds' (10), 'Stratified' (checked), 'Repeat train/test' (10), 'Training set size' (66%), and 'Target Class' (Average over classes). The results table shows performance metrics for Neural Network and Logistic Regression models. The workflow diagram below shows the 'Test and Score' widget (highlighted with an orange box) receiving data from 'Image Embedding' and 'Neural Network' learners, and outputting 'Evaluation Results' to a 'Confusion Matrix' widget, which then outputs 'Selected Data' to an 'Image Viewer (1)' widget.

Model	AUC	CA	F1	Precision	Recall
Neural Network	1.000	0.994	0.994	0.994	0.994
Logistic Regression	1.000	0.994	0.994	0.994	0.994

	Neural N...	Logistic ...
Neural Network		0.569
Logistic Regression	0.431	



Evaluation Results					
Model	AUC	CA	F1	Precision	Recall
kNN	0.997	0.993	0.993	0.993	0.993
SVM	1.000	0.994	0.994	0.994	0.994
Neural Network	1.000	0.994	0.994	0.994	0.994
Logistic Regression	1.000	0.994	0.994	0.994	0.994

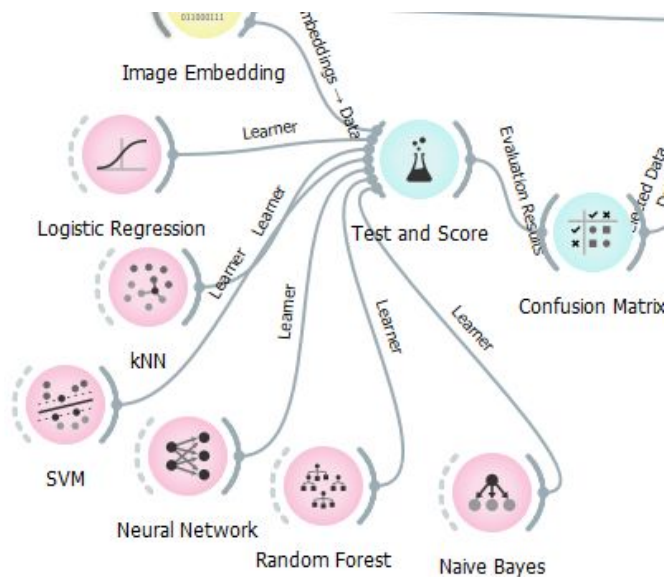
Model	AUC	CA	F1	Precision	Recall
kNN	0.997	0.993	0.993	0.993	0.993
SVM	1.000	0.994	0.994	0.994	0.994
Random Forest	0.999	0.991	0.991	0.991	0.991
Neural Network	1.000	0.994	0.994	0.994	0.994
Naive Bayes	0.989	0.984	0.984	0.985	0.984
Logistic Regression	1.000	0.994	0.994	0.994	0.994

Model Comparison by AUC

kNN
SVM
Neural Network
Logistic Regression

Model Comparison by AUC

	kNN	SVM	Rando...	Neural ...	Naive B...	Logistic...
kNN		0.006	0.025	0.006	1.000	0.007
SVM	0.994		0.938	0.618	1.000	0.641
Random Forest	0.975	0.062		0.068	1.000	0.050
Neural Network	0.994	0.382	0.932		1.000	0.569
Naive Bayes	0.000	0.000	0.000	0.000		0.000
Logistic Regression	0.993	0.359	0.950	0.431	1.000	



Model	AUC	CA	F1	Precision	Recall
kNN	0.997	0.993	0.993	0.993	0.993
SVM	1.000	0.994	0.994	0.994	0.994
Random Forest	0.999	0.991	0.991	0.991	0.991
Neural Network	1.000	0.994	0.994	0.994	0.994
Naive Bayes	0.989	0.984	0.984	0.985	0.984
Logistic Regression	1.000	0.994	0.994	0.994	0.994

Model Comparison by AUC

	kNN	SVM	Rando...	Neural ...	Naive B...	Logistic...
kNN		0.006	0.025	0.006	1.000	0.007
SVM	0.994		0.938	0.618	1.000	0.641
Random Forest	0.975	0.062		0.068	1.000	0.050
Neural Network	0.994	0.382	0.932		1.000	0.569
Naive Bayes	0.000	0.000	0.000	0.000		0.000
Logistic Regression	0.993	0.359	0.950	0.431	1.000	

- [Area under ROC](#) is the area under the receiver-operating curve.
- [Classification accuracy](#) is the proportion of correctly classified examples.
- [F-1](#) is a weighted harmonic mean of precision and recall (see below).
- [Precision](#) is the proportion of true positives among instances classified as positive, e.g. the proportion of *Iris virginica* correctly identified as Iris virginica.
- [Recall](#) is the proportion of true positives among all positive instances in the data, e.g. the number of sick among all diagnosed as sick.

Test and Score

Sampling

Cross validation

Number of folds: 10

Stratified

Cross validation by feature

Random sampling

Repeat train/test: 10

Training set size: 66 %

Stratified

Leave one out

Test on train data

Test on test data

Target Class

(Average over classes)

Click on the table header to select shown columns

Model	AUC	CA	F1	Precision	Recall
Neural Network	1.000	0.994	0.994	0.994	0.994
Logistic Regression	1.000	0.994	0.994	0.994	0.994

Model Comparison by AUC

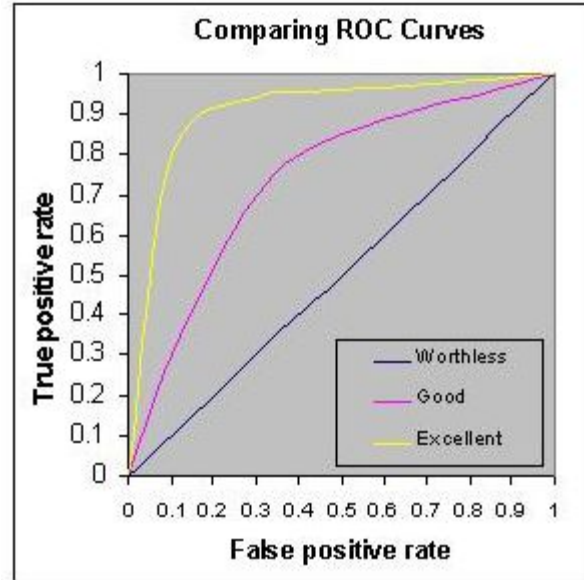
	Neural N...	Logistic ...
Neural Network		0.569
Logistic Regression	0.431	

- **Area under ROC** is the area under the receiver-operating curve.

- .90-1 = excellent (A)
- .80-.90 = good (B)
- .70-.80 = fair (C)
- .60-.70 = poor (D)
- .50-.60 = fail (F)

Click on the table header to see columns

Model	AUC
Neural Network	1.000
Logistic Regression	1.000

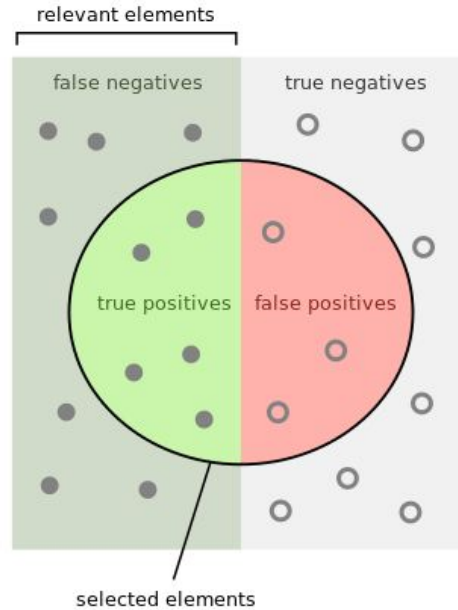


-
- **Classification accuracy** is the proportion of correctly classified examples.

select show	
CA	
0.994	0
0.994	0

accuracy is closeness of the measurements to a specific value, while **precision** is the closeness of the measurements to each other.

- Precision** is the proportion of true positives among instances classified as positive, e.g. the proportion of *Iris virginica* correctly identified as *Iris virginica*.



- Recall** is the proportion of true positives among all positive instances in the data, e.g. the number of sick among all diagnosed as sick.

How many selected items are relevant?

$$\text{Precision} = \frac{\text{true positives}}{\text{true positives} + \text{false positives}}$$

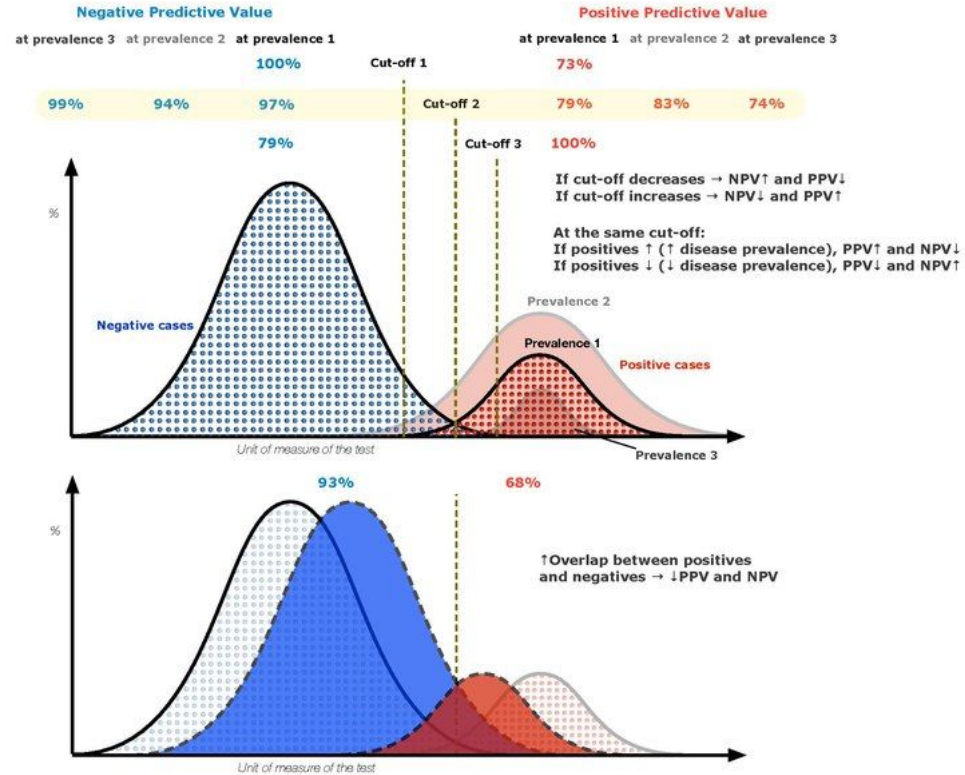
How many relevant items are selected?

$$\text{Recall} = \frac{\text{true positives}}{\text{true positives} + \text{false negatives}}$$

	Ok, got it	X
	Precision	Recall
4	0.994	0.994
4	0.994	0.994

Both precision and recall are therefore based on relevance.

- Precision** is the proportion of true positives among instances classified as positive, e.g. the proportion of *Iris virginica* correctly identified as *Iris virginica*.



precision (also called positive predictive value) is the fraction of relevant instances among the retrieved instances

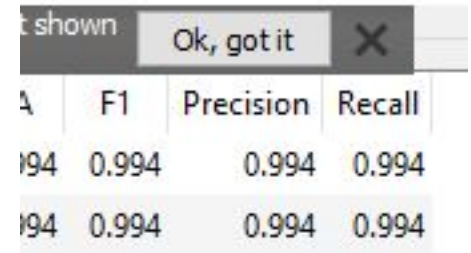
F-score or **F-measure** is a measure of a test's accuracy.

- **F-1** is a weighted harmonic mean of precision and recall .

The traditional F-measure or balanced F-score (**F₁ score**) is the **harmonic mean** of precision and recall:

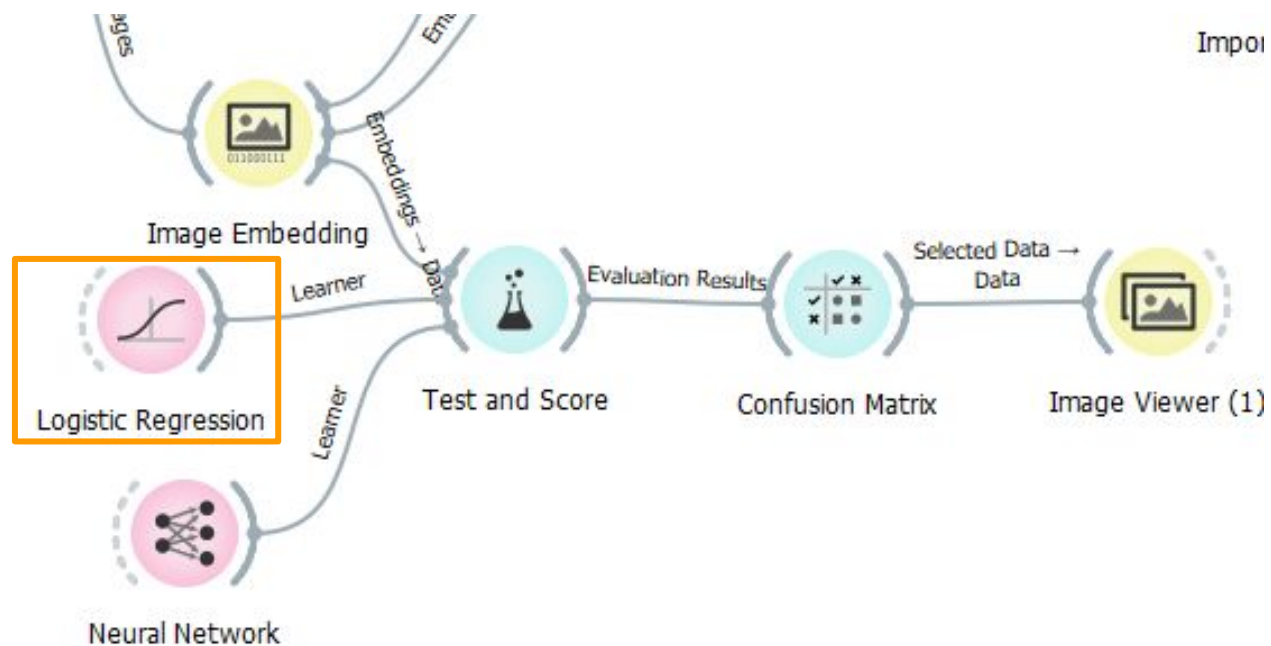
$$F_1 = \frac{2}{\text{recall}^{-1} + \text{precision}^{-1}} = 2 \cdot \frac{\text{precision} \cdot \text{recall}}{\text{precision} + \text{recall}} = \frac{2 \cdot \text{tp}}{2 \cdot \text{tp} + \text{fp} + \text{fn}}$$

The highest possible value of an F-score is 1.0, indicating perfect precision and recall, and the lowest possible value is 0, if either the precision or the recall is zero. The F₁ score is also known as the Dice similarity coefficient (DSC)



The image shows a screenshot of a table with a modal dialog box overlaid on top. The dialog box has a title bar that says "shown" and contains the text "Ok, got it" and a close button (X). The table below has four columns: "A", "F1", "Precision", and "Recall". The first row of the table has the value "0.994" in each of these four columns. The second row of the table has the value "0.994" in each of these four columns.

A	F1	Precision	Recall
0.994	0.994	0.994	0.994
0.994	0.994	0.994	0.994



Logistic Regression

```
def computeLogisticRegressionModel(XTrain, yTrain):
    from sklearn.linear_model import LogisticRegression
    classifier = LogisticRegression(solver='lbfgs')
    classifier.fit(XTrain[0], yTrain)
    return classifier

def computeLogisticRegressionExample(filename):
    X, y, _ = pre.loadDataset(filename, ".")
    X = pre.fillMissingData(X, 2, 3)
    X = pre.computeCategorization(X)
    X = pre.computeCategorization(X)
```

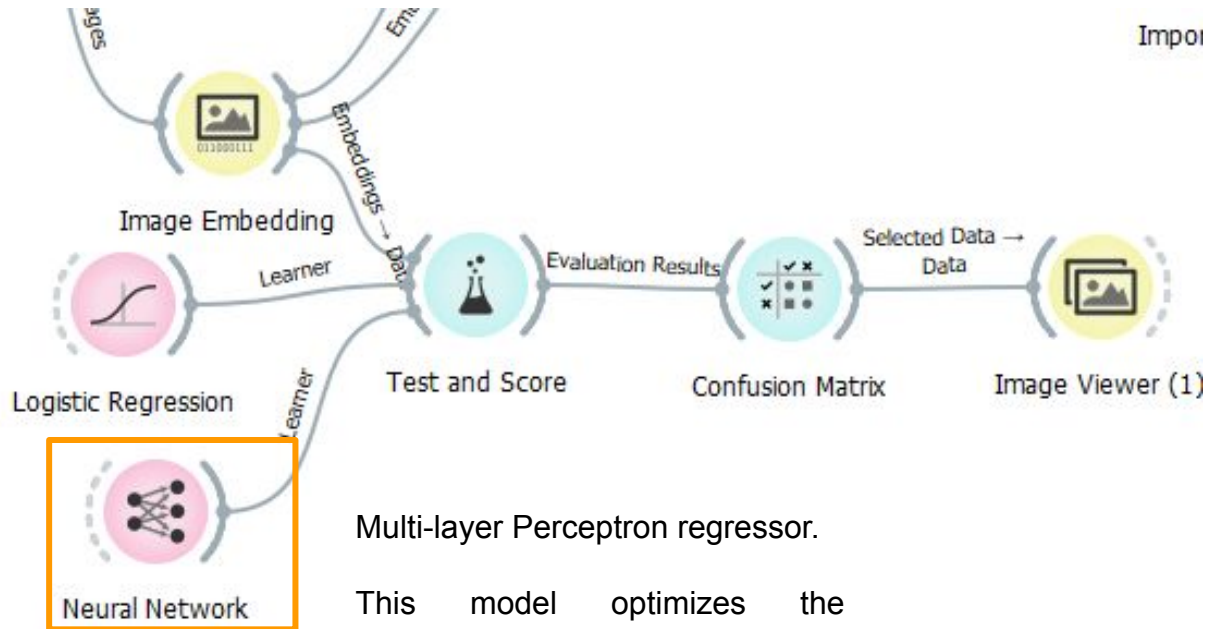
<https://www.youtube.com/watch?v=DMDY0Gar7Fw>

regressão logística (ou, classificador de máxima entropia) é uma técnica estatística que tem como objetivo produzir, a partir de um conjunto de observações, um modelo que permita a predição de valores tomados por uma variável categórica, frequentemente binária, a partir de uma série de variáveis explicativas contínuas e/ou binária

Seja $p(x)$ a probabilidade de êxito quando o valor da variável preditiva é x .

Então,

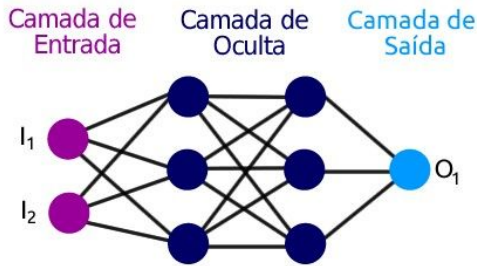
$$p(x) = \frac{1}{1 + e^{-(B_0 + B_1 x)}} = \frac{e^{B_0 + B_1 x}}{1 + e^{B_0 + B_1 x}}.$$



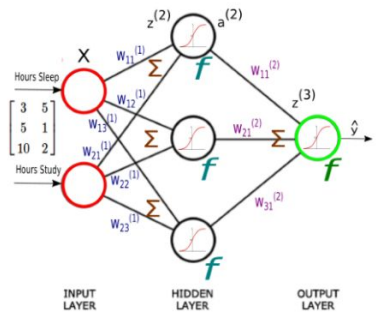
Multi-layer Perceptron regressor.

This model optimizes the squared-loss using LBFGS or stochastic gradient descent.

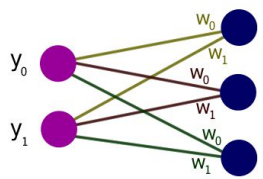
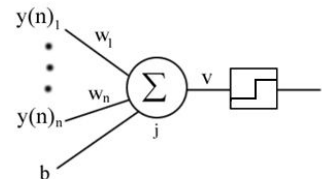
Neural Network



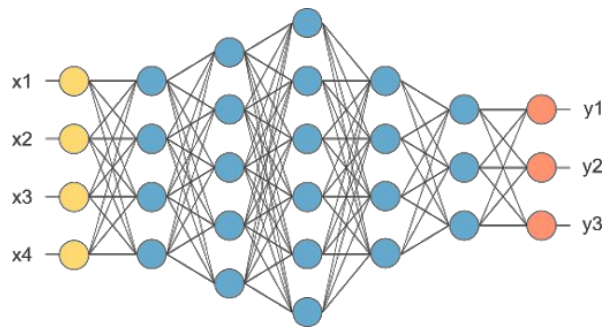
$I = [I_1, I_2] = \text{Vetor de Entrada}$
 $O = [O_1] = \text{Vetor de Saída}$



computador aprende a realizar uma tarefa analisando exemplos de treinamento, que normalmente são indicados manualmente com antecedência.



$$v_j = \sum_{i=0}^m w_i y_i + b$$

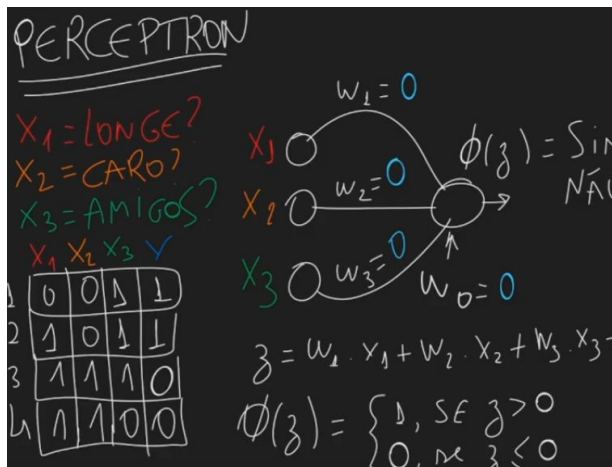


- Processamento de linguagem natural;
- Reconhecimento de fala e imagens;
- Previsão de valores.

Neural Network

$$(XW_1)w = y$$

$$\begin{bmatrix} 1 & x_{11} & \dots & x_{1d} \\ 1 & x_{21} & \dots & x_{2d} \\ \vdots & \vdots & \vdots & \vdots \\ 1 & x_{n1} & \dots & x_{nd} \end{bmatrix} \times \begin{bmatrix} w_{01} & w_{01} & \dots & w_{0m} \\ w_{11} & w_{11} & \dots & w_{1m} \\ \vdots & \vdots & \vdots & \vdots \\ w_{d1} & w_{d1} & \dots & w_{dm} \end{bmatrix} \times \begin{bmatrix} w_{01} \\ w_{11} \\ \vdots \\ w_{d1} \end{bmatrix} = \begin{bmatrix} y_0 \\ y_1 \\ \vdots \\ y_n \end{bmatrix}$$



<https://www.youtube.com/watch?v=tYXGzQs31Og>

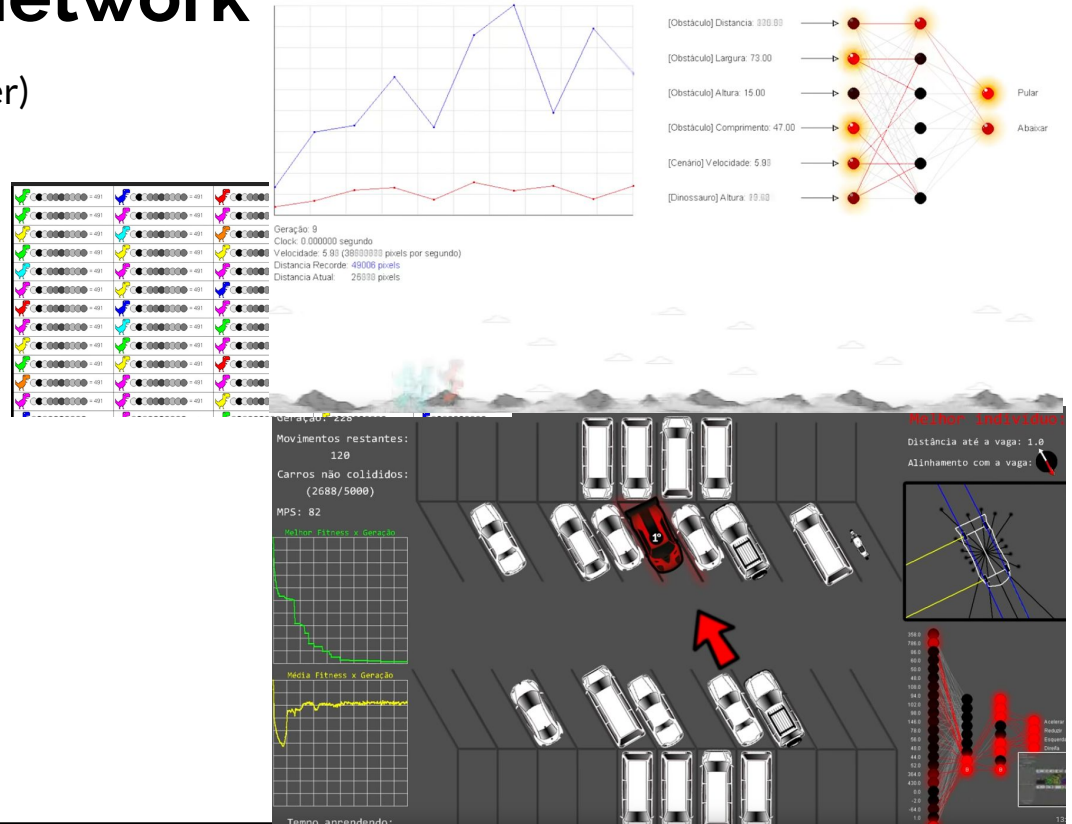
Neural Network

Rede Neural Artificial (Perceptron Multilayer)

Jogando!

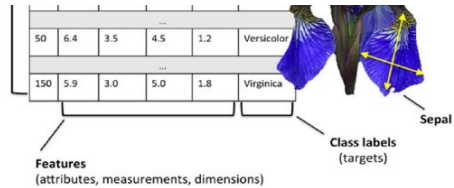
Dino:

Estacionando:



kNN - K-nearest neighbors

[knn colab](#)



To perform some extra checks to the Dataframe, the Pandas library itself may work on any data it is essential to do a preliminary analysis of the data: the type missing values. Then, take the first conclusions and correct any issue* in app

*this is not the case of this well-known data but the checks are to be performed at

```
[ ] import pandas as pd

[ ] pd.value_counts(iris.species)

setosa      50
versicolor  50
virginica   50
Name: species, dtype: int64
```

The dataset is made of 150 flowers (150 rows), 50 of each specie.

MACHINE LEARNING - Predictive Analysis

After performing the exploratory analysis, what is basically, evaluate and reaching some conclusions looking to the charts of distribution and scatter of the data.

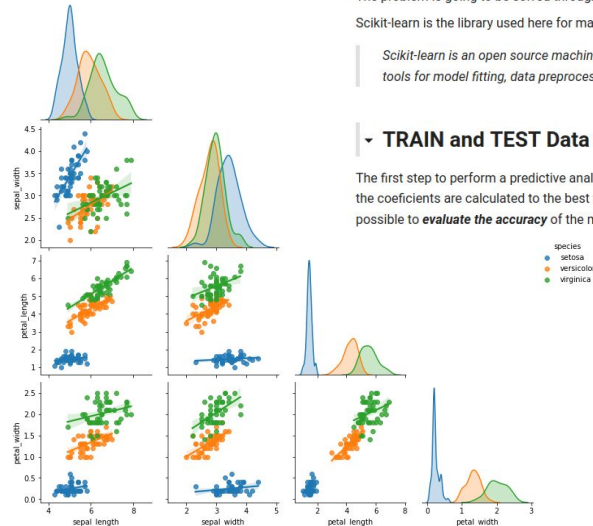
Now, the predictive analysis is going to be performed not by a person itself but with the aid of a computer, mathematical algorithms, machine learning.

The problem is going to be solved through the KNN algorithm. The best practices are going to be performed and explained. Scikit-learn is the library used here for machine learning in Python. From it several modules are imported as needed.

Scikit-learn is an open source machine learning library that supports supervised and unsupervised learning. It also provides various tools for model fitting, data preprocessing, model selection and evaluation, and many other utilities.

TRAIN and TEST Data

The first step to perform a predictive analysis through a Machine Learning algorithm is to separate the data in two parts. One to be trained, so the coefficients are calculated to the best fit to this data, according to the algorithm chosen. The second part is reserved to be tested, so it is possible to evaluate the accuracy of the method with a different data, so that you can extrapolate to any data.



[Atividade com knn no colab](#)

SVM - Support Vector Machine

SVM é um algoritmo de aprendizado de máquina **supervisionado** que pode ser usado para desafios de classificação ou regressão.

Para treinamento e classificação de um dataset.

SVM é uma fronteira que melhor segrega as duas classes (hiperplano / linha).

SVM - [scikit](#) - python

[colab - svm](#)

CLASSIFICAÇÃO com Máquinas de Vetores de Suporte (SVM) | Machine Learning #05

Classes linearmente separáveis podem ser entendidas, grosso modo, como classes que podem ser facilmente separadas com uma linha reta.

```
In [1]: from sklearn import datasets
import pandas as pd
import seaborn as sns

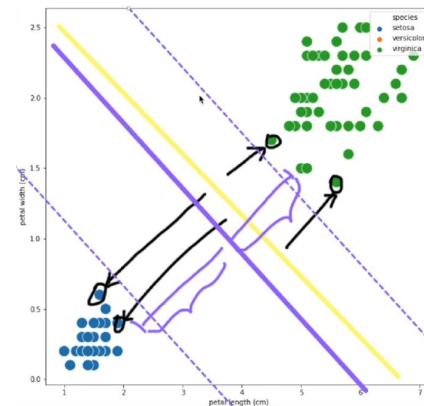
In [2]: iris = datasets.load_iris()
iris_df = pd.DataFrame(data=iris.data, columns=iris.feature_names)
iris_df['label'] = iris.target
iris_df['species'] = pd.Categorical.from_codes(iris.target, iris.target_names)

In [3]: iris_df.head()
```

```
Out[3]:
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	label	species
0	5.1	3.5	1.4	0.2	0	setosa
1	4.9	3.0	1.4	0.2	0	setosa
2	4.7	3.2	1.3	0.2	0	setosa
3	4.6	3.1	1.5	0.2	0	setosa
4	5.0	3.6	1.4	0.2	0	setosa

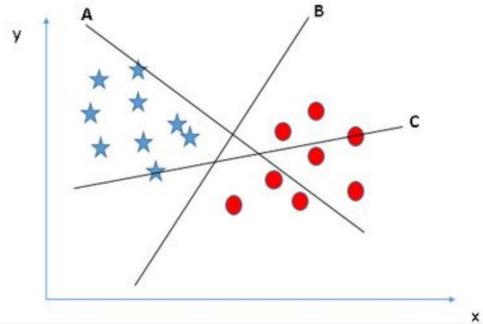
```
In [ ]:
```



SVM - scikit - python

[SVM no site do scikit-learn](#)

supervised learning methods used for classification, regression and outliers detection.



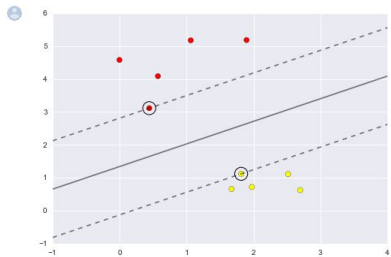
SVM - scikit - python

[SVM no colab](#)

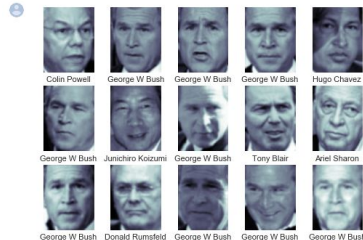
$$\min_w \lambda \|w\|^2 + \sum_{i=1}^n (1 - y_i \langle x_i, w \rangle)_+$$

[Python Data Science handbook](#)

```
from ipywidgets import interact, fixed
interact(plot_svm, N=[10, 200], ax=fixed(None));
```



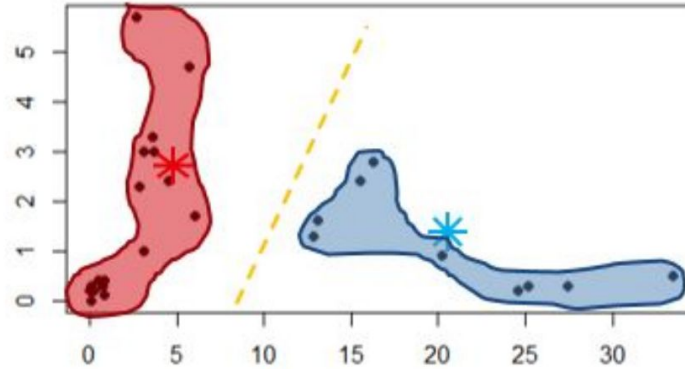
```
fig, ax = plt.subplots(3, 5)
for i, axi in enumerate(ax.flat):
    axi.imshow(faces.images[i], cmap='bone')
    axi.set(xticks=[], yticks=[],
           xlabel=faces.target_names[faces.target[i]])
```



```
from sklearn.metrics import confusion_matrix
res = confusion_matrix(y_test, predictions)
print("Confusion Matrix")
print(res)
print(f"Test Set: {len(X_test)}")
print(f"Accuracy = {percentage*100} %")
```

```
Confusion Matrix
[[63  8]
 [ 1 99]]
Test Set: 171
Accuracy = 94.73684210526315 %
```

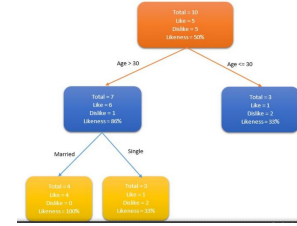
K-Means



Naïve Bayes

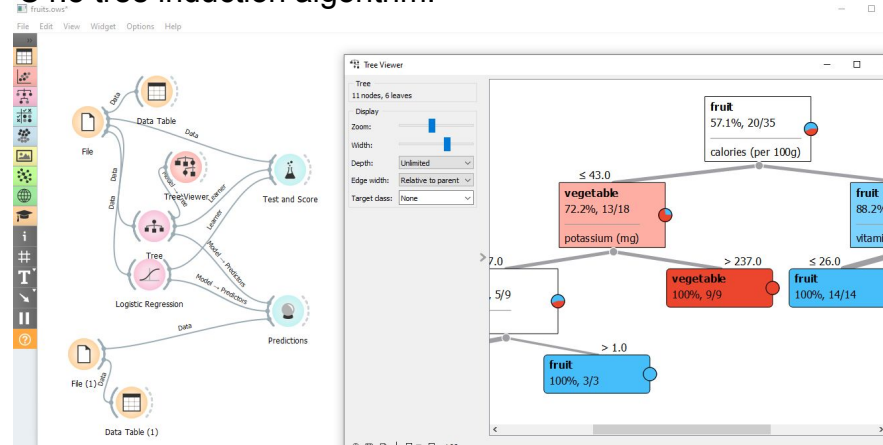
classificadores probabilísticos simples com base na aplicação Bayes 'teorema com forte independência entre as características. A imagem em destaque é a equação – em que $P(A|B)$ é a probabilidade posterior, $P(B|A)$ é a probabilidade, $P(A)$ é a probabilidade prévia e $P(B)$ é preditor de probabilidade prévia.

Árvore de decisão



Uma árvore de decisão é uma ferramenta de apoio que utiliza um gráfico ou modelo de decisões e suas possíveis consequências.

C4.5 tree induction algorithm.



Supervisionado

O aprendizado supervisionado é baseado na regressão básica e classificação. O humano fornece um banco de dados e ensina a máquina a reconhecer o que é uma bicicleta, por exemplo, entre padrões e semelhanças. A cor e tamanho pode variar, mas a máquina aprende que uma bicicleta possui pedais, duas rodas, guidão e outros elementos-chave.

não supervisionado

- Associação;
- Clusterização.

A associação: permite o descobrimento de regras e correlações, identificando conjuntos de itens que frequentemente ocorrem juntos. Os varejistas costumam usar esta análise em carrinhos de compras, para descobrir itens frequentemente comprados em conjunto, desenvolvendo assim estratégias mais eficazes de marketing e merchandising.

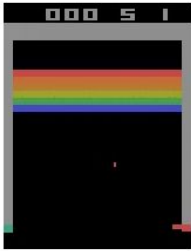
Na clusterização (ou agrupamento): o conjunto todo em análise sofre segmentações em vários grupos, com base nas semelhanças encontradas. É uma técnica que permite dividir automaticamente um conjunto de dados em grupos de acordo com medidas de similaridade ou de distância.

não supervisionado

a máquina começa a analisar, sozinha, os dados e a identificar os padrões — aprendendo a separar o que é uma lata de uma garrafa, por exemplo.

+Demorado

Performance inicial do agente:



Após 15 minutos de treinamento:



Após 30 minutos de treinamento:

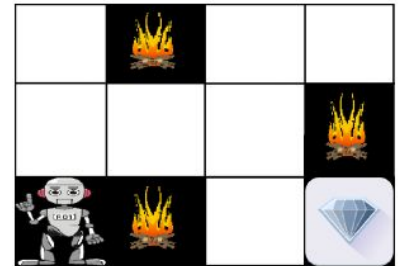


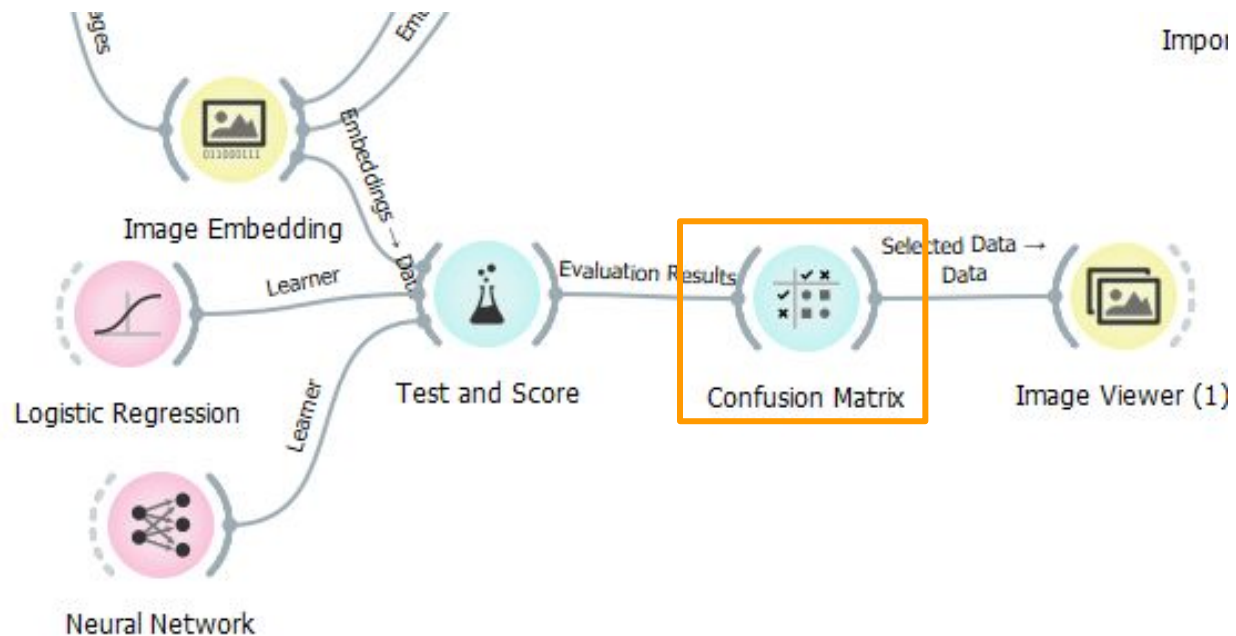
Reforço

No Aprendizado por Reforço o modelo aprende executando ações e avaliando recompensas.

O agente realiza uma ação num dado ambiente, alterando seu estado inicial, o que gera uma recompensa ao agente. De forma cíclica, o agente avalia esta recompensa (que pode ser positiva ou negativa) e age novamente no ambiente, gerando o aprendizado.

- Q-Learning;
- Aproximação por função com atualização por gradiente;
- Multi-Armed Bandits;
- Contextual Bandits;
- k-Armed Bandits





Confusion Matrix

Em análise preditiva, a **matriz de confusão** (às vezes também chamada de '*matriz de erro*' ou '*tabela de confusão*') é uma tabela com duas linhas e duas colunas que relata o número de *falsos positivos*, *falsos negativos*, *verdadeiros positivos* e *verdadeiros negativos*. Isso permite uma análise mais detalhada do que a mera proporção de classificações corretas (precisão). A **precisão** produzirá resultados enganosos se o conjunto de dados estiver desequilibrado; isto é, quando o número de observações em diferentes classes variam muito.

Por exemplo, se houver 95 gatos e apenas 5 cachorros nos dados, um determinado classificador pode classificar todas as observações como gatos. A precisão geral seria de 95%, mas com mais detalhes o classificador teria uma taxa de reconhecimento de 100% para a classe de gatos, mas uma taxa de reconhecimento de 0% para a classe de cães.

O **F1 Score (ou f-measure)** é ainda mais confiável em tais casos, e aqui renderia mais de 97,4%, enquanto remove o viés e produz 0 como a probabilidade de uma decisão informada para qualquer forma de suposição (supondo gato).

Confusion Matrix

```
import numpy as np

# 1 para grávida, 0 para não grávida
valores_reais = [1, 0, 1, 0, 0, 0, 1, 0, 1, 0]
valores_preditos = [1, 0, 0, 1, 0, 0, 1, 1, 1, 0]

def get_confusion_matrix(reais, preditos, labels):
    ..
    ..
    ...

get_confusion_matrix(reais=valores_reais, preditos=valores_preditos, labels=[1,0])
# array([[3, 1], [2, 4]])
```

Confusion Matrix

Learners
Logistic Regression

Clicking on cells or in headers outputs the corresponding data instances Ok, got it Show: Number of instances

		Predicted		Σ
		cats	dogs	
Actual	cats	3972	28	4000
	dogs	19	3986	4005
Σ		3991	4014	8005

Predictions Probabilities
 Apply Automatically

1x8005 3972 | 8005

Select Correct Select Misclassified Clear Selection

Import

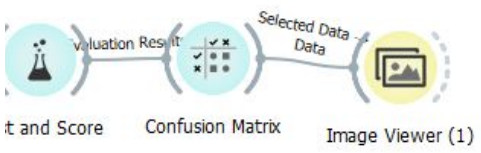
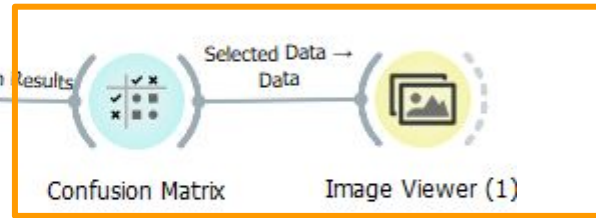



Image Viewer (1)


Image Filename Attribute
image

Title Attribute
category

Image Size



cats



cats

Confusion Matrix

Clicking on cells or in headers outputs the corresponding data instances Ok, got it

Learners

- Logistic Regression
- Neural Network

		Predicted		Σ
		cats	dogs	
Actual	cats	3971	29	4000
	dogs	20	3985	4005
Σ		3991	4014	8005

Predictions Probabilities
 Apply Automatically

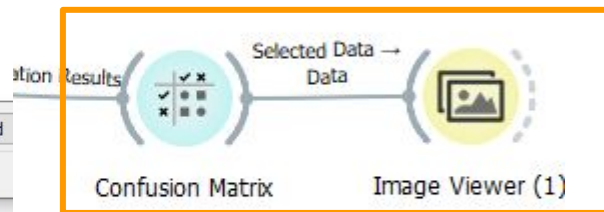
Select Correct Select Misclassified

? | 2x8005 | 29 | 8005

category

Image Size

Impo



Learners

- Logistic Regression
- Neural Network
- kNN
- SVM

Predictions Probabilities

Clicking on cells or in headers outputs the corresponding data instances Ok, got it

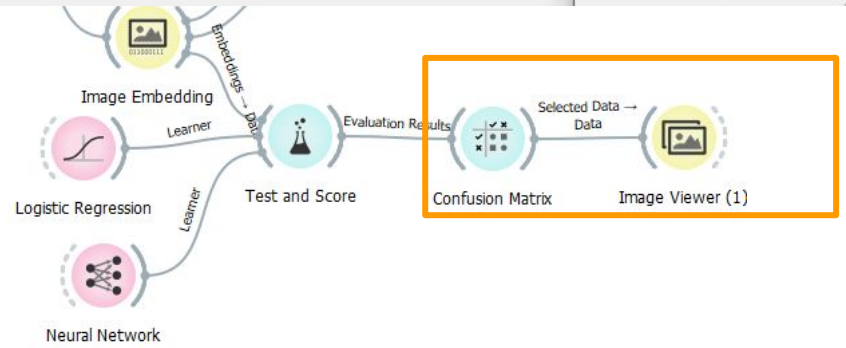
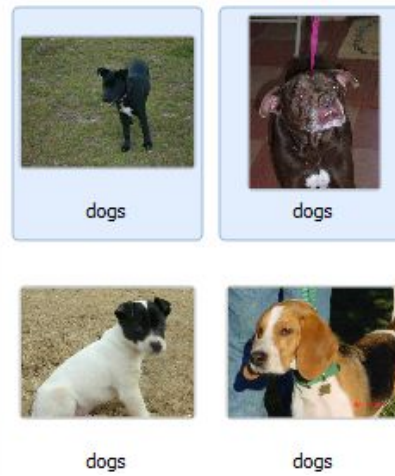
Show: Number of instances

		Predicted		Σ
		cats	dogs	
Actual	cats	3957	43	4000
	dogs	14	3991	4005
Σ		3971	4034	8005

Image Filename Attribute
S image

Title Attribute
C category

Image Size



Clicking on cells or in headers outputs the corresponding data instances

Ok, got it

Show: Number of instances

Learners

- Logistic Regression
- Neural Network
- kNN
- SVM

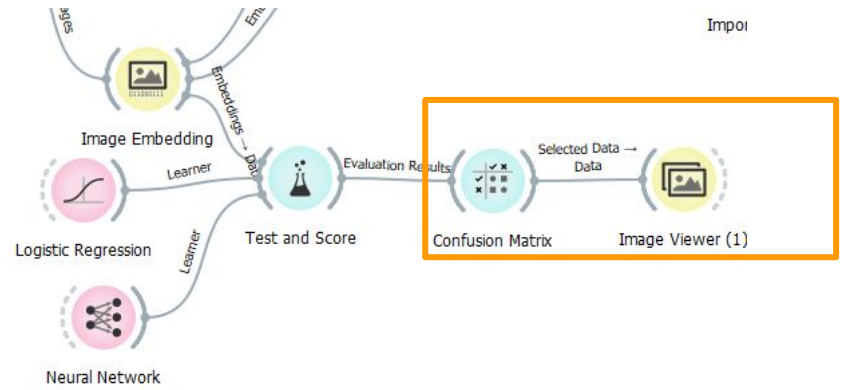
		Predicted		
		cats	dogs	Σ
Actual	cats	3969	31	4000
	dogs	18	3987	4005
Σ		3987	4018	8005

Image Filename Attribute: image

Title Attribute: category

Image Size: [Slider]

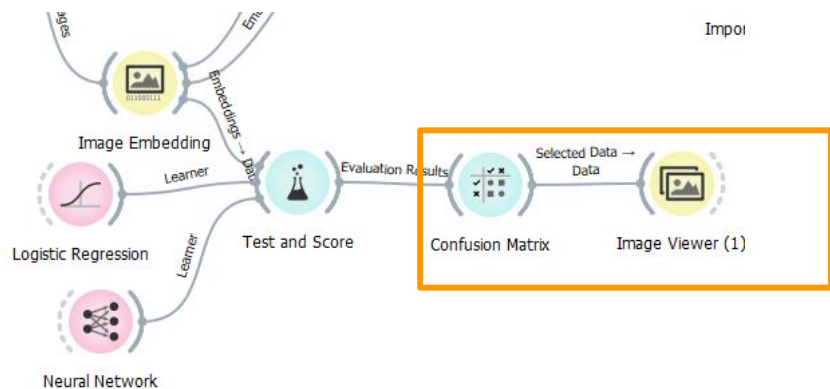
cats cats



corresponding data instances

Logistic Regression
 Neural Network
 kNN
 SVM
Random Forest
 Naive Bayes

		Predicted		
		cats	dogs	Σ
Actual	cats	3951	49	4000
	dogs	22	3983	4005
Σ		3973	4032	8005

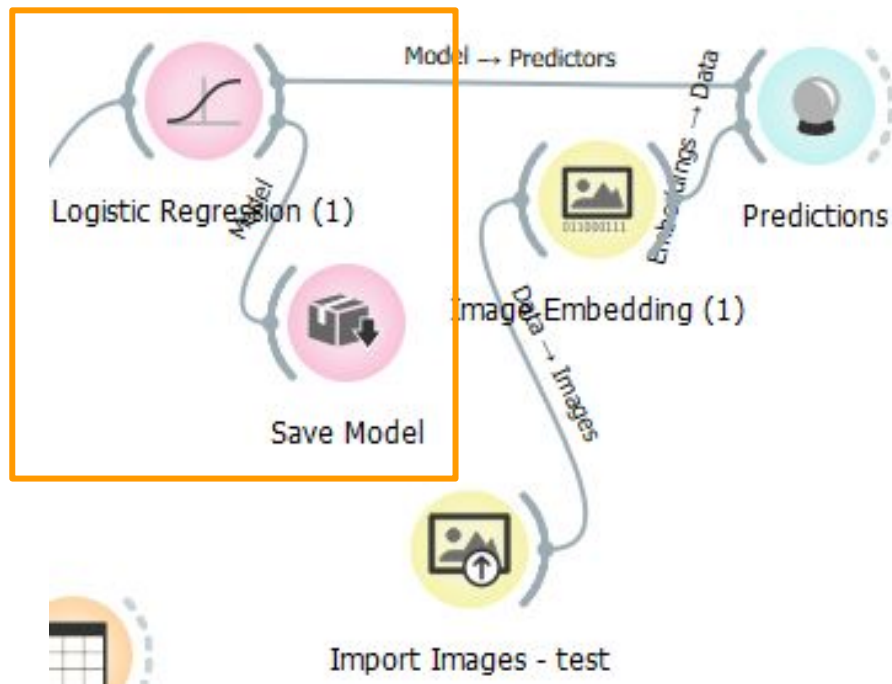


Clicking on cells or in headers outputs the corresponding data instances

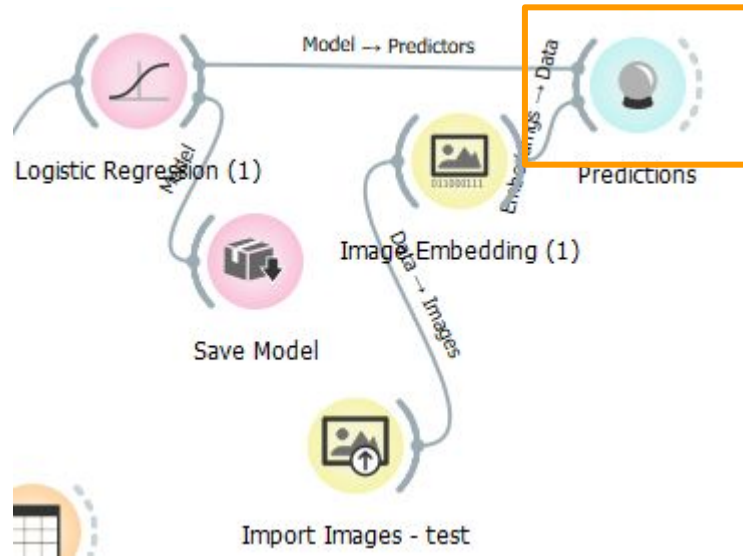
Logistic Regression
 Neural Network
 kNN
 SVM
 Random Forest
Naive Bayes

		Predicted		
		cats	dogs	Σ
Actual	cats	3880	120	4000
	dogs	7	3998	4005
Σ		3887	4118	8005

Model



Predictor



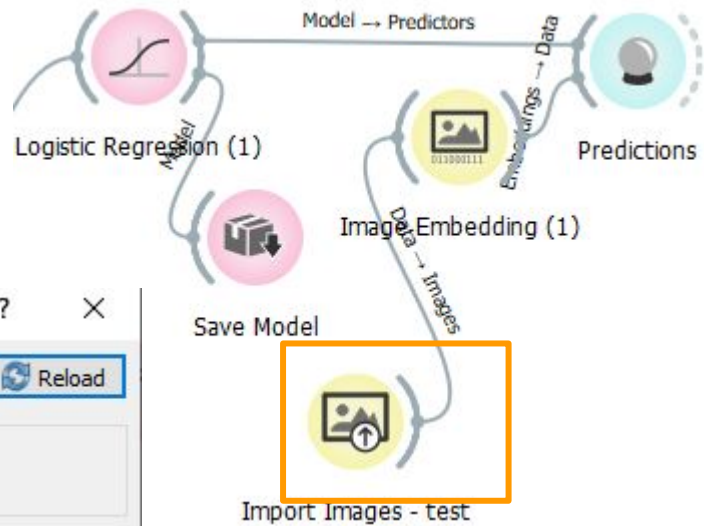
Import Images - test

misturados

Info

40 images

40

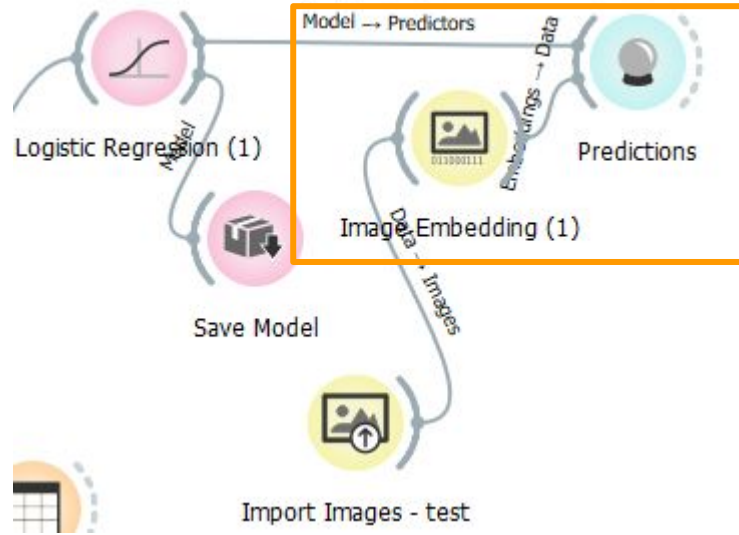


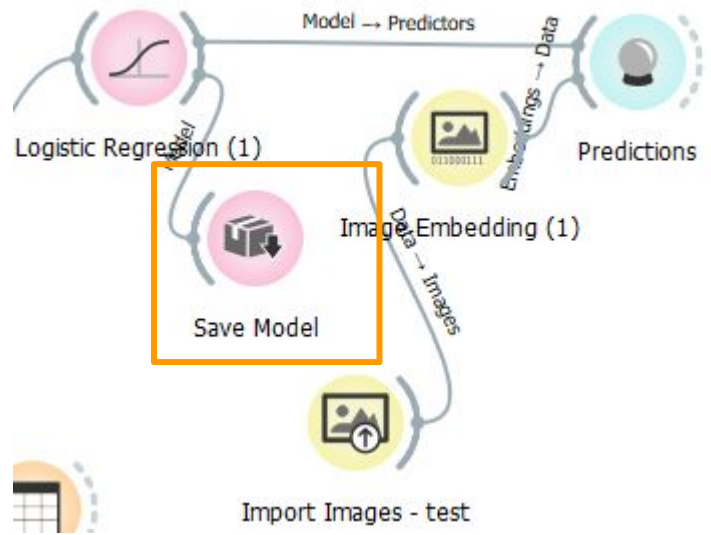
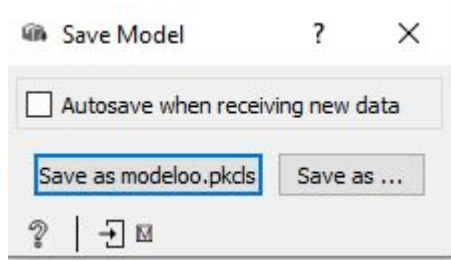
Predictions

Show probabilities for
cats
dogs

	Logistic Regression	image name	image	size
22	cats	cat.435	cat.435.jpg	15110
23	dogs	dog.4072	dog.4072.jpg	22541
24	dogs	dog.4073	dog.4073.jpg	16045
25	dogs	dog.4074	dog.4074.jpg	25053
26	dogs	dog.4075	dog.4075.jpg	30518
27	dogs	dog.4076	dog.4076.jpg	20626
28	dogs	dog.4077	dog.4077.jpg	24940
29	dogs	dog.4080	dog.4080.jpg	31661
30	dogs	dog.4081	dog.4081.jpg	20310
31	dogs	dog.4082	dog.4082.jpg	29628
32	dogs	dog.4083	dog.4083.jpg	13272
33	dogs	dog.4084	dog.4084.jpg	14323
34	dogs	dog.4085	dog.4085.jpg	25691
35	dogs	dog.4088	dog.4088.jpg	37892
36	dogs	dog.4089	dog.4089.jpg	14107
37	dogs	dog.4090	dog.4090.jpg	6497
38	dogs	dog.4091	dog.4091.jpg	7081
39	dogs	dog.4092	dog.4092.jpg	18295
40	dogs	dog.4093	dog.4093.jpg	20787

Restore Original Order

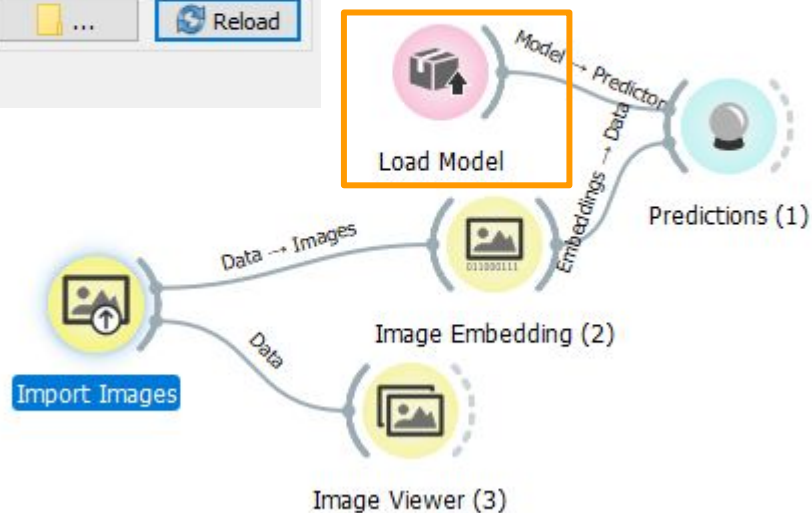
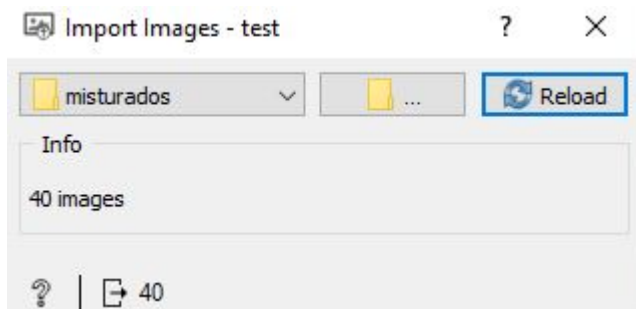




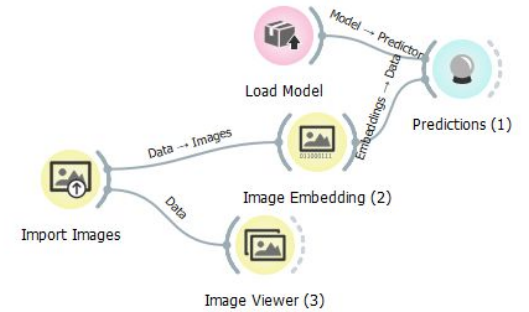
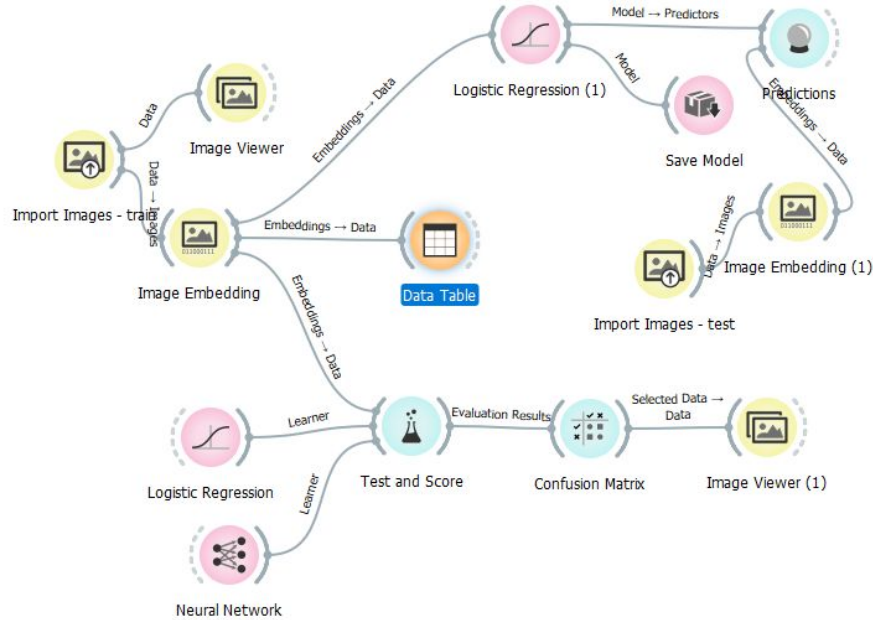
After trained....

use

load model and use



complete view



pause orange

coding an example - cat or dog

[Colab + python + keras + matplotlib + numpy](#) = dogs vc cats

2,000 JPG pictures of cats and dogs

```
base_dir = '/tmp/cats_and_dogs_filtered'
train_dir = os.path.join(base_dir, 'train')
validation_dir = os.path.join(base_dir, 'validation')

# Directory with our training cat pictures
train_cats_dir = os.path.join(train_dir, 'cats')

# Directory with our training dog pictures
train_dogs_dir = os.path.join(train_dir, 'dogs')

# Directory with our validation cat pictures
validation_cats_dir = os.path.join(validation_dir, 'cats')

# Directory with our validation dog pictures
validation_dogs_dir = os.path.join(validation_dir, 'dogs')
```

```
[7] train_cat_fnames = os.listdir(train_c
print(train_cat_fnames[:10])
```

```
train_dog_fnames = os.listdir(train_c
train_dog_fnames.sort()
print(train_dog_fnames[:10])
```

```
['cat.427.jpg', 'cat.256.jpg', 'cat.4
['dog.0.jpg', 'dog.1.jpg', 'dog.10.jp
```

Let's find out the total number of cat and dog ir

```
▶ print('total training cat images:', 1
print('total training dog images:', 1
print('total validation cat images:',
print('total validation dog images:',
```

```
total training cat images: 1000
total training dog images: 1000
total validation cat images: 500
total validation dog images: 500
```

```
[11] from tensorflow.keras import layers
from tensorflow.keras import Model
```

```
▶ # Our input feature map is 150x150x3: 150x1.
# the three color channels: R, G, and B
img_input = layers.Input(shape=(150, 150, 3
```

```
# First convolution extracts 16 filters tha
# Convolution is followed by max-pooling la
x = layers.Conv2D(16, 3, activation='relu')
x = layers.MaxPooling2D(2)(x)
```

```
# Second convolution extracts 32 filters th
# Convolution is followed by max-pooling la
x = layers.Conv2D(32, 3, activation='relu')
x = layers.MaxPooling2D(2)(x)
```

```
# Third convolution extracts 64 filters tha
# Convolution is followed by max-pooling la
x = layers.Conv2D(64, 3, activation='relu')
x = layers.MaxPooling2D(2)(x)
```

```
▶ from tensorflow.keras.preprocessing.image import ImageDataGenerator

# All images will be rescaled by 1./255
train_datagen = ImageDataGenerator(rescale=1./255)
val_datagen = ImageDataGenerator(rescale=1./255)

# Flow training images in batches of 20 using train_datagen generat
train_generator = train_datagen.flow_from_directory(
    train_dir, # This is the source directory for training ima
    target_size=(150, 150), # All images will be resized to 15
    batch_size=20,
    # Since we use binary_crossentropy loss, we need binary lab
    class_mode='binary')

# Flow validation images in batches of 20 using val_datagen generat
validation_generator = val_datagen.flow_from_directory(
    validation_dir,
    target_size=(150, 150),
    batch_size=20,
    class_mode='binary')
```

```
Found 2000 images belonging to 2 classes.
Found 1000 images belonging to 2 classes.
```

Training

Let's train on all 2,000 images available, for 15 epochs, and validate on all 1,000 validation images.

```
history = model.fit_generator(  
    train_generator,  
    steps_per_epoch=100, # 2000 images = batch_size * steps  
    epochs=15,  
    validation_data=validation_generator,  
    validation_steps=50, # 1000 images = batch_size * steps  
    verbose=2)
```

```
/usr/local/lib/python3.7/dist-packages/tensorflow/python/keras/engine/training.py:1  
warnings.warn("`Model.fit_generator` is deprecated and '  
Epoch 1/15  
100/100 - 40s - loss: 0.8383 - acc: 0.5505 - val_loss: 0.6629 - val_acc: 0.6150  
Epoch 2/15  
100/100 - 8s - loss: 0.6415 - acc: 0.6420 - val_loss: 0.6087 - val_acc: 0.6670  
Epoch 3/15  
100/100 - 8s - loss: 0.5803 - acc: 0.7095 - val_loss: 0.6058 - val_acc: 0.6770  
Epoch 4/15  
100/100 - 8s - loss: 0.5173 - acc: 0.7775 - val_loss: 0.7317 - val_acc: 0.6690  
Epoch 5/15  
100/100 - 8s - loss: 0.4405 - acc: 0.8040 - val_loss: 0.7667 - val_acc: 0.6730  
Epoch 6/15  
100/100 - 8s - loss: 0.3744 - acc: 0.8405 - val_loss: 0.8034 - val_acc: 0.6960  
Epoch 7/15  
100/100 - 8s - loss: 0.2713 - acc: 0.8905 - val_loss: 0.8882 - val_acc: 0.6930  
Epoch 8/15  
100/100 - 8s - loss: 0.2026 - acc: 0.9190 - val_loss: 0.8854 - val_acc: 0.7070  
Epoch 9/15  
100/100 - 8s - loss: 0.1391 - acc: 0.9495 - val_loss: 1.3200 - val_acc: 0.6850  
Epoch 10/15  
100/100 - 8s - loss: 0.1057 - acc: 0.9695 - val_loss: 1.7593 - val_acc: 0.6970  
Epoch 11/15
```

Visualizing Intermediate Representations

To get a feel for what kind of features our convnet has learned, one fun thing to do is to visualize through the convnet.

Let's pick a random cat or dog image from the training set, and then generate a figure where each row is a specific filter in that output feature map. Rerun this cell to generate intermediate repre

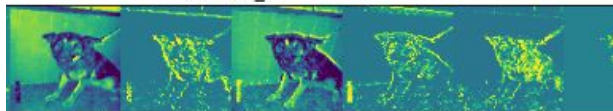
```
import numpy as np  
import random  
from tensorflow.keras.preprocessing.image import img_to_array, load_img  
  
# Let's define a new Model that will take an image as input, and will output  
# intermediate representations for all layers in the previous model after  
# the first.  
successive_outputs = [layer.output for layer in model.layers[1:]]  
visualization_model = Model(img_input, successive_outputs)  
  
# Let's prepare a random input image of a cat or dog from the training set.  
cat_img_files = [os.path.join(train_cats_dir, f) for f in train_cat_filenames]  
dog_img_files = [os.path.join(train_dogs_dir, f) for f in train_dog_filenames]  
img_path = random.choice(cat_img_files + dog_img_files)  
  
img = load_img(img_path, target_size=(150, 150)) # this is a PIL image  
x = img_to_array(img) # Numpy array with shape (150, 150, 3)  
x = x.reshape((1,) + x.shape) # Numpy array with shape (1, 150, 150, 3)  
  
# Rescale by 1/255  
x /= 255
```

er.py:43: RuntimeWarning: invalid value encounte:

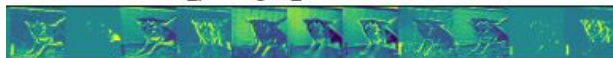
input_2



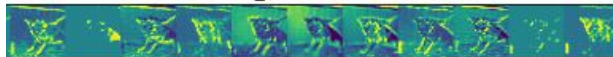
conv2d_3



max_pooling2d_3



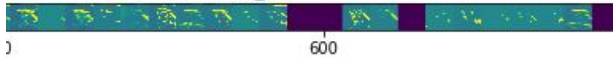
conv2d_4



max_pooling2d_4



conv2d_5



▼ Evaluating Accuracy and Loss for the Model

Let's plot the training/validation accuracy and loss as collected du

```
▶ # Retrieve a list of accuracy results on training and
# sets for each training epoch
acc = history.history['acc']
val_acc = history.history['val_acc']

# Retrieve a list of list results on training and vali
# sets for each training epoch
loss = history.history['loss']
val_loss = history.history['val_loss']

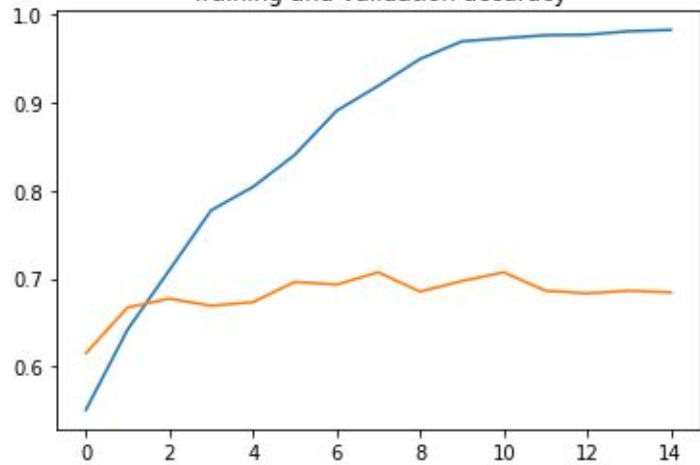
# Get number of epochs
epochs = range(len(acc))

# Plot training and validation accuracy per epoch
plt.plot(epochs, acc)
plt.plot(epochs, val_acc)
plt.title('Training and validation accuracy')

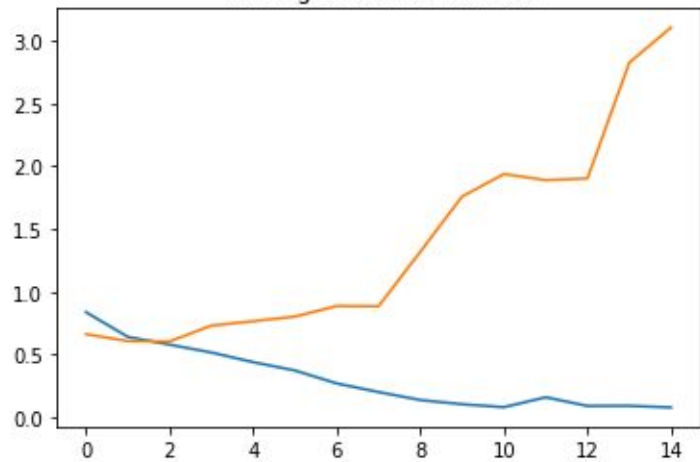
plt.figure()

# Plot training and validation loss per epoch
plt.plot(epochs, loss)
plt.plot(epochs, val_loss)
plt.title('Training and validation loss')
```

Training and validation accuracy



Training and validation loss



Aprenda no colab:

[Exercise 1: Building a Convnet from Scratch](#)

[Exercise 2: Reducing Overfitting](#)

[Feature Extraction and Fine-Tuning](#)

little more

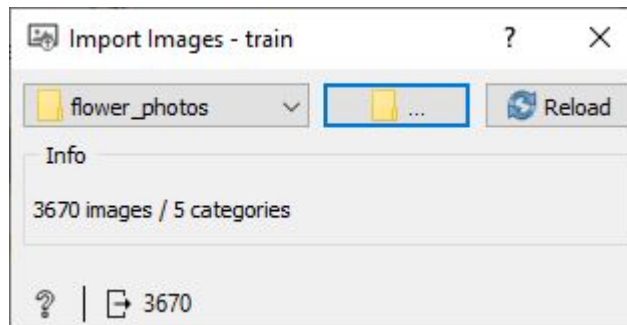
With flowers?

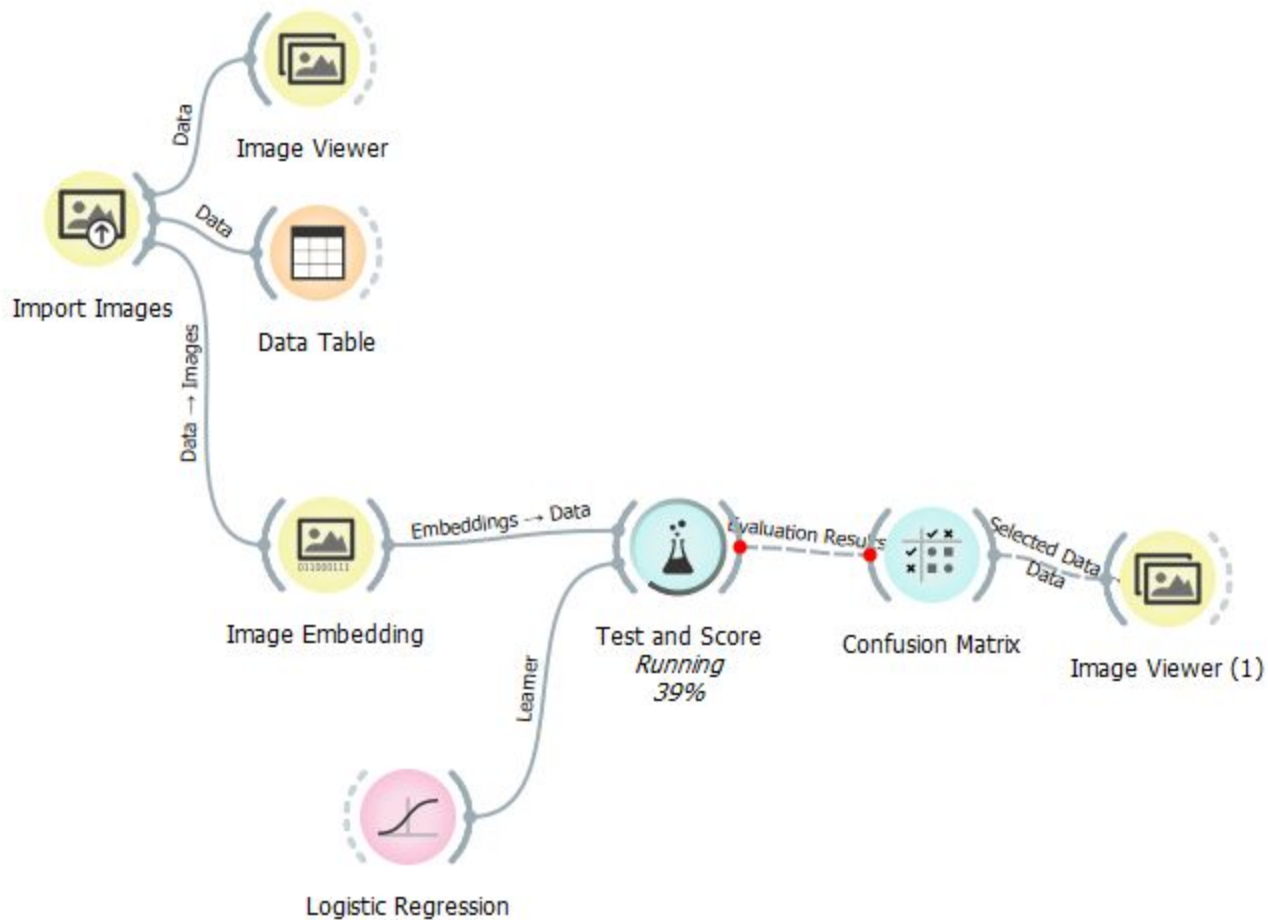
Other

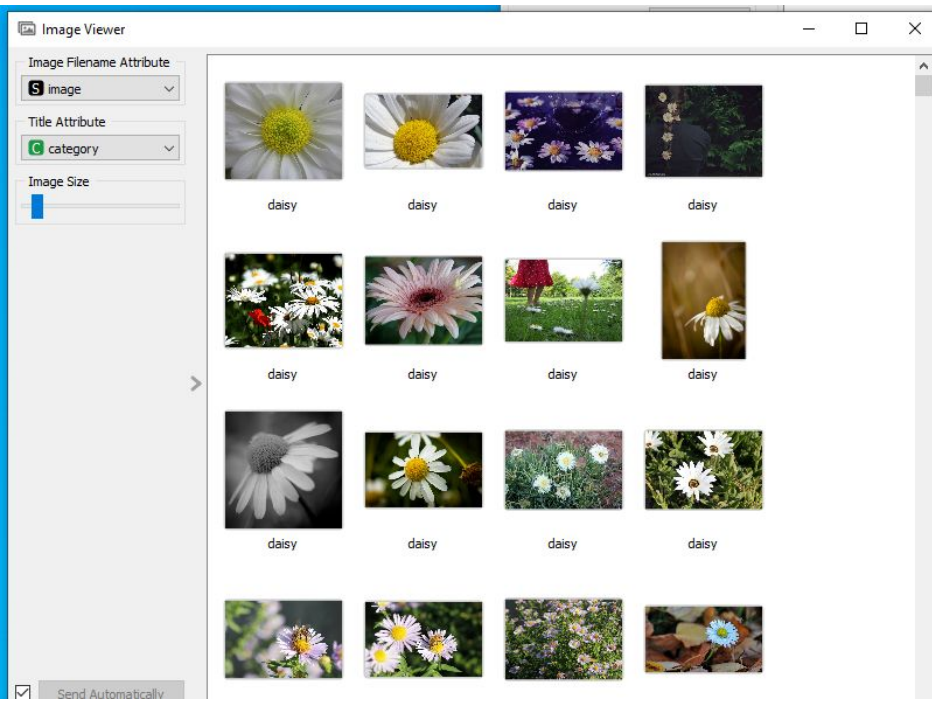
try yourself with other example

now, with this imagens

https://storage.googleapis.com/download.tensorflow.org/example_images/flower_photos.tgz







Data Table

Info
 3540 instances (no missing data)
 No features
 Target with 5 values
 5 meta attributes

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

Selection
 Select full rows

origir	category	image name	image	ange test/flower_tf	size	width
1	daisy	100080576_f52e...	daisy\10008057...	image	26797	320
2	daisy	10140303196_b...	daisy\10140303...		117247	500
3	daisy	10172567486_27...	daisy\10172567...		102862	500
4	daisy	10172636503_21...	daisy\10172636...		27419	320
5	daisy	102841525_bd6...	daisy\10284152...		132803	500
6	daisy	1031799732_e7f...	daisy\10317997...		102618	500
7	daisy	10391248763_1...	daisy\10391248...		51688	320
8	daisy	10437754174_22...	daisy\10437754...		13946	171
9	daisy	10437770546_8...	daisy\10437770...		13518	240
10	daisy	10437929963_b...	daisy\10437929...		84219	500
11	daisy	10466290366_cc...	daisy\10466290...		172328	500
12	daisy	10466558316_a7...	daisy\10466558...		129584	500
13	daisy	10555749515_13...	daisy\10555749...		86878	500
14	daisy	10555815624_d...	daisy\10555815...		93133	500
15	daisy	10555826524_42...	daisy\10555826...		59028	320
16	daisy	10559679065_50...	daisy\10559679...		100584	500
17	daisy	105806915_a9c1...	daisy\10580691...		25620	320
18	daisy	10712722853_56...	daisy\10712722...		63389	500
19	daisv	107592979_aaa9...	daisv\10759297...		31691	240

Restore Original Order

Send Automatically:

Image attribute dialog:

- Image attribute: image
- Embedder: Inception v3
- Google's Inception v3 model trained on ImageNet.
- Apply Automatically
- Cancel
- 3540

Logistic Regression dialog:

- Name: Logistic Regression
- Regularization type: Ridge (L2)
- Strength: Weak to Strong (C=1)
- Balance class distribution:
- Apply Automatically:

Test and Score dialog:

- Sampling: Cross validation
- Number of folds: 10
- Stratified
- Cross validation by feature
- Random sampling
- Repeat train/test: 10
- Training set size: 66 %
- Stratified
- Leave one out
- Test on train data
- Test on test data
- Target Class

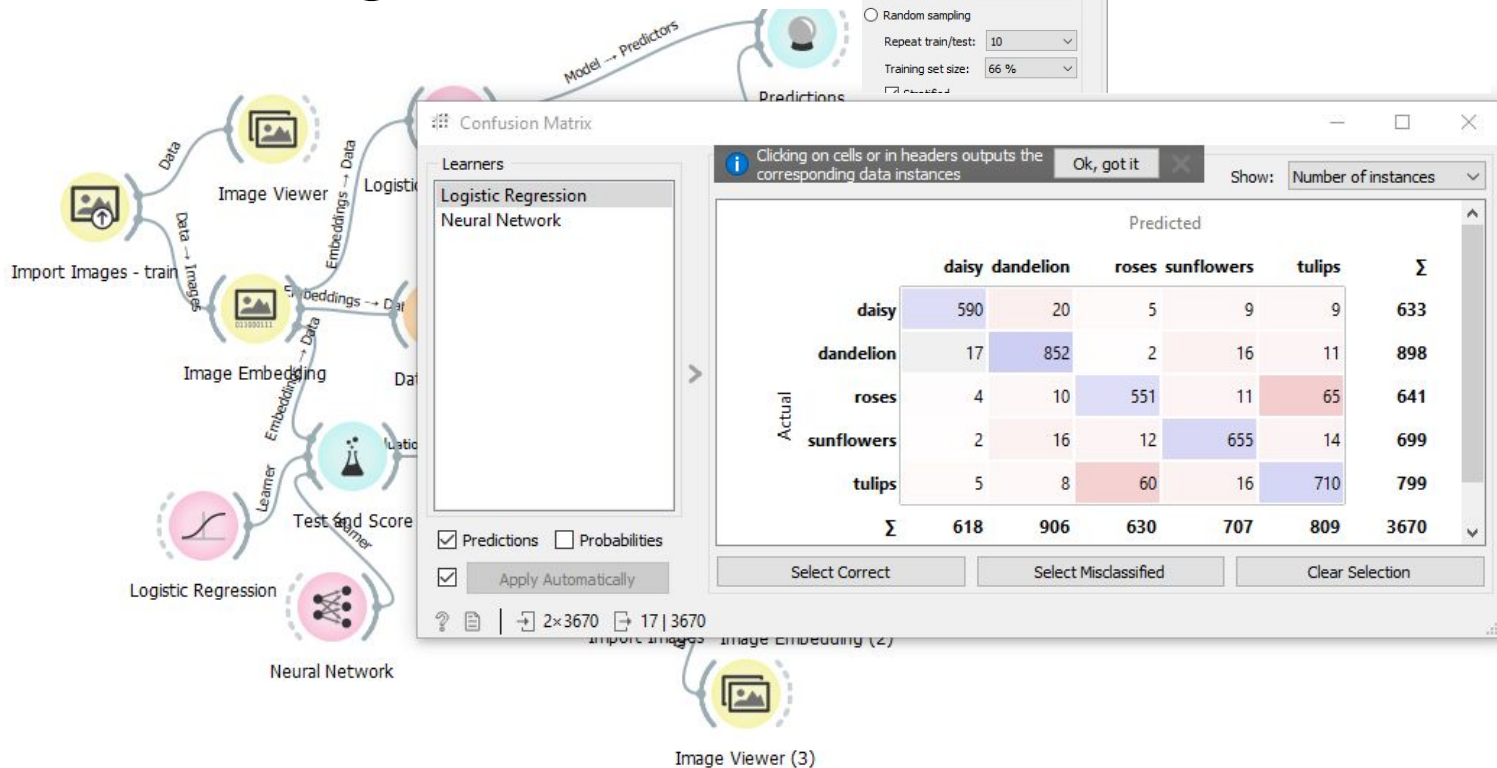
Test and Score Results Table:

Model	AUC	CA	F1	Precision	Recal
Logistic Regression	0.990	0.913	0.913	0.913	0.913

Model Comparison by AUC:

Model	AUC
Logistic Regression	0.990

orange



Test and Score

Sampling

Cross validation

Number of folds: 10

Stratified

Cross validation by feature

Random sampling

Repeat train/test: 10

Training set size: 66 %

Click on the table header to select shown columns

Model	AUC	CA	F1	Precision	Recall
Neural Network	0.991	0.913	0.913	0.914	0.913
Logistic Regression	0.990	0.915	0.915	0.915	0.915

Confusion Matrix

Learners

- Logistic Regression
- Neural Network

Predictions Probabilities

Apply Automatically

Clicking on cells or in headers outputs the corresponding data instances

Show: Number of instances

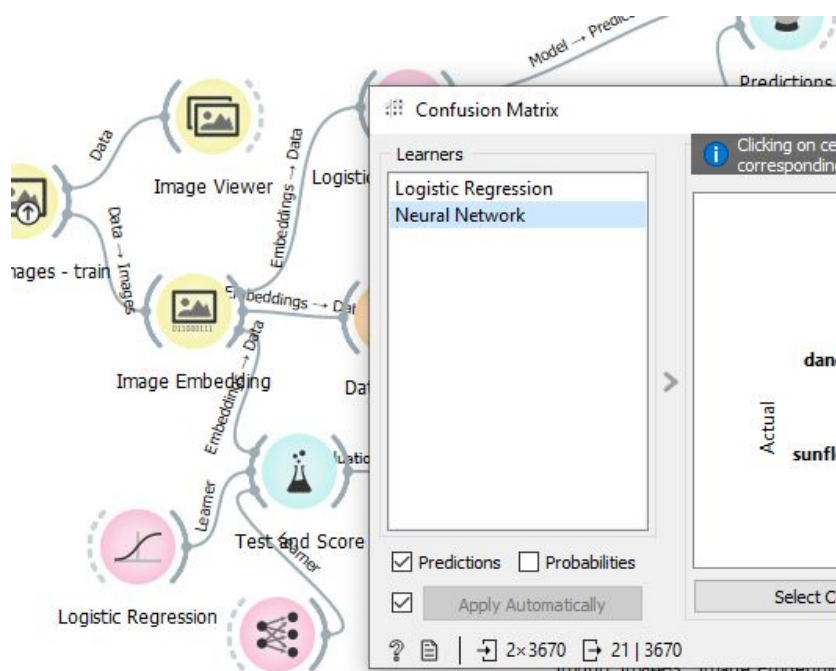
		Predicted					Σ
		daisy	dandelion	roses	sunflowers	tulips	
Actual	daisy	590	20	5	9	9	633
	dandelion	17	852	2	16	11	898
	roses	4	10	551	11	65	641
	sunflowers	2	16	12	655	14	699
	tulips	5	8	60	16	710	799
Σ		618	906	630	707	809	3670

Select Correct Select Misclassified Clear Selection

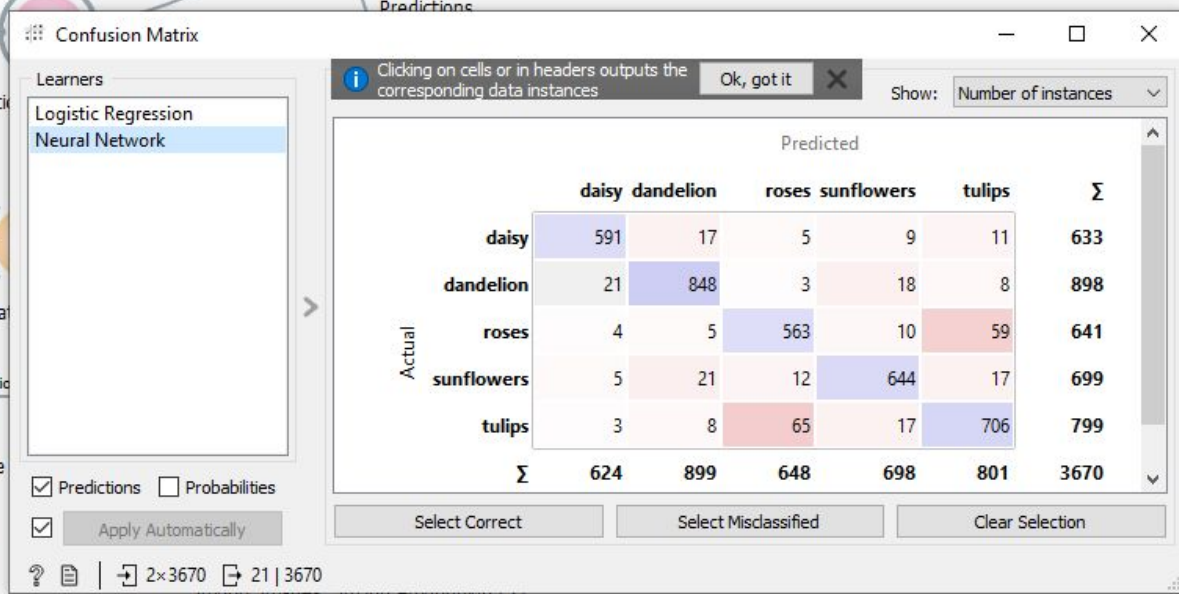
2x3670 17 | 3670

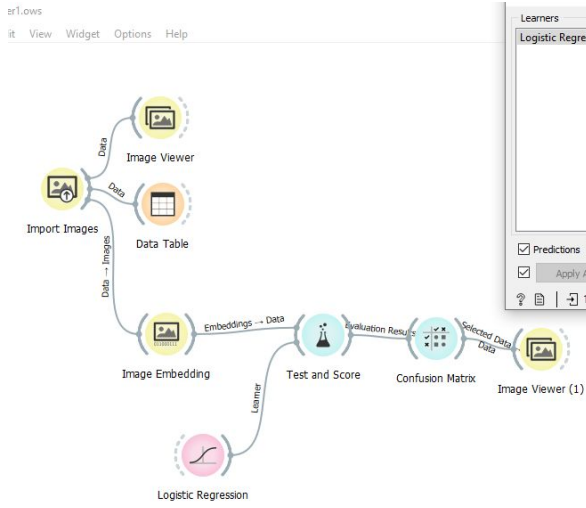
Image Viewer (3)

orange



category	image name	image	size	width
daisy	1031799732_e7f...	daisy\10317997...	102618	500
daisy	10391248763_1...	daisy\10391248...	51688	320
daisy	10437754174_22...	daisy\10437754...	13946	171
daisy	10437770546_8...	daisy\10437770...	13518	240
daisy	10437929963_b...	daisy\10437929...	84219	500
daisy	10466290366_cc...	daisy\10466290...	172328	500
daisy	10466558316_a7...	daisy\10466558...	129584	500
daisy	10555749515_13...	daisy\10555749...	86878	500
daisy	10555815624_d...	daisy\10555815...	93133	500
daisy	10555826524_47...	daisy\10555826...	50928	320





Clicking on cells or in headers outputs the corresponding data instances Ok, got it Show: Number of instances

		Predicted				Σ	
		daisy	dandelion	roses	sunflowers		tulips
Actual	daisy	565	20	3	10	607	
	dandelion	17	826	2	20	872	
	roses	3	7	533	13	615	
	sunflowers	4	16	13	627	673	
	tulips	3	10	65	15	680	
Σ		592	879	616	685	768	3540

Predictions Probabilities
 Apply Automatically

1x3540 17 | 3540

Select Correct Select Misclassified Clear Selection

Image Viewer (1)

Image Filename Attribute: image

Title Attribute: category

Image Size: [Slider]

dandelion dandelion dandelion dandelion

dandelion dandelion dandelion dandelion

dandelion dandelion dandelion dandelion

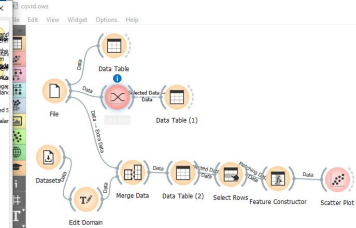
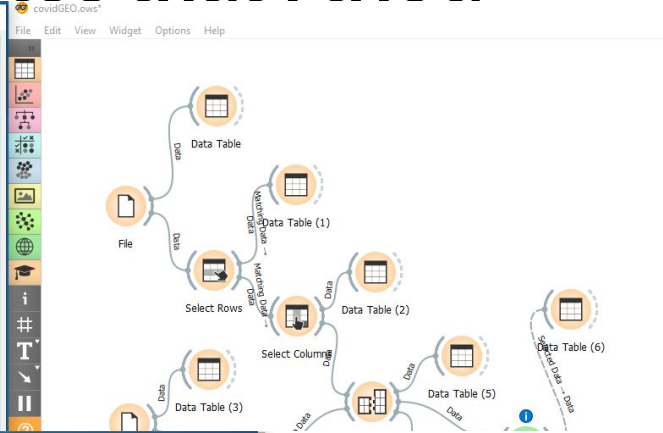
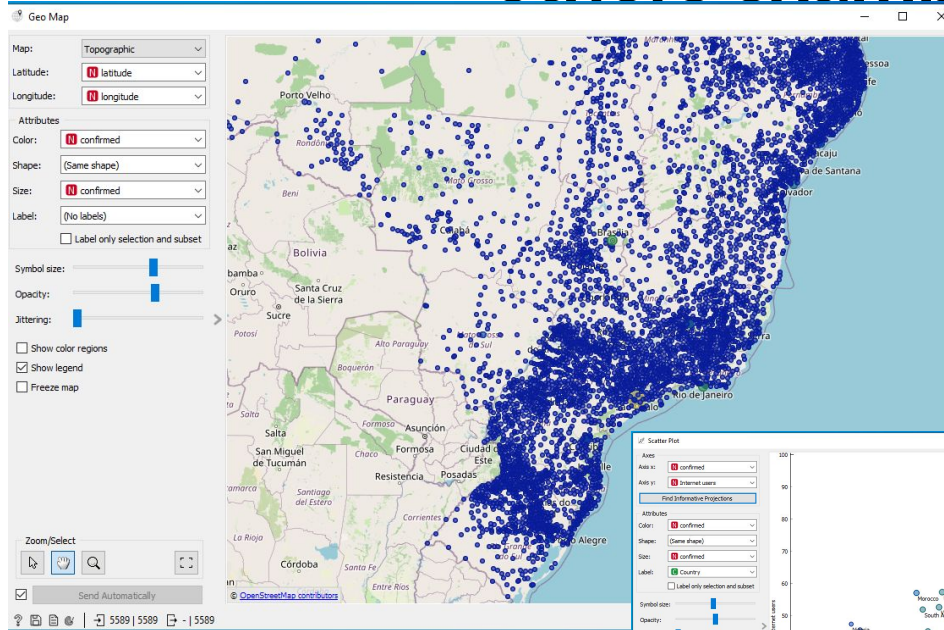
coding an example - flowers

[Colab + python + keras + matplotlib + numpy - aprendendo e classificando flores](#)

—

others

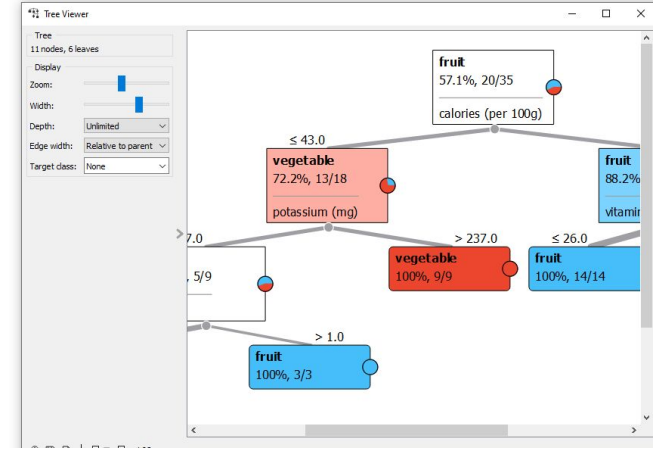
others examples that i tried



others examples that i tried

The screenshot shows the Orange3 data mining software interface. A workflow is visible with the following widgets: File, Data Table, Tree, Logistic Regression, and Test and Score. Several configuration windows are open:

- Tree Configuration:**
 - Name: Tree
 - Parameters:
 - Induce binary tree:
 - Min. number of instances in leaves: 2
 - Do not split subsets smaller than: 5
 - Limit the maximal tree depth to: 100
 - Classification:
 - Stop when majority reaches [%]: 95
 - Buttons: Apply Automatically, Save, Undo, Redo, Refresh, Close
- Logistic Regression Configuration:**
 - Name: Logistic Regression
 - Regularization type: Ridge (L2)
 - Strength: C=1 (slider between Weak and Strong)
 - Balance class distribution:
- Test and Score Configuration:**
 - Sampling:
 - Cross validation:
 - Number of folds: 10
 - Stratified:
 - Cross validation by feature: (dropdown)
 - Random sampling:
 - Repeat train/test: 10
 - Training set size: 66%
 - Stratified:
 - Leave one out:
 - Test on train data:
 - Test on test data:
 - Target Class: (Average over classes)
 - Model Comparison: Area under ROC curve
 - Negligible difference: 0.1



Predictions

Show probabilities for

	Logistic Regression	Tree	name	vitamin
1	fruit	fruit	?	1.0
2	vegetable	fruit	?	15.0
3	fruit	fruit	?	0.0

fruit
vegetable

others examples that i tried

The screenshot displays the diabetes.ows software interface. The main workspace shows a workflow diagram with nodes for File, Data Table, Data, Logistic Regression, Naive Bayes, Neural Network, Learner, Predictions, and Save Model. Overlaid windows provide detailed information:

- Test and Score:** Shows sampling options (Cross validation, Random sampling) and a table of model performance metrics.
- Predictions:** Displays a table of predicted outcomes for individual data points, including probabilities and feature values.
- Model Comparison:** Compares the performance of Neural Network, Naive Bayes, and Logistic Regression models based on AUC, CA, F1, Precision, and Recall.

	Neural Network	Naive Bayes	Logistic Regression
Neural Network		0.721	0.473
Naive Bayes	0.279		0.312
Logistic Regression	0.527	0.688	
(Average over classes)			

	Naive Bayes	Neural Network	Outcome	Pregnancies	Glucose	B ⁺
0						
1	0.04 : 0.96 ...	0.40 : 0.60 - 1	1	6	148	72
2	0.98 : 0.02 ...	0.96 : 0.04 - 0	0	1	85	66
3	0.20 : 0.80 ...	0.07 : 0.93 - 1	1	8	183	64
4	0.99 : 0.01 ...	0.99 : 0.01 - 0	0	1	89	66
5	0.17 : 0.83 ...	0.26 : 0.74 - 1	1	0	137	40
6	0.90 : 0.10 ...	0.85 : 0.15 - 0	0	5	116	74
7	0.99 : 0.01 ...	0.96 : 0.04 - 0	1	3	78	50
8	0.49 : 0.51 ...	0.39 : 0.61 - 1	0	10	115	0
9	0.14 : 0.86 ...	0.04 : 0.96 - 1	1	2	197	70
10	0.54 : 0.46 ...	0.89 : 0.11 - 0	1	8	125	96
11	0.47 : 0.53 ...	0.82 : 0.18 - 0	0	4	110	92
12	0.05 : 0.95 ...	0.07 : 0.93 - 1	1	10	168	74
13	0.30 : 0.70 ...	0.63 : 0.37 - 0	0	10	139	80
14	0.38 : 0.62 ...	0.06 : 0.94 - 1	1	1	189	60

Model	AUC	CA	F1	Precision	Recall
Naive Bayes	0.829	0.749	0.752	0.757	0.749
Neural Network	0.893	0.827	0.824	0.824	0.827

others examples that i tried

The screenshot displays a workflow application interface with a main workspace and two floating windows.

Main Workspace: A workflow diagram showing the following steps:

- Import Images** (Data source) feeds into **Image Embedding** (Data).
- Image Embedding** feeds into **Data Table** (Data).
- Data Table** feeds into **Neighbors (1)** (Data).
- Neighbors (1)** feeds into **Image Viewer (1)** (Data).

Image Viewer (Main): A window titled "Image Viewer" with a sidebar containing the following controls:

- Image Filename Attribute:** dropdown menu set to "image".
- Title Attribute:** dropdown menu set to "image name".
- Image Size:** slider control.

The main area of this window displays a grid of 12 image thumbnails with their corresponding names:

Albert Edelfelt	Antoine Vollon	Anton Karinger	Canaletto
Claude Monet	Dario de Regoyos	Dominik Skutecky	Jan Hackaert
Johann Christian Vollerdt			

Image Viewer (1): A smaller window titled "Image Viewer (1)" with a sidebar containing the following controls:

- Image Filename Attribute:** dropdown menu set to "image".
- Title Attribute:** dropdown menu set to "image name".
- Image Size:** slider control.

The main area of this window displays a grid of 2 image thumbnails with their corresponding names:

Claude Monet	Dario de Regoyos
--------------	------------------

others examples that i tried

The image displays several screenshots of a software interface for hierarchical clustering of images. The main components shown are:

- Hierarchical Clustering Panel:** Shows a dendrogram with a linkage method of 'Average'. The 'Annotations' dropdown is set to 'image name'. The 'Selection' radio button is set to 'Manual'. The 'Height ratio' is set to 75.0% and 'Top N' is set to 5. The 'Zoom' is set to 20x-20.
- Image Viewer (1):** Displays a grid of images for various animal classes: calf, cow, foal, goat, horse, kid, lamb, ox, and sheep. The 'Image Name Attribute' is set to 'image name'.
- Image Viewer (2):** Displays a grid of images for classes: cat and rabbit. The 'Image Name Attribute' is set to 'image name'.
- Image Viewer (3):** Displays a grid of images for classes: cat and rabbit. The 'Image Name Attribute' is set to 'image name'.
- Image Viewer (4):** Displays a grid of images for classes: cat and rabbit. The 'Image Name Attribute' is set to 'image name'.

At the bottom, two flowcharts illustrate the data processing pipeline:

```
graph TD
    subgraph Pipeline 1
        A[Import Images] --> B[Data Table]
        B --> C[Image Embedding]
        C --> D[Distances]
        D --> E[Hierarchical Clustering]
        E --> F[Image Viewer (1)]
    end
    subgraph Pipeline 2
        G[Import Images] --> H[Data Table]
        H --> I[Image Embedding]
        I --> J[Distances]
        J --> K[Hierarchical Clustering]
        K --> L[Image Viewer (1)]
    end
```

Conclusion

compare

KNN

		Predicted		
		cats	dogs	Σ
Actual	cats	3957	43	4000
	dogs	14	3991	4005
Σ		3971	4034	8005

SVM

		Predicted		
		cats	dogs	
Actual	cats	3969	31	4000
	dogs	18	3987	4005
Σ		3987	4018	8005

NEURAL NETWORK

		Predicted		
		cats	dogs	Σ
Actual	cats	3971	29	4000
	dogs	20	3985	4005
Σ		3991	4014	8005

Random Forest

		Predicted		
		cats	dogs	Σ
Actual	cats	3951	49	4000
	dogs	22	3983	4005
Σ		3973	4032	8005

LOGIC REGRESSION

		Predicted		
		cats	dogs	Σ
Actual	cats	3972	28	4000
	dogs	19	3986	4005
Σ		3991	4014	8005



Naive Bayes

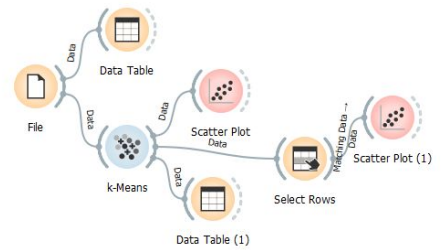
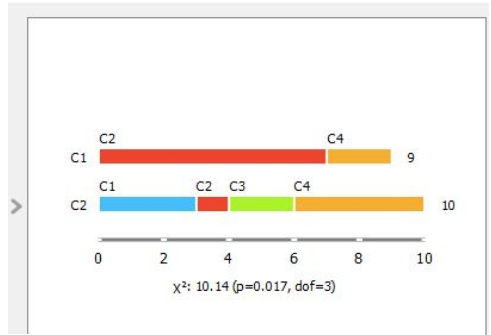
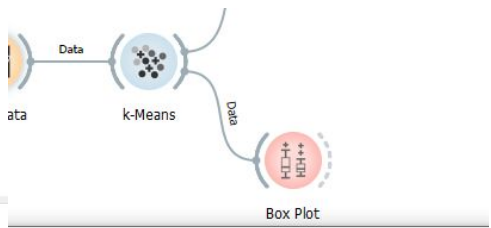
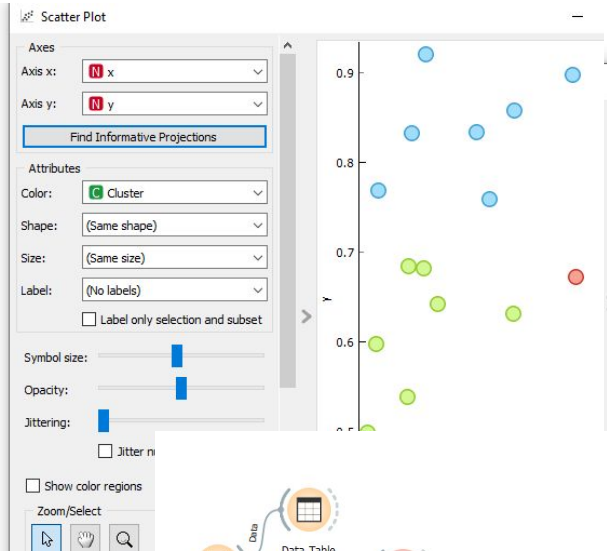
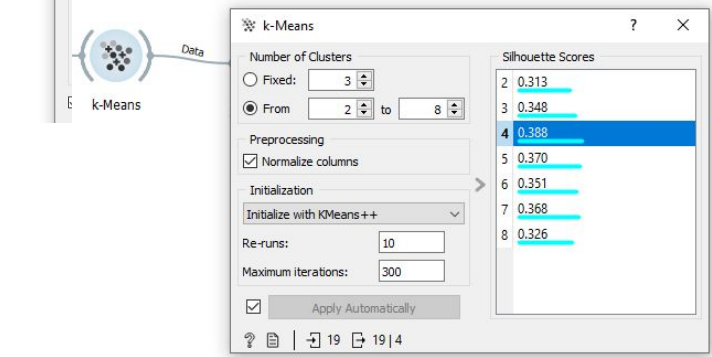
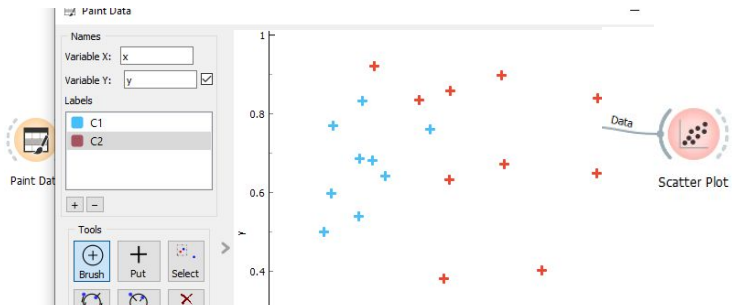
		Predicted		
		cats	dogs	Σ
Actual	cats	3880	120	4000
	dogs	7	3998	4005
Σ		3887	4118	8005



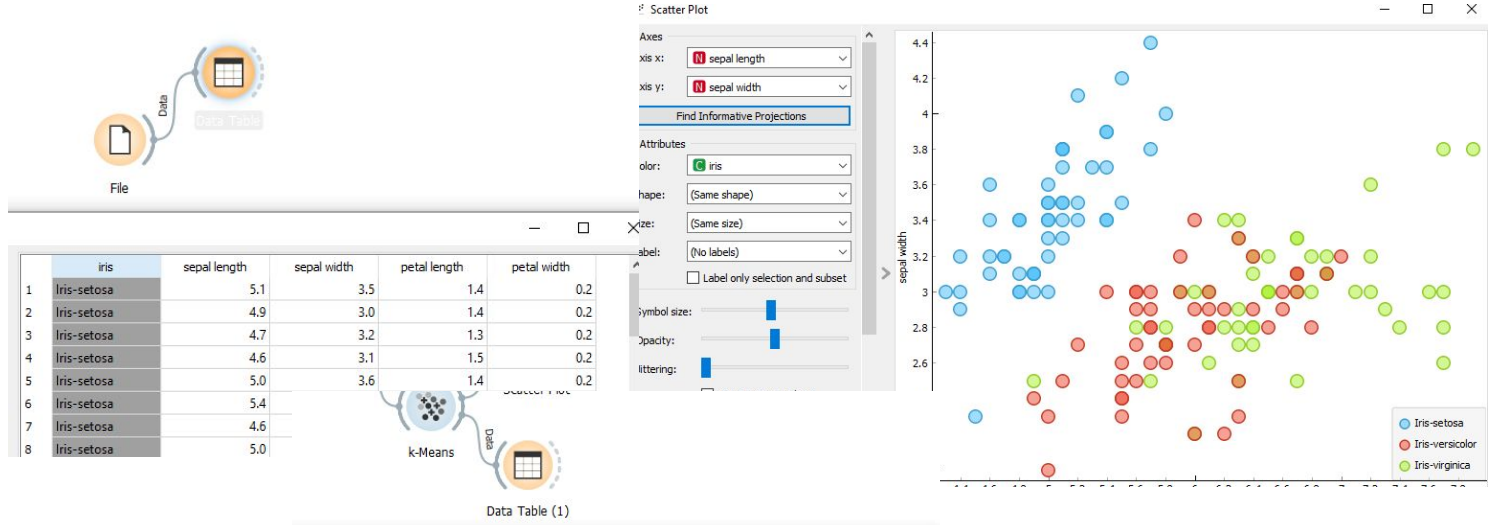
—

+

K-means

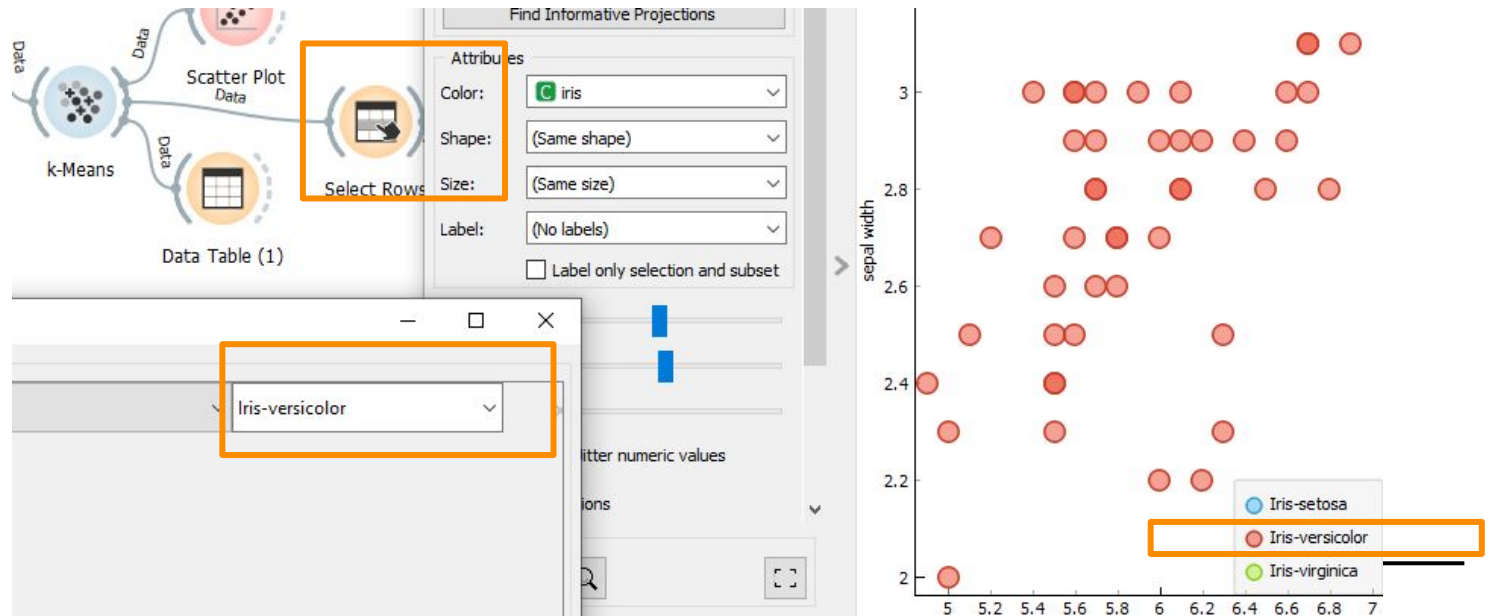


Agrupamento - K-means - iris



iris	Cluster	Silhouette	sepal length	sepal width	pe
Iris-setosa	C1	0.706706	5.1	3.5	
Iris-setosa	C1	0.683687	4.9	3.0	
Iris-setosa	C1	0.700571	4.7	3.2	
Iris-setosa	C1	0.691771	4.6	3.1	
Iris-setosa	C1	0.704669	5.0	3.6	
Iris-setosa	C1	0.677309	5.4	3.9	
Iris-setosa	C1	0.701974	4.6	3.4	
Iris-setosa	C1	0.707426	5.0	3.4	

Agrupamento - K-means - iris



iris

confusion matrix

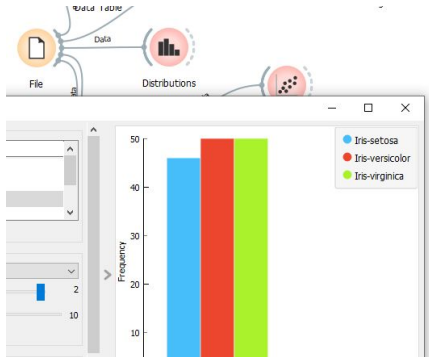
		Predicted			Σ
		Iris-setosa	Iris-versicolor	Iris-virginica	
Actual	Iris-setosa	50	0	0	50
	Iris-versicolor	0	46	4	50
	Iris-virginica	0	3	47	50
Σ		50	49	51	150

random forest e logic regression

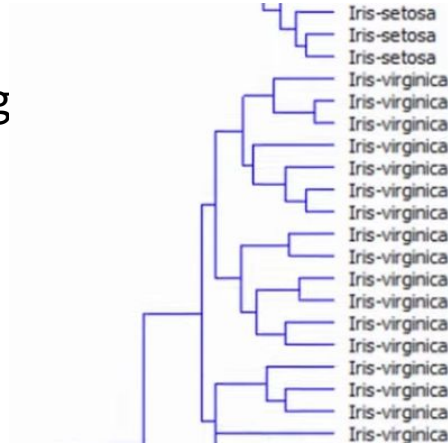
<https://orange3.readthedocs.io/projects/orange-data-mining-library/en/latest/reference/regression.html#random-forest>

Method	AUC	CA	F1	Precision	Recall
Logistic Regression	0.970	0.960	0.960	0.962	0.960
RF Classification Learner	0.965	0.953	0.953	0.953	0.953

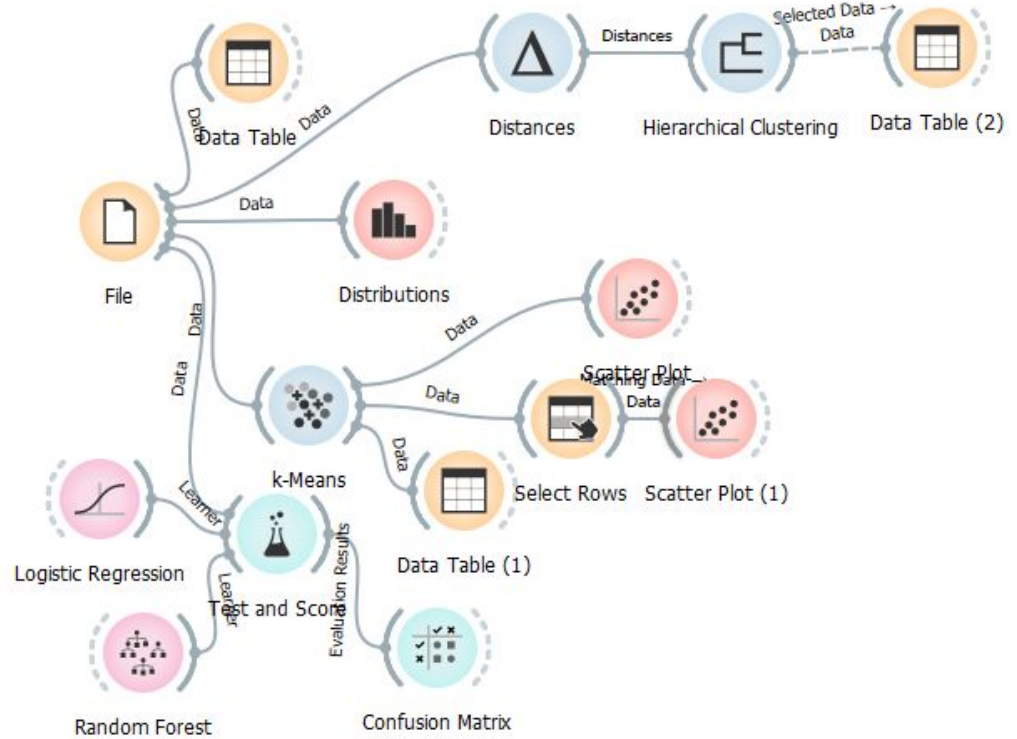
distributions



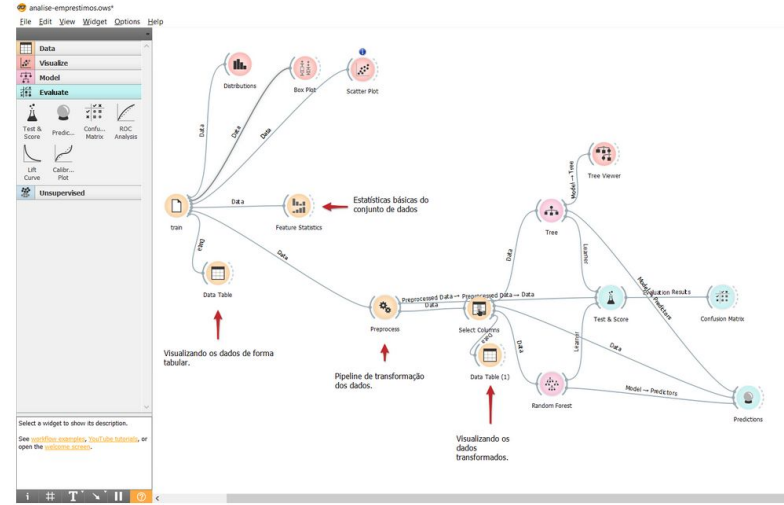
iris clustering



iris



example loan - orange



Conhecendo as previsões de cada um dos modelos

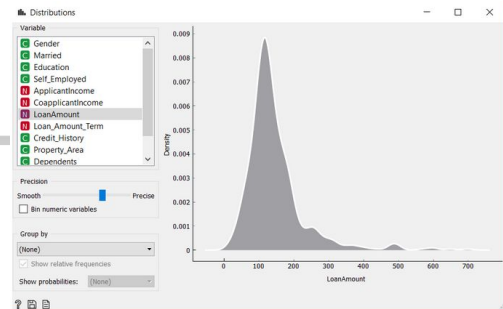
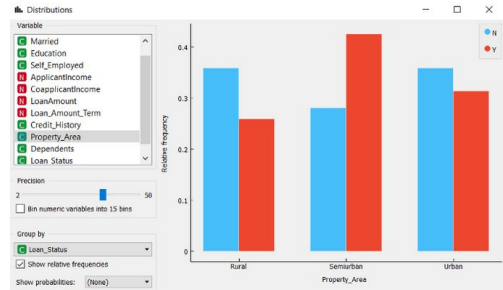
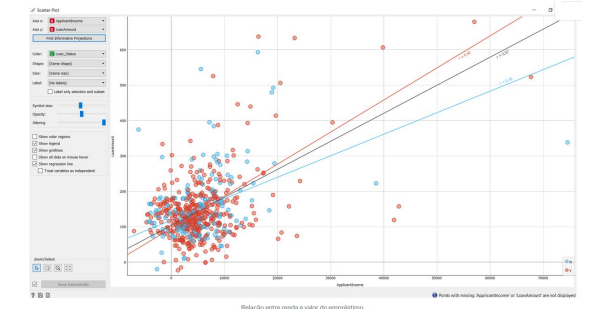
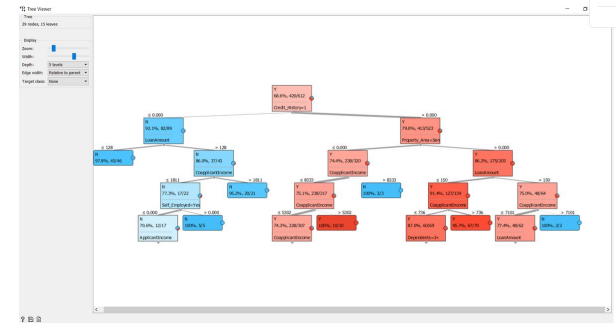


Gráfico de densidade do valor de empréstimo solicitado.

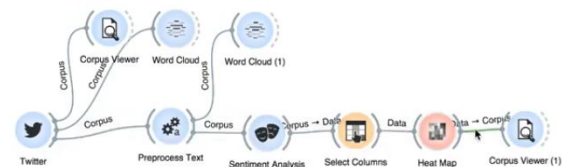


Relação entre renda e o valor do empréstimo.

videos

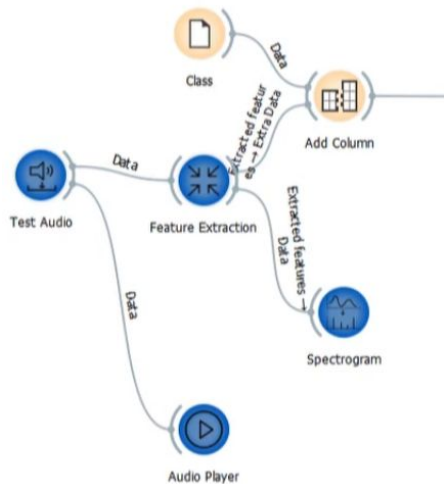
moods on twitter

<https://www.youtube.com/watch?v=1stp41wz3LA>



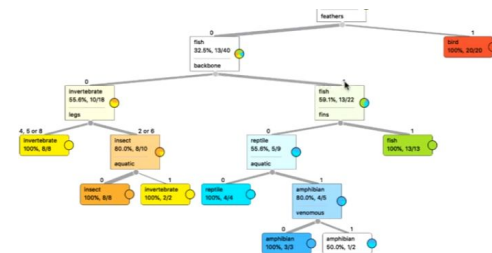
classify tree - fish - reptile - bird - insect

https://www.youtube.com/watch?v=eTe8r5tnfRA&list=PLZ3V9XyVA52_XSu059rS9eqi7P2nY6Qe9&index=7



instrument sound classifier

<https://www.youtube.com/watch?v=BO0LUIMx6qs>



svm - unicamp - orange

<https://www.youtube.com/watch?v=1KFx3861Djs>

videos - image challenge



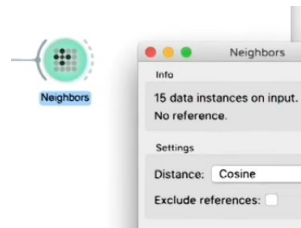
Canaletto Nikola Petrov Peder Severin Kroyer Dario de Regoyos



Albert Edelfelt Rudolf Bernt Monet Jan Hackaert



Johann Christian Vollerdt Antoine Vollon Claude Monet Dominik Skutecky



Neighbors

Info

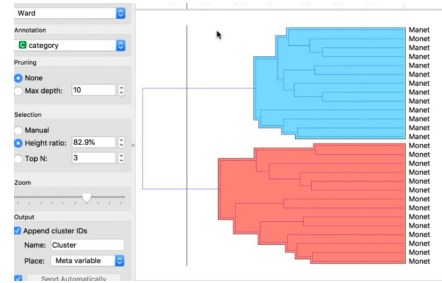
15 data instances on input.
No reference.

Settings

Distance: Cosine

Exclude references:

monet or manet - inceptionv3 (google)



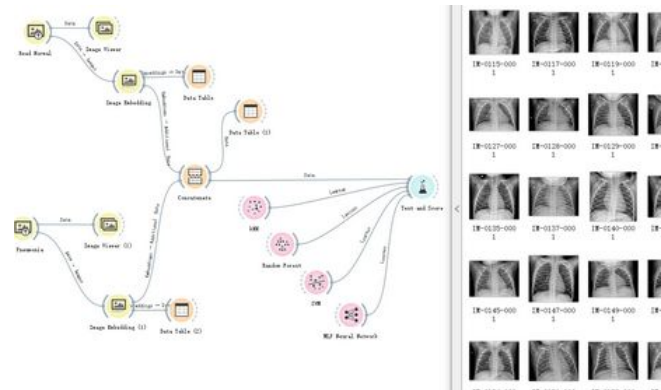
Monet Claude Monet

the lost monet - model pre trained painters

links

Detecting Pneumonia in Chest X-Ray Images under Orange Machine Learning/Deep Learning Platform

<https://towardsdatascience.com/detecting-pneumonia-in-chest-x-ray-images-under-orange-machine-learning-deep-learning-platform-dd7b6ca6bd4c>



curiosity

article in nature communications: **Democratized image analytics by visual programming through integration of deep models and small-scale machine learning.**

<https://www.nature.com/articles/s41467-019-12397-x>

Use of Orange Data Mining Toolbox for Data Analysis in Clinical Decision Making: The Diagnosis of Diabetes Disease.

https://www.researchgate.net/publication/329707993_Use_of_Orange_Data_Mining_Toolbox_for_Data_Analysis_in_Clinical_Decision_Making_The_Diagnosis_of_Diabetes_Disease

Analysis of Heart Disease using in Data Mining Tools Orange and Weka

https://globaljournals.org/GJCST_Volume18/4-Analysis-of-Heart-Disease.pdf

sources

<https://orangedatamining.com/>

—

colab - CIFAR10

Convolutional Neural Network (CNN) - CIFAR10

```
[ ] import tensorflow as tf

from tensorflow.keras import datasets, layers, models
import matplotlib.pyplot as plt
```

Download and prepare the CIFAR10 dataset

The CIFAR10 dataset contains 60,000 color images in 10 classes, with 6,000 images in each class. The dataset is divided into 50,000 training images and 10,000 testing images. The classes are mutually exclusive and there is no overlap between them.

```
[ ] ((train_images, train_labels), (test_images, test_labels)) = datasets.cifar10.load_data()

# Normalize pixel values to be between 0 and 1
train_images, test_images = train_images / 255.0, test_images / 255.0
```

Verify the data

To verify that the dataset looks correct, let's plot the first 25 images from the training set and display the class names

```
▶ class_names = ['airplane', 'automobile', 'bird', 'cat', 'deer',
                 'dog', 'frog', 'horse', 'ship', 'truck']

plt.figure(figsize=(10,10))
for i in range(25):
    plt.subplot(5,5,i+1)
    plt.xticks([])
```

https://colab.research.google.com/drive/1Le_aq2TNmLRnKXr-5t10oua8qiF68WfK?usp=sharing

Create the convolutional base

The 6 lines of code below define the convolutional base using a common pattern: a stack of [Conv2D](#) and [MaxPooling2D](#) layers.

As input, a CNN takes tensors of shape (image_height, image_width, color_channels), ignoring the batch size. If you are not using color images, color_channels refers to (R,G,B). In this example, you will configure your CNN to process inputs of shape (32, 32, 3) format of CIFAR images. You can do this by passing the argument `input_shape` to your first layer.

```
▶ model = models.Sequential()
model.add(layers.Conv2D(32, (3, 3),
                        input_shape=(32, 32, 3)))
model.add(layers.MaxPooling2D((2, 2)))
model.add(layers.Conv2D(64, (3, 3)))
model.add(layers.MaxPooling2D((2, 2)))
model.add(layers.Conv2D(64, (3, 3)))
```

<https://www.cs.toronto.edu/~kriz/cifar.html>

CIFAR-10 dataset consists of 60000 32x32 colour images in 10 classes, with 6000 images per class. There are 50000 training images and 10000 test images.

Compile and train the model

```
[ ] model.compile(optimizer='adam',
                 loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
                 metrics=['accuracy'])

history = model.fit(train_images, train_labels, epochs=10,
                    validation_data=(test_images, test_labels))
```

Evaluate the model

```
[ ] plt.plot(history.history['accuracy'], label='accuracy')
plt.plot(history.history['val_accuracy'], label='val_accuracy')
plt.xlabel('Epoch')
plt.ylabel('Accuracy')
plt.ylim([0.5, 1])
plt.legend(loc='lower right')

test_loss, test_acc = model.evaluate(test_images, test_labels, verbose=0)

[ ] print(test_acc)
```

Your simple CNN has achieved a test accuracy of over 70%. Not bad for a few lines of code! For an advanced example that uses the Keras subclassing API and `tf.GradientTape`, see [this](#).

—

fim
colab

others

[Power BI](#)

[Tableau](#)



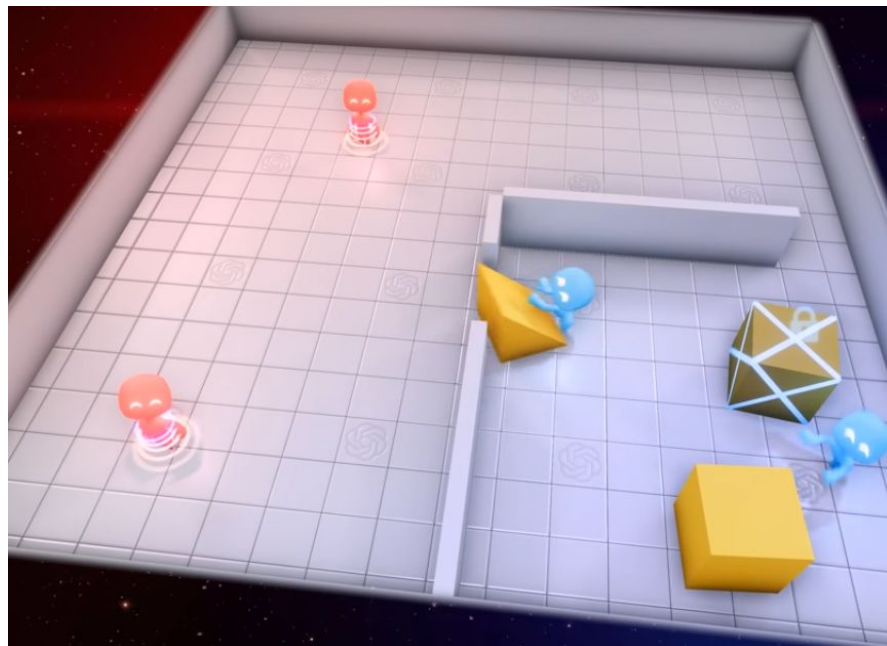
[Modelo Preditivo - Parte 1 - Prevendo a demanda por aluguel de bicicletas usando R e Azure Machine Learning!](#)



open IA

instituição sem fins lucrativos de pesquisa em inteligência artificial (IA), que tem como objetivo promover e desenvolver IA amigável, de tal forma a beneficiar a humanidade como um todo

[Inteligência Artificial brincando de Pique-Esconde](#)



Teachable Machine

<https://teachablemachine.withgoogle.com/>

<https://www.youtube.com/watch?v=T2qQGqZxkD0>

Class 1 

4000 Image Samples



Webcam



Upload



Class 2 

File

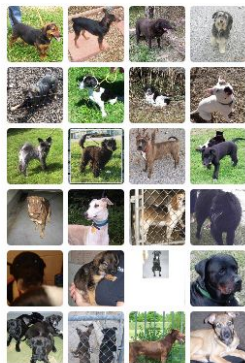
 Choose images from your files,
or drag & drop here


 Import images from
Google Drive



Images will be cropped to square

4005 Image Samples



 Add a class

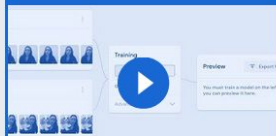
Training

Train Model

Advanced 

2. Train your Model 

Now that you have two classes, you
can train your model here (or add
more classes).



Preview

 Export Model

You must train a model on the left
before you can preview it here.

155 min depois...

Training

Train Model

Advanced ^

Epochs: 50

Batch Size: 16

Learning Rate:

0.001

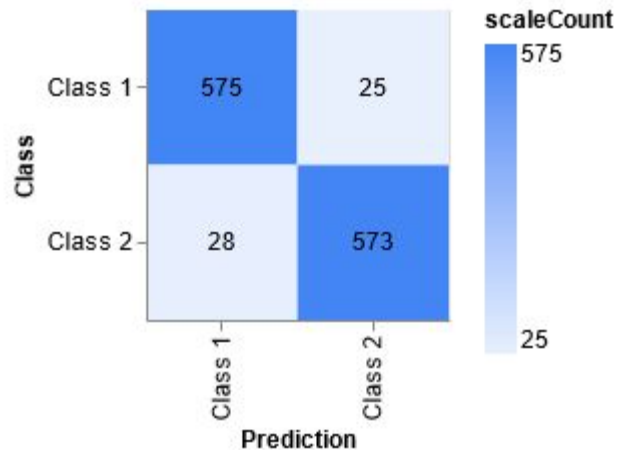
Reset Defaults

Under the hood

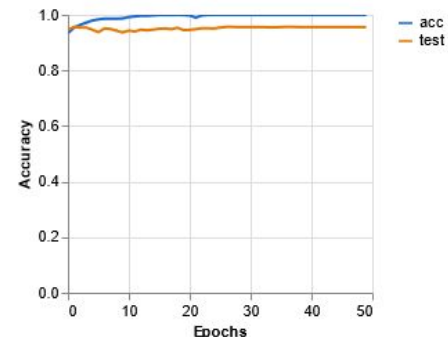
Accuracy per class

CLASS	ACCURACY	# SAMPLES
Class 1	0.96	600
Class 2	0.95	601

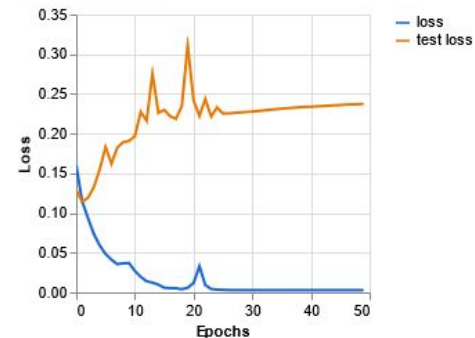
Confusion Matrix



Accuracy per epoch



Loss per epoch

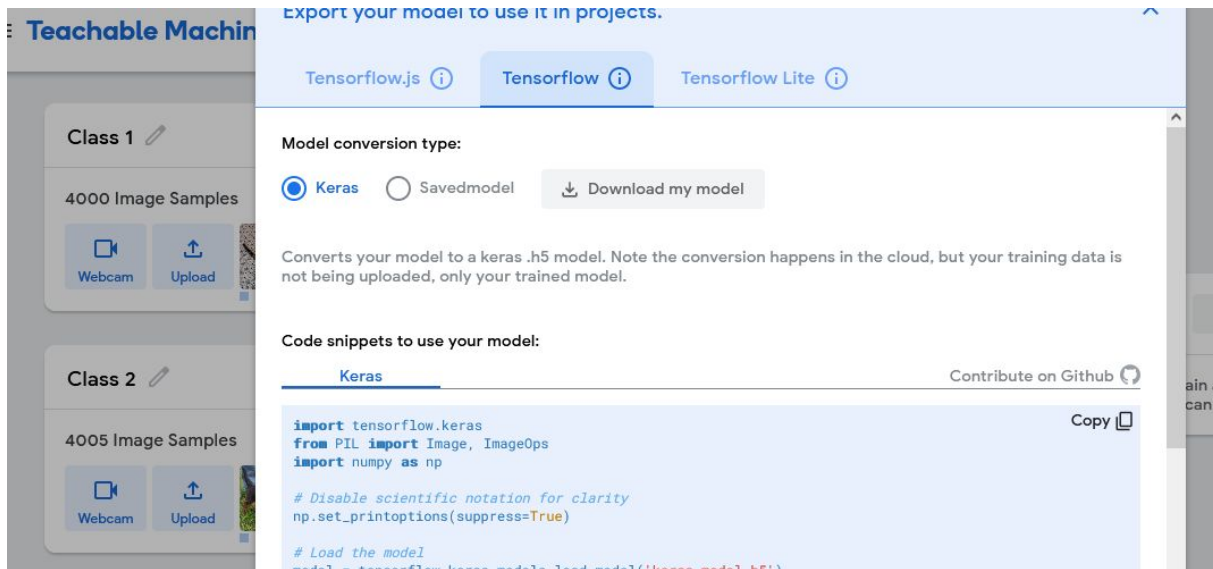


modelo gerado

```
Tensorflow.js ⓘ   Tensorflow ⓘ   Tensorflow Lite ⓘ  
  
<button type="button" onClick="init()">Start</button>  
<div id="webcam-container"></div>  
<div id="label-container"></div>  
<script src="https://cdn.jsdelivr.net/npm/@tensorflow/tfjs@1.3.1/dist/tf.min.js"></script>  
<script src="https://cdn.jsdelivr.net/npm/@teachablemachine/image@0.8/dist/teachablemachine-image.min.js"></script>  
<script type="text/javascript">  
  // More API functions here:  
  // https://github.com/googlecreativelab/teachablemachine-community/tree/master/libraries/image  
  
  // the link to your model provided by Teachable Machine export panel  
  const URL = "https://teachablemachine.withgoogle.com/models/q6FVCD1jo/";  
  
  let model, webcam, labelContainer, maxPredictions;  
  
  // Load the image model and setup the webcam  
  async function init() {  
    const modelURL = URL + "model.json";  
    const metadataURL = URL + "metadata.json";  
  
    // load the model and metadata  
    // Refer to tmImage.loadFromFiles() in the API to support files from a file picker  
    // or files from your local hard drive  
    // Note: the pose library adds "tmImage" object to your window (window.tmImage)  
    model = await tmImage.load(modelURL, metadataURL);  
    maxPredictions = model.getTotalClasses();  
  
    // Convenience function to setup a webcam  
    // https://github.com/googlecreativelab/teachablemachine-community/tree/master/libraries/webcam
```

<https://teachablemachine.withgoogle.com/models/q6FVCD1jo/>

modelo gerado



The screenshot shows the Teachable Machine interface. On the left, there are two classes: 'Class 1' with 4000 image samples and 'Class 2' with 4005 image samples. Each class has 'Webcam' and 'Upload' buttons. The main area is titled 'export your model to use it in projects.' and has three tabs: 'Tensorflow.js', 'Tensorflow', and 'Tensorflow Lite'. The 'Tensorflow' tab is selected. Under 'Model conversion type:', there are three options: 'Keras' (selected with a radio button), 'Savedmodel', and a 'Download my model' button. Below this, a note states: 'Converts your model to a keras .h5 model. Note the conversion happens in the cloud, but your training data is not being uploaded, only your trained model.' Under 'Code snippets to use your model:', there is a 'Keras' section with a 'Contribute on Github' link. A code editor shows the following Python code:

```
import tensorflow.keras
from PIL import Image, ImageOps
import numpy as np

# Disable scientific notation for clarity
np.set_printoptions(suppress=True)

# Load the model
model = tensorflow.keras.models.load_model('keras_model.h5')
```

A 'Copy' button is located to the right of the code editor.

[modelo - teachable machine](#)

[colab - teachable machine](#)

—

FIM

Teachable Machine

obrigado

prof.jefer@gmail.com



datasets

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fique de olho

[pytorch](#)

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[Caffe](#)

[Tensorflow - keras](#)

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[colab - intro deep learning - boston housing price - handwritten](#)

[colab - intro scikit - iris](#)

[Previendo aluguel de bicicleta](#)

[Previendo valor - aluguel RJ](#)

[Previendo valor passagem - NZ](#)

[Previendo passagens ORANGE](#)

orange no python

<https://colab.research.google.com/drive/1zbdbTeDWmBzpEA9VpDhWddW0IFZAUDYp?usp=sharing>

```
!pip install Orange3
import Orange

data = Orange.data.Table("housing")
learner = Orange.regression.linear.LinearRegressionLearner()
model = learner(data)

print ("pred obs")
for d in data[:3]:
    print ("%0.1f %0.1f" % (model(d), d.get_class()))
```

```
pred obs
30.0 24.0
25.0 21.6
30.6 34.7
```

```
!pip install Orange3
import Orange
from Orange.data import Table
from Orange.distance import Euclidean
iris = Table('iris')
dist_matrix = Euclidean(iris)
# Distance between first two examples
dist_matrix.X[0, 1]
```

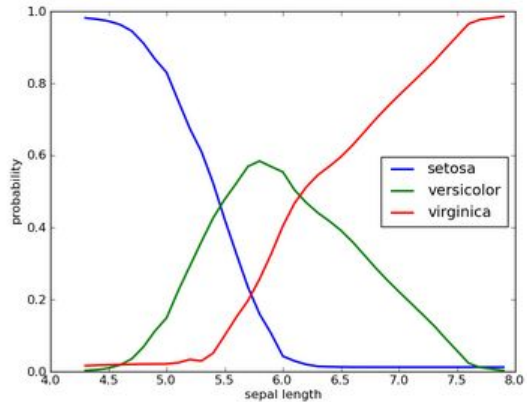
0.5385164807134628

```
iris = Orange.data.Table("iris.tab")
nb = Orange.classification.bayes.NaiveLearner(iris)

sepal_length, probabilities = zip(*nb.conditional_distributions[0].items())
p_setosa, p_versicolor, p_virginica = zip(*probabilities)

pylab.xlabel("sepal length")
pylab.ylabel("probability")
pylab.plot(sepal_length, p_setosa, label="setosa", linewidth=2)
pylab.plot(sepal_length, p_versicolor, label="versicolor", linewidth=2)
pylab.plot(sepal_length, p_virginica, label="virginica", linewidth=2)

pylab.legend(loc="best")
pylab.savefig("bayes-iris.png")
```



```
data = Orange.data.Table("housing.tab")
tree = Orange.regression.tree.TreeLearner(data, m_pruning=2., min_instances=20)
print tree.to_string()
```

The script outputs the tree:

```
RM<=6.941: 19.9
RM>6.941
|   RM<=7.437
|   |   CRIM>7.393: 14.4
|   |   CRIM<=7.393
|   |   |   DIS<=1.886: 45.7
|   |   |   DIS>1.886: 32.7
|   RM>7.437
|   |   TAX<=534.500: 45.9
|   |   TAX>534.500: 21.9
```

```
data = Orange.data.Table("housing.tab")
tree = Orange.regression.tree.TreeLearner(data, m_pruning=2., min_instances=20)
print tree.to_string()
```

The script outputs the tree:

```
RM<=6.941: 19.9
RM>6.941
|   RM<=7.437
|   |   CRIM>7.393: 14.4
|   |   CRIM<=7.393
|   |   |   DIS<=1.886: 45.7
|   |   |   DIS>1.886: 32.7
|   RM>7.437
|   |   TAX<=534.500: 45.9
|   |   TAX>534.500: 21.9
```

<https://docs.biolab.si/orange/2/tutorial/rst/regression.html>

```

>>> from Orange import data, distance
>>> from Orange.clustering import hierarchical
>>> data = data.Table('iris')
>>> dist_matrix = distance.Euclidean(data)
>>> hierar = hierarchical.HierarchicalClustering(n_clusters=3)
>>> hierar.linkage = hierarchical.AVERAGE
>>> hierar.fit(dist_matrix)
>>> hierar.labels
array([[ 1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.
         1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.
         1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.
         1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  1.,  0.
         0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.
         0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.
         0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  2.,  0.,  2.
         2.,  2.,  0.,  2.,  2.,  2.,  2.,  2.,  2.,  0.,  0.,  2.
         2.,  2.,  0.,  2.,  0.,  2.,  0.,  2.,  2.,  0.,  0.,  2.
         2.,  2.,  2.,  0.,  2.,  2.,  2.,  2.,  0.,  2.,  2.,  2.
         2.,  2.,  2.,  0.,  2.,  2.,  0.]])

```

Linear Regression

```
>>> from Orange.regression.linear import LinearRegressionLearner
>>> mpg = Orange.data.Table('auto-mpg')
>>> mean_ = LinearRegressionLearner()
>>> model = mean_(mpg[40:110])
>>> print(model)
LinearModel LinearRegression(copy_X=True, fit_intercept=True, normal:
>>> mpg[20]
Value('mpg', 25.0)
>>> model(mpg[0])
Value('mpg', 24.6)
```

cross validation

```
data = Orange.data.Table("titanic")
lr = Orange.classification.LogisticRegressionLearner()
res = Orange.evaluation.CrossValidation(data, [lr], k=5)
print("Accuracy: %.3f" % Orange.evaluation.scoring.CA(res)[0])
print("AUC:    %.3f" % Orange.evaluation.scoring.AUC(res)[0])
```

Accuracy: 0.779
AUC: 0.704

<https://orange3.readthedocs.io/projects/orange-data-mining-library/en/latest/tutorial/classification.html#learners-and-classifiers>

classification tree

```
>>> import Orange
>>> iris = Orange.data.Table('iris')
>>> tr = Orange.classification.TreeLearner()
>>> classifier = tr(iris)
>>> printed_tree = classifier.print_tree()
>>> for i in printed_tree.split('\n'):
>>>     print(i)
[50.  0.  0.] petal length ≤ 1.9
[ 0. 50. 50.] petal length > 1.9
[ 0. 49.  5.]     petal width ≤ 1.7
[ 0. 47.  1.]     petal length ≤ 4.9
  [0.  2.  4.]     petal length > 4.9
    [0.  0.  3.]     petal width ≤ 1.5
    [0.  2.  1.]     petal width > 1.5
      [0.  2.  0.]     sepal length ≤ 6.7
      [0.  0.  1.]     sepal length > 6.7
[ 0.  1. 45.]     petal width > 1.7
```

—

python no orange

```
import numpy as np
out_data = in_data.copy()
#copy, otherwise input data will be overwritten
np.round(out_data.X, 0, out_data.X)
```



The screenshot shows a window titled "Python Script" with a teal header. On the left, there is a sidebar with "Info" (Execute python script), "Input variables" (in_data, in_learner, in_classifier, in_object), "Output variables" (out_data, out_learner, out_classifier, out_object), and "Library" (round). The main area contains the Python code from the previous block. Below the code is a "Console" window showing the output of the script, which consists of several rows of numerical data in list format, such as [14.000, 6.000, 2.000, 20.000, 95.000, 2.000, 1.000, 1.000, 1.000, 8.000, 1.000, 2.000, 740.000 | 3], and so on. At the bottom, there are buttons for "+", "-", "Update", "More", and "Execute".

IA [údica

atividades

[Shy Panda](#)

[Ocean IA](#)
