

Supervised Learning: Support Vector Machines & Naive Bayes Classifier

AAA-Python Edition



Plan

- 1- Support Vector Machines
- 2- Kernel SVM Regressor
- 3- Face recognition with SVM
- 4- Naive Bayes Classification
- 5- Gaussian Naive Bayes
- 6- Multinomal Naive Bayes



Principles

- The support Vector Machine classification (SVM) is a classification where the mathematical model is the optimal hyperplane that delimits the classes of the data.
- The optimal hyperplane, is the one that maximizes the distance from each class.
- In a binary classification, with 2 features, the hyperplane is a **line** that maximizes the distance from the two classes.
- To identify the optimal line in this binary classification, a margin is drawn around each separating line up to the nearest point of each class.
- The optimal line, is the line that maximizes this margin.



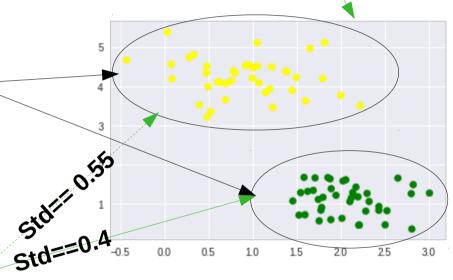
1- Support Vector Machines

Example

from sklearn.datasets.samples_generator import make_blobs
generating random data with 2 classes, standard deviation = 0.55, and random seed ==0
x, y = make_blobs(n_samples=80, centers=2, random_state=0,cluster_std=[0.55,0.4])
class 0 points are yellow, class 1 points are green
yc= ["green" if i else "yellow" for i in y]
plt.scatter(x[:, 0], x[:, 1],c=yc);

This function will generate:

- **80** points
- The points are centered around 2 clusters.
- the seed (a number used by the pseudo random generator. To have different data each time, you have to change the seed at each execution == a random one)== 0
- The standard deviation (indicates how spares is the data for each cluster)
- The coordinates of these points are in x: they represent the features values. The labels are in y



[By Amina Delali]



1- Support Vector Machines

importing support vector Example (suite) 2 from sklearn.svm import SVC 3 # creating an instance of an SVC classifier mvModel =SVC(kernel='linear') 5 # fiting the model to the data # make prediction about some values mvModel.fit(x,v) toPred=np.array([[1,1],[1,3],[3,5]]) # the optimal calculated support vectors vPred= mvModel.predict(toPred) vectXY=myModel.support vectors print("Predicted classes are:\n",yPred) # plotting the results ## plotting the decisions (classes) regions with the boundary line ## also hilighting the points with predicted classes Support vector from mlxtend.plotting import plot decision regions classifier with a plot_decision_regions(X=x,y=y,clf=myModel, legend=2,colors="yellow,green", markers="oo", X highlight=toPred) linear regressor ## plot the suport vector values plt.scatter(vectXY[:,0],vectXY[:,1],marker="+",c="red") Predicted classes are: [1 0 0] The separating line Support vectors: samples that touch the margins of the classifier

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gressor

Deep into SVM

```
# the parameters of the regressor
  = mvModel.coef [0]
b = mvModel.intercept [0]
  In general, the linear regressor model is as follow
# v = a1*x1 + a2 *x2 + b
# y represents the distances from the boundary line
# at the boudary line: y == 0
                                                           2
# => a1*x1+a2*x2+b=0 ==> a line's equation
\# => x2 = (-a1*x1 - b)/a2 = -a1/a2 * x1 - b/a2
# we will draw the boundary by initialing x1 values,
# and applying the previous formula, to calculate the
# x2 corresponding values defining the line
x1 = np.linspace(-1,4,1000)
                                                                     0
                                                                                             3
x2= (-a[0]*x1-b)/a[1]
plt.scatter(x[:, 0], x[:, 1],c=yc);
plt.plot(x1,x2, "r-", linewidth=2)
# plot the suport vector (the red "+" markers)
plt.scatter(vectXY[:,0],vectXY[:,1],marker="+",c="red")
# plot the margin lines
# the margin lines are paralles to the boundary line
# and their distance from the boundary line |v|== 1
# for the upper line: y==1, for the lower one: y==-1
#upper line equation: ==>a1*x1+a2*x2+b=1 ==> x2==(1-b-a1*x1)/a2
# lower line equation==> a1*x1+a2*x2+b=-1 ==>x2==( -1-b-a1*x1)/a2
# we will apply the previous formulas, to calculate the
# xmar1 and xmar2 corresponding values defining the margin lines
x2mar1 = (1 - a[0] * x1 - b)/a[1]
x2mar2 = (-1 - a[0] * x1 - b)/a[1]
plt.plot(x1,x2mar1, "b--", linewidth=1)
plt.plot(x1,x2mar2, "b--", linewidth=1)
```



2- Kernel SVM Regressor

Regression Example

```
# sklearn.datasets utilities doload data samples
# and to use data generators
from sklearn import datasets
# import a support vector machine regressor
from sklearn.svm import SVR
from sklearn.model selection import train_test_split
from sklearn.metrics import mean_squared_error
# shuffle wil
from sklearn.utils import shuffle
# load the data
myData = datasets.load_boston()
# the data is a dictionnary where : the x eatures values are designated by "data" key
# the y labels values are designated by "target" key
myData
```

```
'data': array([[6.3200e-03, 1.8000e+01, 2.3100e+00, ..., 1.5300e+01, 3.9690e+02, 4.9800e+00], [2.7310e-02, 0.0000e+00, 7.0700e+00, ..., 1.7800e+01, 3.9690e+02, 9.1400e+00], [2.7290e-02, 0.0000e+00, 7.0700e+00, ..., 1.7800e+01, 3.9283e+02, 4.0200e+001
```

	CRIM	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO	В	LSTAT
0	0.00632	18.0	2.31	0.0	0.538	6.575	65.2	4.09	1.0	296.0	15.3	396.9	4.98

{'DESCR': "Boston House Prices dataset\n=====================\n\nNotes\n-----\nData Set

[By Amina Delali]





Regression Example (suite)

```
x = myData.data
2 v = myData.target
                                                                            We can use
3 # spliting the data into training and testing sets
4 x_train, x_test, y_train, y_test = train_test_split(x, y,test_size=0.2
                                                                            different
5 # instantiate a svr with a linear regressor
6 myModel2 = SVR(kernel='linear')
                                                                            kernels for
7 # training phase
                                                                            regression ==>
8 myModel2.fit(x_train, y_train)
9 # testting the model
                                                                            Kernel SVM
10 vPred = myModel2.predict(x test)
11 # Evaluating the model
13 print( "mean squared error :", mean_squared_error(y_test,yPred))
                                                        mean squared error : 22.74219384815589
```

- In SVM, the model tries to maximizes the margin between the classe
- In an SVR, the model tries to fit as many as possible of samples (points) into that margin.

[By Amina Delali]



The data

- The statement "fetch Ifw people(min faces per person=10)" will import a dictionary with the following keys:
 - images: 3-D array with **4324** images. Each image, is described by a 2-D array of 62 rows, 47 columns of pixel values.
 - data: 2-D array with 4324 samples. Each sample, is described by a 1-D array of 62*47==2914 pixel values (reshape of "images").
 - target: labels of the images. Integer **code** indicating the name

of the person represented by a data row.

target names: names corresponding to codes.

```
1 # look closer to the images
                                                (2914,)
2 plt.imshow(faces.images[16],cmap='bone')
                                                              158
                                                (62, 47)
print(faces.data[16].shape)
 print(faces.images[16].shape)
                                                              people
                                                (158.)
5 print(faces.target[16])
                                                Catherine Zeta-Jones
print(faces.target names.shape)
 print(faces.target names[faces.target[16]])
print("images[0,0,\overline{0}]==", faces.images[0,0,0]," data[0,0]==", faces.data[0,0])
 images[0,0,0] == 95.333336 data[0,0] == 95.333336
                          Same values
```



Visualizing the data

Creating 3*5 == 15 subplots

```
ploting some images
import matplotlib.image as mpimg
fig, ax = plt.subplots(3, 5)
for i, axi in zip(range(10,25),ax.flat):
  axi.imshow(faces.images[i],cmap="bone")
  axi.set(xticks=[], yticks=[],xlabel=faces.target names[faces.target[i]][:11])
```

Returns an iterator over ax: in this case, it's like ax[k,l]

Displaying the images in each subplot

- Gloria Maca Dominique d









Lindsav Dav











Catherine Z





Gerhard Sch George W Bu George W Bu Winona Ryde

- Disabling the x and y ticks
- xlabel for each subplot corresponds to the person's name whose the face is displayed in that subplot
- we display only 11 characters for each name

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Training and Testing

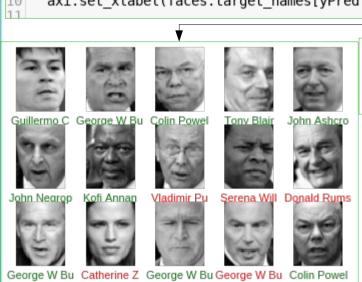
```
# predicting the values
yPred = myModel3.predict(x_test)
# defining the colors for correct and incorrect labels

CY = ["green" if j == i else "red" for i, j in zip(y_test,yPred)]

# ploting some test samples
fig, ax = plt.subplots(3, 5)
for i, axi in zip(range(88,103), ax.flat):
    axi.imshow(x_test[i].reshape(62, 47), cmap='gist_gray')
    axi.set(xticks=[], yticks=[])
axi.set xlabel(faces.target names[yPred[i]][:11],color=cY[i])
```

Displaying some test prediction: the **red** labels are **wrong** labels, the **green** ones are **good**.

fl-scoro



	precision	recure	11 30010	Support
Abdullah Gul	1.00	0.25	0.40	4
Adrien Brody	0.67	0.50	0.57	4

nrocision

recall.

avg / total 0.50 0.49 0.46 865

The worst results were related to "rbf" kernl. The "poly" kernel (default degree 3) scored piratically same as the "linear" one.

from sklearn.metrics import classification_report
the classification report
print(classification_report(y_test, yPred,target_names=faces.target_names))

[By Amina Delali]

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4- Naive Bayes Classification

Concept

- Naive Bayes models, are a group a of classification algorithms suitable for high dimensional datasets.
- They can handle multiple classes directly.
- The Naive Bayes classification methods rely on Bayes's theorem (Bayesian classification).
 - In **Bayesian classification**, we try to determine the **probability** of a **label** knowing the **features values**. We note: **P(L|Features)**
- **P(L|features)** is defined by as follow: $\frac{P(Features | L) \times P(L)}{P(Features)}$
- To decide between 2 labels, we calculate the ratio:

 $\frac{P(Features | L_1) \times P(L_1)}{P(Features | L_2) \times P(L_2)} _{12}$





Naive Bayes Classifiers

- So, we have to use a model capable of computing the: (features|L)
- This kind of models is called: generative model: it is able to generate data for each label.
- We have to define a generative model for each label.
- It is difficult to define a "general" model ==> so we make assumptions about the model ==> we define a "rough **approximation**" of the general model.
- Because of this simplification, we call such classifier: "Naive Bayes Classifier"
- The assumption in "Naive Bayes classifiers is that all features are independent given the value of the class variable.

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Different Naive Bayes Classifiers

Gaussian Naive Bayes

- Other assumption == the data for each label follows a "simple Guassian Distribution"
- ==> the model is simply defined by the mean and the standard deviation of the samples of each label

Multinomial Naive Bayes

- Other assumption == the data for each label follows a "simple Multinomial Distribution"
- ==> the model is simply defined by the probability of observing counts, among a number of categories





Gaussian Distribution in scikitlearn

$$P(x_i|y) = \frac{1}{\sqrt{2 \cdot \pi \cdot \sigma_y^2}} \cdot \exp\left(-\frac{(x_i - \mu_y)^2}{2 \cdot \sigma_y^2}\right)$$

 $\overline{\mu_y}$ \longrightarrow Is the features mean value, according to label y (labled y)

Is the standard deviation (its square is the variance) mean value, according to label y (labled y)

The two parameters are estimated using **maximum likelihood estimation**



5- Gaussian Naive Bayes

```
1 from sklearn import datasets
  2 iris = datasets.load_iris()
  3 print(iris.keys())
    print(iris.DESCR)
                      dict keys(['data', 'target', 'target names', 'DESCR', 'feature names'])
                Iris Plants Database
                Notes
classes
                Data Set Characteristics:
and 4
                     :Number of Instances: 150 (50 in each of three classes)
                     :Number of Attributes: 4 numeric, predictive attributes and the class
features
                     :Attribute Information:
                         - sepal length in cm
                         - sepal width in cm
                         - petal length in cm
                         - petal width in cm
                         - class:
                                 - Iris-Setosa
                                 - Iris-Versicolour
                                 - Iris-Virginica
```

Data

[By Amina Delali]



Training and Testing

```
from sklearn.model_selection import train_test_split

#import the gaussian naive bayes model
from sklearn.naive_bayes import GaussianNB
myModel4 = GaussianNB()
# spliting the data
7 x_train, x_test, y_train, y_test =train_test_split(iris.data, iris.target,test_size=0.2)
# training the model
myModel4.fit(x_train, y_train)

# testing the model
yPred= myModel4.predict(x_test)

from sklearn.metrics import classification_report

# print the classification_report(y_test,yPred,target_names=iris.target_names))
```

precision recall f1-score support setosa 1.00 1.00 1.00 11 versicolor 0.91 1.00 0.95 10 virginica 1.00 0.89 0.94 9 avg / total 0.97 0.97 0.97 30						
versicolor 0.91 1.00 0.95 10 virginica 1.00 0.89 0.94 9		precision	recall	f1-score	support	
virginica 1.00 0.89 0.94 9						
avg / total 0.97 0.97 0.97 30						
	avg / total	0.97	0.97	0.97	30	





Deep into Multinomial Distribution

- Used in text classification, where the data is represented as vectors of words count.
- It is parametrized by vectors: $\theta_y = (\theta_{y1}, ..., \theta_{yn})$ for each class y. n is the number of features (size of the vocabulary in a text classification).
 - θ_i is P(xi|y) == the probability that the feature i appears in a sample belonging to class y.
 - $heta_{
 m yi}$ is computed as follow: $\hat{ heta}_{yi} = rac{N_{yi} + lpha}{N_y + lpha n}$

 $N_{yi} = \sum_{x \in T} x_i$: number times that the feature i appears in a sample belonging to class y in the training set T.

- $N_y = \sum_{i=1}^n N_{yi}$: is the total count for all features appearing in class y
- $\alpha \ge 0$: a smoothing parameter, allows to avoid 0 probability.



Bayes 6- Mul Naive

Data

```
'data': <11314x130107 sparse matrix of type '<class 'numpy.float64'>'
      with 1787565 stored elements in Compressed Sparse Row format>,
'target': array([17, 7, 10, ..., 14, 12, 11]),
                                                                print(myTextD.data.shape)
'target names': ['alt.atheism',
 'comp.graphics',
                                                              (11314, 130107)
 'comp.os.ms-windows.misc',
 'comp.sys.ibm.pc.hardware',
 'comp.sys.mac.hardware',
                                If we used: fetch_20newsgroups, we will have to
 'comp.windows.x',
                                vectorize the data, using for example:
 'misc.forsale',
                                from sklearn.feature extraction.text import
 'rec.autos',
                                TfidfVectorizer
'rec.motorcvcles',
 'rec.sport.baseball',
 'rec.sport.hockey',
 'sci.crypt',
                                 from sklearn.datasets import fetch_20newsgroups_vectorized
 'sci.electronics'.
                                 myTextD = fetch 20newsgroups vectorized()
'sci.med',
                                 myTextD.target names
 'sci.space',
 'soc.religion.christian',
                                 myTextD
 'talk.politics.guns',
 'talk.politics.mideast',
 'talk.politics.misc'.
 'talk.religion.misc']}
```





Training and testing

```
1 #import the multinomial naive bayes model
 2 from sklearn.naive bayes import MultinomialNB
 3 myModel5 = Multinomia(NB()
 4 # spliting the data
 5 # spliting the data
 6 x train, x test, y train, y test =train test split(myTextD.data, myTextD.target,test size=0.2)
 7 # training the model
 8 myModel5.fit(x train, y train)
10 # testing the model
11 vPred= mvModel5.predict(x test)
13 from sklearn.metrics import classification_report
15 # print the classification report
16 print (classification report(y test, yPred, target names=myTextD.target names))
```

			precision	recall	f1-score	support	
	alt.atheism		0.89	0.40	0.55	101	
	comp.graphics			0.54	0.66	127	
	comp.os.ms-windows.misc			0.74	0.77	117	
	comp.	sys.ibm.pc.hardware	0.58	0.81	0.68	113	
	co	mp.sys.mac.hardware	0.95	0.55	0.70	137	
	comp.windows.x		0.87	0.81	0.84	113	
By Amina D)elali1	avg / total	0.80	0.73	0.71	2263	Π

[By



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Thank you!

FOR ALL YOUR TIME