

# APA-L5-python

September 7, 2018

## 1 APA Laboratori 5 - LDA/QDA/NBayes/RegLog

```
In [1]: # Uncomment to upgrade packages
        # !pip install pandas --upgrade
        # !pip install numpy --upgrade
        # !pip install scipy --upgrade
        # !pip install statsmodels --upgrade
        # !pip install scikit-learn --upgrade
        %load_ext autoreload

In [2]: %%matplotlib notebook
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sn
import pandas as pd
pd.set_option('precision', 3)
from IPython.core.interactiveshell import InteractiveShell
InteractiveShell.ast_node_interactivity = "all"

In [3]: # Extra imports
from pandas import read_csv
from sklearn.metrics import confusion_matrix, \
    classification_report, accuracy_score
from pandas.api.types import CategoricalDtype
from pandas.plotting import scatter_matrix
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
from sklearn.model_selection import LeaveOneOut
from sklearn.discriminant_analysis import QuadraticDiscriminantAnalysis
from sklearn.preprocessing import Imputer
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import BernoulliNB
from sklearn.datasets import load_iris
from sklearn.neighbors import KNeighborsClassifier
from numpy.random import normal, binomial
from statsmodels.genmod.generalized_linear_model import GLM
from statsmodels.genmod.families.family import Binomial
from statsmodels.tools.tools import add_constant
```

```

from sklearn.feature_selection import RFECV
from sklearn.linear_model import LogisticRegression

```

```

In [4]: def confusion(true, pred, classes):
        """
        Function for pretty printing confusion matrices
        """
        cm = pd.DataFrame(confusion_matrix(true, pred),
                           index=classes,
                           columns=classes)
        cm.index.name = 'Actual'
        cm.columns.name = 'Predicted'
        return cm

```

## 1.1 Example 1: Visualizing and classifying wines with LDA and QDA

We have the results of an analysis on wines grown in a region in Italy but derived from three different cultivars. The analysis determined the quantities of 13 chemical constituents found in each of the three types of wines. The goal is to separate the three types of wines:

```

In [5]: wine = read_csv("wine.data", delimiter=',', header=None)
        wine_classes = ['cultivar %d'%(i+1) for i in range(3)]
        wine.shape
        wine.columns = ['Wine_type', 'Alcohol', 'Malic_acid', 'Ash',
                        'Alcalinity_of_ash', 'Magnesium', 'Total_phenols',
                        'Flavanoids', 'Nonflavanoid_phenols',
                        'Proanthocyanins', 'Color_intensity', 'Hue',
                        'OD280/OD315', 'Proline']

```

```
Out[5]: (178, 14)
```

```

In [6]: wine.Wine_type = wine.Wine_type.astype(CategoricalDtype(categories=[1, 2, 3],
                                                                    ordered=True))
        wine.describe(include='all')

```

```

Out[6]:

```

	Wine_type	Alcohol	Malic_acid	Ash	Alcalinity_of_ash	Magnesium \
count	178.0	178.000	178.000	178.000	178.000	178.000
unique	3.0	NaN	NaN	NaN	NaN	NaN
top	2.0	NaN	NaN	NaN	NaN	NaN
freq	71.0	NaN	NaN	NaN	NaN	NaN
mean	NaN	13.001	2.336	2.367	19.495	99.742
std	NaN	0.812	1.117	0.274	3.340	14.282
min	NaN	11.030	0.740	1.360	10.600	70.000
25%	NaN	12.362	1.603	2.210	17.200	88.000
50%	NaN	13.050	1.865	2.360	19.500	98.000
75%	NaN	13.678	3.083	2.558	21.500	107.000
max	NaN	14.830	5.800	3.230	30.000	162.000

```

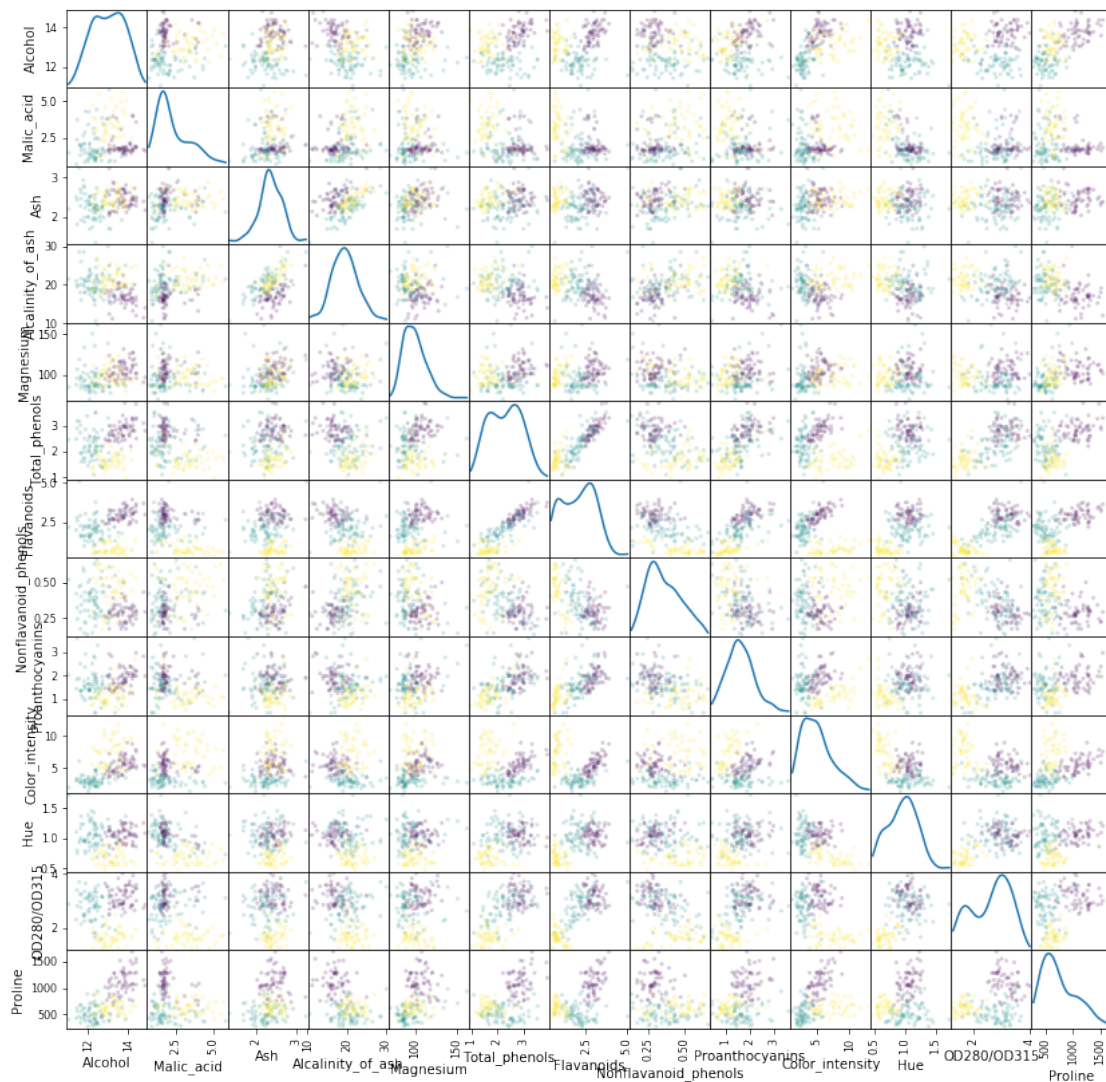
        Total_phenols  Flavanoids  Nonflavanoid_phenols  Proanthocyanins \

```

count	178.000	178.000	178.000	178.000
unique	NaN	NaN	NaN	NaN
top	NaN	NaN	NaN	NaN
freq	NaN	NaN	NaN	NaN
mean	2.295	2.029	0.362	1.591
std	0.626	0.999	0.124	0.572
min	0.980	0.340	0.130	0.410
25%	1.742	1.205	0.270	1.250
50%	2.355	2.135	0.340	1.555
75%	2.800	2.875	0.438	1.950
max	3.880	5.080	0.660	3.580

	Color_intensity	Hue	OD280/OD315	Proline
count	178.000	178.000	178.000	178.000
unique	NaN	NaN	NaN	NaN
top	NaN	NaN	NaN	NaN
freq	NaN	NaN	NaN	NaN
mean	5.058	0.957	2.612	746.893
std	2.318	0.229	0.710	314.907
min	1.280	0.480	1.270	278.000
25%	3.220	0.782	1.938	500.500
50%	4.690	0.965	2.780	673.500
75%	6.200	1.120	3.170	985.000
max	13.000	1.710	4.000	1680.000

```
In [7]: scatter_matrix(wine.loc[:, 'Alcohol': 'Proline'],
                        alpha=0.2, figsize=(14, 14),
                        diagonal='kde', marker='.',
                        c=wine.Wine_type);
```



For this example let's practice a different call mode to `lda()`, using a formula; this is most useful when our data is in a dataframe format:

```
In [8]: lda_model = LinearDiscriminantAnalysis().fit(wine.loc[:, 'Alcohol': 'Proline'],
                                                    wine.Wine_type)

print('Priors:', lda_model.priors_)
print('Means:\n')
means = pd.DataFrame(lda_model.means_)
means.columns = wine.columns[1:]
means
print('Coefs:')
coefs = pd.DataFrame(lda_model.scalings_)
coefs.index = wine.columns[1:]
coefs
```

```
print('Explained Variance Ratio')
pd.DataFrame(lda_model.explained_variance_ratio_ )
```

Priors: [0.33146067 0.3988764 0.26966292]

Means:

```
Out[8]:
```

	Alcohol	Malic_acid	Ash	Alcalinity_of_ash	Magnesium	Total_phenols	\
0	13.745	2.011	2.456	17.037	106.339	2.840	
1	12.279	1.933	2.245	20.238	94.549	2.259	
2	13.154	3.334	2.437	21.417	99.312	1.679	

	Flavanoids	Nonflavanoid_phenols	Proanthocyanins	Color_intensity	Hue	\
0	2.982		0.290	1.899	5.528	1.062
1	2.081		0.364	1.630	3.087	1.056
2	0.781		0.447	1.154	7.396	0.683

	OD280/OD315	Proline
0	3.158	1115.712
1	2.785	519.507
2	1.684	629.896

Coefs:

```
Out[8]:
```

	0	1
Alcohol	-0.403	8.718e-01
Malic_acid	0.165	3.054e-01
Ash	-0.369	2.346e+00
Alcalinity_of_ash	0.155	-1.464e-01
Magnesium	-0.002	-4.628e-04
Total_phenols	0.618	-3.221e-02
Flavanoids	-1.661	-4.920e-01
Nonflavanoid_phenols	-1.496	-1.631e+00
Proanthocyanins	0.134	-3.071e-01
Color_intensity	0.355	2.532e-01
Hue	-0.818	-1.516e+00
OD280/OD315	-1.158	5.118e-02
Proline	-0.003	2.853e-03

Explained Variance Ratio

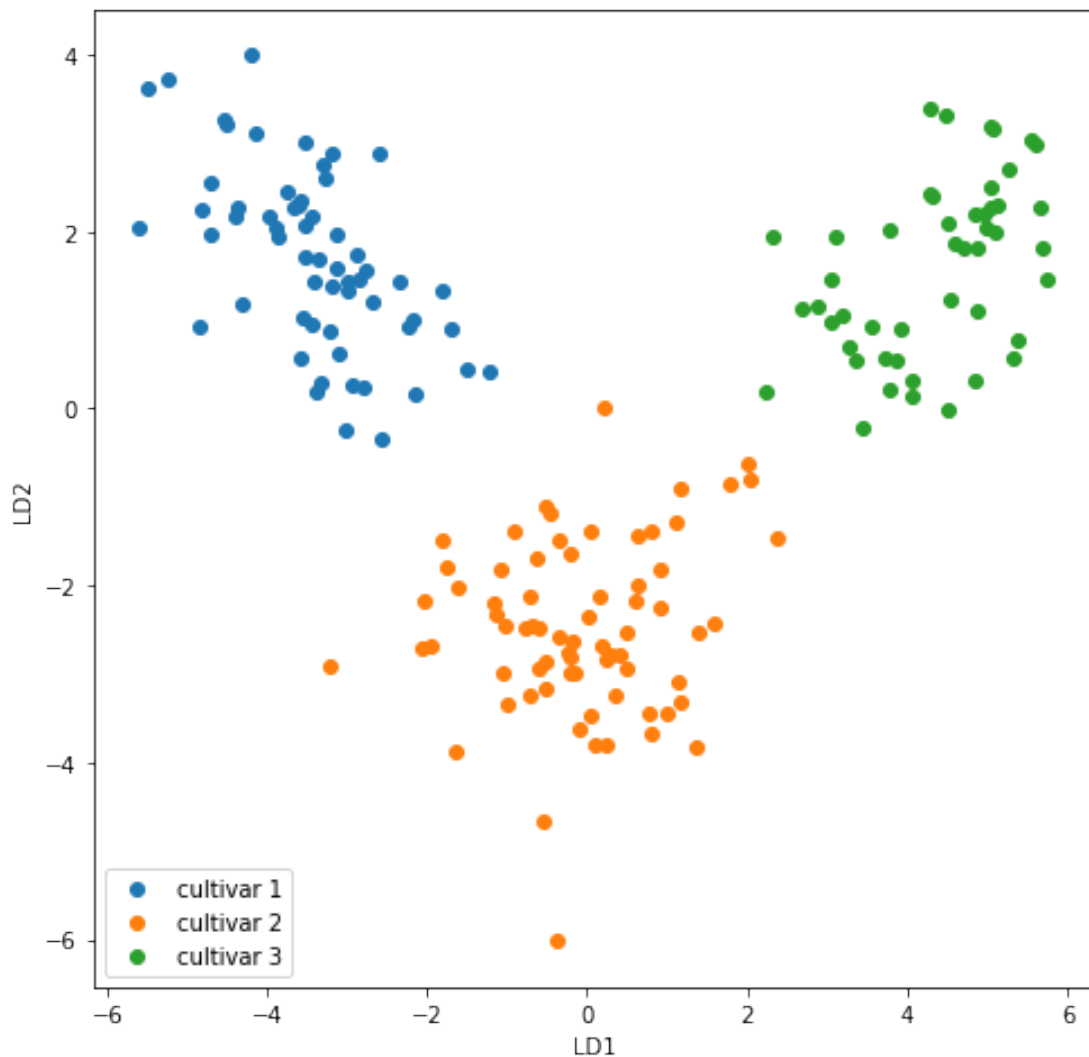
```
Out[8]:
```

	0
0	0.687
1	0.313

We can see that neither Magnesium or Proline seem useful to separate the wines; while Flavanoids and Nonflavanoid.phenols do. Ash is mainly used in the LD2.

Plot the projected data in the first two LDs  
We can see that the discrimination is very good

```
In [9]: wine_trans = lda_model.transform(wine.loc[:, 'Alcohol': 'Proline'])
fig, ax = plt.subplots(figsize=(8,8))
for i in wine.Wine_type.unique():
    plt.scatter(wine_trans[:,0][wine.Wine_type==i],
                wine_trans[:,1][wine.Wine_type==i],
                label='cultivar %d'%i)
ax.set_xlabel('LD1')
ax.set_ylabel('LD2')
plt.legend();
```

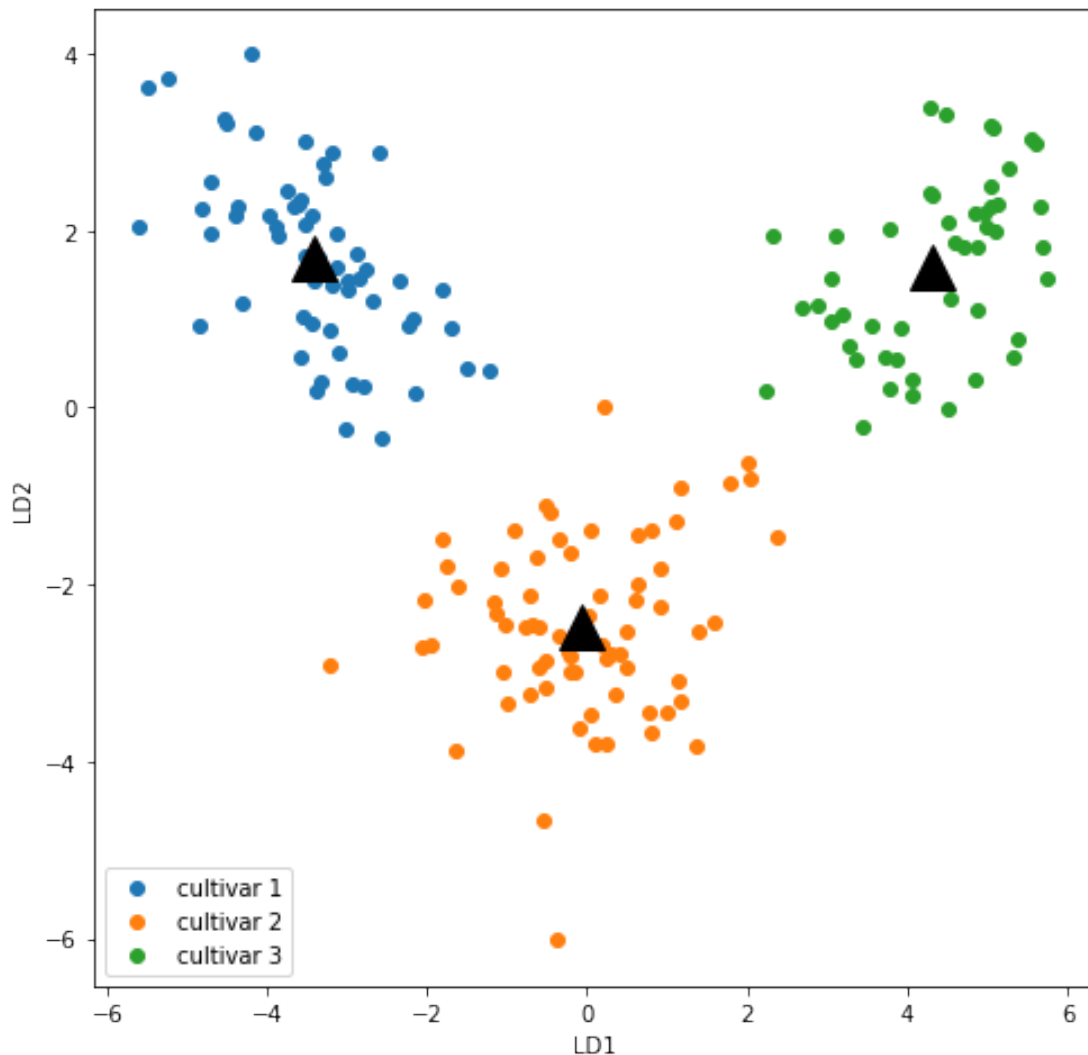


If need be, we can add the (projected) means to the plot

```

In [10]: fig, ax = plt.subplots(figsize=(8,8))
         for i in wine.Wine_type.unique():
             plt.scatter(wine_trans[:,0][wine.Wine_type==i],
                         wine_trans[:,1][wine.Wine_type==i],
                         label='cultivar %d'%i)
             plt.plot(wine_trans[:,0][wine.Wine_type==i].mean(),
                     wine_trans[:,1][wine.Wine_type==i].mean(),
                     'k^',markersize=20)
         ax.set_xlabel('LD1')
         ax.set_ylabel('LD2')
         plt.legend();

```



indeed classification is perfect

```

In [11]: confusion(wine.Wine_type, lda_model.predict(wine.loc[:, 'Alcohol': 'Proline']),
                  classes=wine_classes)

```

```
Out[11]: Predicted   cultivar 1   cultivar 2   cultivar 3
Actual
cultivar 1         59           0           0
cultivar 2          0          71           0
cultivar 3          0           0          48
```

Let us switch to leave-one-out cross-validation

```
In [12]: def loocv(X,y,model,classes):
        loo = LeaveOneOut()
        pred=[]
        for train_index, test_index in loo.split(X):
            X_tr, X_ts = X[train_index], X[test_index]
            y_tr, _ = y[train_index], y[test_index]
            model.fit(X_tr,y_tr)
            pred.append(model.predict(X_ts)[0])
        return confusion(y,pred,classes), 1-accuracy_score(y,pred)
```

```
In [13]: cm, err = loocv(wine.loc[:, 'Alcohol': 'Proline'].values,
                        wine.Wine_type,
                        LinearDiscriminantAnalysis(),
                        wine_classes)
```

```
cm
```

```
err*100
```

```
Out[13]: Predicted   cultivar 1   cultivar 2   cultivar 3
Actual
cultivar 1         59           0           0
cultivar 2          1          69           1
cultivar 3          0           0          48
```

```
Out[13]: 1.1235955056179803
```

2 mistakes (on 178 observations): 1.12% error  
 Quadratic Discriminant Analysis is the same  
 problems may arise if for some class there are less (or equal) observations than dimensions (is not the case for the wine data)

```
In [14]: qda_model = QuadraticDiscriminantAnalysis().fit(wine.loc[:, 'Alcohol': 'Proline'],
                                                         wine.Wine_type)
```

```
print('Priors:\n')
pd.DataFrame(qda_model.priors_)
print('Means:\n')
means =pd.DataFrame(qda_model.means_)
means.columns=wine.columns[1:]
means
```



Priors:

```
Out[14]:      0
0  0.331
1  0.399
2  0.270
```

Means:

```
Out[14]:  Alcohol  Malic_acid  Ash  Alcalinity_of_ash  Magnesium  Total_phenols  \
0   13.745      2.011  2.456          17.037    106.339      2.840
1   12.279      1.933  2.245          20.238     94.549      2.259
2   13.154      3.334  2.437          21.417    99.312      1.679

      Flavanoids  Nonflavanoid_phenols  Proanthocyanins  Color_intensity  Hue  \
0         2.982             0.290          1.899          5.528  1.062
1         2.081             0.364          1.630          3.087  1.056
2         0.781             0.447          1.154          7.396  0.683

      OD280/OD315  Proline
0         3.158  1115.712
1         2.785   519.507
2         1.684   629.896
```

There is no projection this time (because projection is a linear operator and the QDA boundaries are quadratic ones)

but let's have a look at classification:

```
In [15]: confusion(wine.Wine_type, qda_model.predict(wine.loc[:, 'Alcohol': 'Proline']),
                  classes=wine_classes)
```

```
Out[15]: Predicted   cultivar 1   cultivar 2   cultivar 3
Actual
cultivar 1          59           0           0
cultivar 2           1          70           0
cultivar 3           0           0          48
```

Let us switch to leave-one-out cross-validation

```
In [16]: cm, err = loocv(wine.loc[:, 'Alcohol': 'Proline'].values,
                        wine.Wine_type,
                        QuadraticDiscriminantAnalysis(),
                        wine_classes)
```

cm

err\*100

```
Out[16]: Predicted   cultivar 1   cultivar 2   cultivar 3
Actual
cultivar 1         59           0           0
cultivar 2          1          70           0
cultivar 3          0           0          48
```

```
Out[16]: 0.5617977528089901
```

1 mistake (on 178 observations): 0.56% error

it would be nice to ascertain which wine is the "stubborn" one: it is a wine of type '2' classified as class '1'. Maybe there is something special with this wine ...

In the event of numerical errors (insufficient number of observations per class), we can use regularization.

in this case the regularization parameter (0.1) is applied to the covariance matrix (Sigma) so it is not ill conditioned in this fashion

```
(1-reg_param)*Sigma + reg_param*np.eye(n_features)
```

```
In [17]: qda_model = QuadraticDiscriminantAnalysis(reg_param=0.1).\
          fit(wine.loc[:, 'Alcohol': 'Proline'],
              wine.Wine_type)
```

```
print('Priors:', qda_model.priors_)
print('Means:\n')
means = pd.DataFrame(qda_model.means_)
means.columns=wine.columns[1:]
means
```

```
Priors: [0.33146067 0.3988764  0.26966292]
```

```
Means:
```

```
Out[17]:
```

	Alcohol	Malic_acid	Ash	Alcalinity_of_ash	Magnesium	Total_phenols	\
0	13.745	2.011	2.456	17.037	106.339	2.840	
1	12.279	1.933	2.245	20.238	94.549	2.259	
2	13.154	3.334	2.437	21.417	99.312	1.679	

	Flavanoids	Nonflavanoid_phenols	Proanthocyanins	Color_intensity	Hue	\
0	2.982		0.290	1.899	5.528	1.062
1	2.081		0.364	1.630	3.087	1.056
2	0.781		0.447	1.154	7.396	0.683

	OD280/OD315	Proline
0	3.158	1115.712
1	2.785	519.507
2	1.684	629.896

```
In [18]: confusion(wine.Wine_type, qda_model.predict(wine.loc[:, 'Alcohol': 'Proline']),
                  classes=wine_classes)
```

```
Out[18]: Predicted   cultivar 1   cultivar 2   cultivar 3
Actual
cultivar 1         59           0           0
cultivar 2          0          69           2
cultivar 3          0           0          48
```

---

## 1.2 Example 2: The Naïve Bayes classifier

Naive Bayes Classifier for Discrete Predictors: we use the 1984 United States Congressional Voting Records;

This data set includes votes for each of the U.S. House of Representatives Congressmen on 16 key votes In origin they were nine different types of votes:

- voted for, paired for, and announced for (these three simplified to yea or 'y'),
- voted against, paired against, and announced against (these three simplified to nay or 'n'),
- voted present, voted present to avoid conflict of interest, and did not vote or otherwise make a position known (these three simplified to an 'unknown' disposition)

The goal is to classify Congressmen as Republican or Democrat as a function of their voting profiles, which is not immediate because in the US Congressmen have a large freedom of vote (obviously linked to their party but also to their own feelings, interests and compromises with voters)

```
In [19]: HouseVotes84 = read_csv("house-votes-84.data",
                                delimiter=',',
                                header=None,na_values='?')

house_classes = ['n','y']
```

add meaningful names to the votes

```
In [20]: HouseVotes84.columns=["Class","handicapped_infants","water_project_sharing",
                                "budget_resolution","physician_fee_freeze",
                                "el_salvador_aid","religious_groups_in_schools",
                                "anti_satellite_ban", "aid_to_nicaraguan_contras",
                                "mx_missile","immigration","synfuels_cutback",
                                "education_spending","superfund","crime",
                                "duty_free_exports","export_South_Africa"]

HouseVotes84.describe()
```

```
Out[20]:
```

	Class	handicapped_infants	water_project_sharing	budget_resolution	\
count	435	423	387	424	
unique	2	2	2	2	
top	democrat	n	y	y	
freq	267	236	195	253	

	physician_fee_freeze	el_salvador_aid	religious_groups_in_schools	\
count	424	420	424	

unique	2	2	2
top	n	y	y
freq	247	212	272

	anti_satellite_ban	aid_to_nicaraguan_contras	mx_missile	immigration	\
count	421	420	413	428	
unique	2	2	2	2	
top	y	y	y	y	
freq	239	242	207	216	

	synfuels_cutback	education_spending	superfund	crime	duty_free_exports	\
count	414	404	410	418	407	
unique	2	2	2	2	2	
top	n	n	y	y	n	
freq	264	233	209	248	233	

	export_South_Africa
count	331
unique	2
top	y
freq	269

```
In [21]: for v in HouseVotes84.columns:
          HouseVotes84[v].value_counts(dropna=False)
```

```
Out[21]: democrat      267
republican      168
Name: Class, dtype: int64
```

```
Out[21]: n      236
y      187
NaN      12
Name: handicapped_infants, dtype: int64
```

```
Out[21]: y      195
n      192
NaN      48
Name: water_project_sharing, dtype: int64
```

```
Out[21]: y      253
n      171
NaN      11
Name: budget_resolution, dtype: int64
```

```
Out[21]: n      247
y      177
NaN      11
Name: physician_fee_freeze, dtype: int64
```

```

Out[21]: y      212
         n      208
         NaN     15
         Name: el_salvador_aid, dtype: int64

Out[21]: y      272
         n      152
         NaN     11
         Name: religious_groups_in_schools, dtype: int64

Out[21]: y      239
         n      182
         NaN     14
         Name: anti_satellite_ban, dtype: int64

Out[21]: y      242
         n      178
         NaN     15
         Name: aid_to_nicaraguan_contras, dtype: int64

Out[21]: y      207
         n      206
         NaN     22
         Name: mx_missile, dtype: int64

Out[21]: y      216
         n      212
         NaN      7
         Name: immigration, dtype: int64

Out[21]: n      264
         y      150
         NaN     21
         Name: synfuels_cutback, dtype: int64

Out[21]: n      233
         y      171
         NaN     31
         Name: education_spending, dtype: int64

Out[21]: y      209
         n      201
         NaN     25
         Name: superfund, dtype: int64

Out[21]: y      248
         n      170
         NaN     17
         Name: crime, dtype: int64

```

```
Out[21]: n      233
        y      174
        NaN     28
        Name: duty_free_exports, dtype: int64
```

```
Out[21]: y      269
        NaN    104
        n       62
        Name: export_South_Africa, dtype: int64
```

1 = democrat, 0 = republican Note "unknown dispositions" have been treated as missing values!

The naive bayes implementations of scikit-learn do not allow missing values and also need binary data, so we will preprocess first changing *y* for 1 and *n* for 0 and then we perform missing data imputation. Another option would be to eliminate all rows with missing, but that will discard half of the data

```
In [22]: HouseVotes84.replace({'y':1, 'n':0},inplace=True)
        HouseVotes84.head()
```

```
Out[22]:
```

	Class	handicapped_infants	water_project_sharing	budget_resolution	\
0	republican	0.0	1.0	0.0	
1	republican	0.0	1.0	0.0	
2	democrat	NaN	1.0	1.0	
3	democrat	0.0	1.0	1.0	
4	democrat	1.0	1.0	1.0	

	physician_fee_freeze	el_salvador_aid	religious_groups_in_schools	\
0	1.0	1.0	1.0	
1	1.0	1.0	1.0	
2	NaN	1.0	1.0	
3	0.0	NaN	1.0	
4	0.0	1.0	1.0	

	anti_satellite_ban	aid_to_nicaraguan_contras	mx_missile	immigration	\
0	0.0	0.0	0.0	1.0	
1	0.0	0.0	0.0	0.0	
2	0.0	0.0	0.0	0.0	
3	0.0	0.0	0.0	0.0	
4	0.0	0.0	0.0	0.0	

	synfuels_cutback	education_spending	superfund	crime	duty_free_exports	\
0	NaN	1.0	1.0	1.0	0.0	
1	0.0	1.0	1.0	1.0	0.0	
2	1.0	0.0	1.0	1.0	0.0	
3	1.0	0.0	1.0	0.0	0.0	
4	1.0	NaN	1.0	1.0	1.0	

export\_South\_Africa

0	1.0
1	NaN
2	0.0
3	1.0
4	1.0

We use the most frequent value from each column for imputation

```
In [23]: HouseVotes84.loc[:, 'handicapped_infants'] = Imputer(strategy='most_frequent').\
fit_transform(HouseVotes84.loc[:, 'handicapped_infants'])
HouseVotes84.head()
```

```
Out [23]:
```

	Class	handicapped_infants	water_project_sharing	budget_resolution	\
0	republican	0.0	1.0	0.0	
1	republican	0.0	1.0	0.0	
2	democrat	0.0	1.0	1.0	
3	democrat	0.0	1.0	1.0	
4	democrat	1.0	1.0	1.0	

	physician_fee_freeze	el_salvador_aid	religious_groups_in_schools	\
0	1.0	1.0	1.0	
1	1.0	1.0	1.0	
2	0.0	1.0	1.0	
3	0.0	1.0	1.0	
4	0.0	1.0	1.0	

	anti_satellite_ban	aid_to_nicaraguan_contras	mx_missile	immigration	\
0	0.0	0.0	0.0	1.0	
1	0.0	0.0	0.0	0.0	
2	0.0	0.0	0.0	0.0	
3	0.0	0.0	0.0	0.0	
4	0.0	0.0	0.0	0.0	

	synfuels_cutback	education_spending	superfund	crime	duty_free_exports	\
0	0.0	1.0	1.0	1.0	0.0	
1	0.0	1.0	1.0	1.0	0.0	
2	1.0	0.0	1.0	1.0	0.0	
3	1.0	0.0	1.0	0.0	0.0	
4	1.0	0.0	1.0	1.0	1.0	

	export_South_Africa
0	1.0
1	1.0
2	0.0
3	1.0
4	1.0

```
In [24]: np.random.seed(1111)
N = HouseVotes84.shape[0]
```

We first split the available data into learning and test sets, selecting randomly 2/3 and 1/3 of the data.

We do this for a honest estimation of prediction performance

```
In [25]: train, test = train_test_split(HouseVotes84, test_size=N//3)
```

We use the BernoulliNB estimator because we have binary data

```
In [26]: model = BernoulliNB().fit(train.loc[:, 'handicapped_infants'],
                                     train.Class)
```

To obtain the probabilities from the model is a little bit tricky.

The attribute `class_log_prior_` stores the prior log probabilities for the classes, so we can compute the probabilities doing:

```
In [27]: np.e**model.class_log_prior_
```

```
Out [27]: array([0.62068966, 0.37931034])
```

For the attributes/class probabilities is trickier because only one of the probabilities is stored (the other is the complement) and also are the log probabilities

```
In [28]: probs=pd.DataFrame({'Democrat Y':np.e**model.feature_log_prob_.T[:,0],
                             'Democrat N':1-np.e**model.feature_log_prob_.T[:,0],
                             'Republican Y':np.e**model.feature_log_prob_.T[:,1],
                             'Republican N':1-np.e**model.feature_log_prob_.T[:,1]},
                             index=HouseVotes84.columns[1:])
```

probs

```
Out [28]:
```

	Democrat Y	Democrat N	Republican Y \
handicapped_infants	0.533	0.467	0.205
water_project_sharing	0.549	0.451	0.518
budget_resolution	0.896	0.104	0.196
physician_fee_freeze	0.055	0.945	0.955
el_salvador_aid	0.236	0.764	0.929
religious_groups_in_schools	0.511	0.489	0.875
anti_satellite_ban	0.775	0.225	0.313
aid_to_nicaraguan_contras	0.835	0.165	0.259
mx_missile	0.797	0.203	0.170
immigration	0.505	0.495	0.545
synfuels_cutback	0.473	0.527	0.116
education_spending	0.143	0.857	0.750
superfund	0.352	0.648	0.839
crime	0.363	0.637	0.982
duty_free_exports	0.610	0.390	0.125
export_South_Africa	0.962	0.038	0.679

Republican N



handicapped_infants	0.795
water_project_sharing	0.482
budget_resolution	0.804
physician_fee_freeze	0.045
el_salvador_aid	0.071
religious_groups_in_schools	0.125
anti_satellite_ban	0.687
aid_to_nicaraguan_contras	0.741
mx_missile	0.830
immigration	0.455
synfuels_cutback	0.884
education_spending	0.250
superfund	0.161
crime	0.018
duty_free_exports	0.875
export_South_Africa	0.321

predict the outcome of the first 20 Congressmen

```
In [29]: model.predict(HouseVotes84.loc[0:20, 'handicapped_infants':])
```

```
Out[29]: array(['republican', 'republican', 'republican', 'democrat', 'democrat',
                'republican', 'republican', 'republican', 'republican', 'democrat',
                'republican', 'republican', 'democrat', 'democrat', 'republican',
                'republican', 'democrat', 'democrat', 'republican', 'democrat',
                'democrat'], dtype='<U10')
```

```
In [30]: pred=pd.DataFrame(model.predict_proba(HouseVotes84.loc[0:20, 'handicapped_infants':]))
```

```
pred.columns=['democrat', 'republican']
pred
```

```
Out[30]:
```

	democrat	republican
0	2.084e-07	1.000e+00
1	2.439e-07	1.000e+00
2	3.163e-02	9.684e-01
3	9.740e-01	2.604e-02
4	9.492e-01	5.077e-02
5	3.829e-01	6.171e-01
6	4.804e-05	1.000e+00
7	4.390e-06	1.000e+00
8	2.439e-07	1.000e+00
9	1.000e+00	7.242e-09
10	2.483e-06	1.000e+00
11	2.995e-05	1.000e+00
12	1.000e+00	3.079e-07
13	1.000e+00	7.603e-10
14	2.439e-07	1.000e+00
15	2.084e-07	1.000e+00

```

16  1.000e+00  1.239e-05
17  1.000e+00  6.372e-09
18  3.707e-07  1.000e+00
19  1.000e+00  9.700e-11
20  1.000e+00  1.246e-08

```

form and display confusion matrix & overall error

```
In [31]: confusion(train.Class, model.predict(train.loc[:, 'handicapped_infants:']),
               classes=house_classes)
```

```

(1-accuracy_score(train.Class,
                   model.predict(train.loc[:, 'handicapped_infants:']))) * 100

```

```
Out[31]: Predicted   n   y
Actual
n           159   21
y            9  101
```

```
Out[31]: 10.344827586206895
```

compute the test (prediction) error

```
In [32]: confusion(test.Class,
                   model.predict(test.loc[:, 'handicapped_infants:']),
                   classes=house_classes)
```

```

(1-accuracy_score(test.Class,
                   model.predict(test.loc[:, 'handicapped_infants:']))) * 100

```

```
Out[32]: Predicted   n   y
Actual
n           77   10
y            2   56
```

```
Out[32]: 8.275862068965523
```

note how most errors (10/12) correspond to democrats wrongly predicted as republicans  
in the event of **empty empirical probabilities**, there is an alpha parameter (0-1) that can be use  
for performing Laplace correction (aka smoothing) (0 = no smoothing)

```
In [33]: model = BernoulliNB(alpha=0.9).fit(train.loc[:, 'handicapped_infants:'],
                                             train.Class)
```

### 1.3 Example 3: The kNN classifier

We are going to use the famous (Fisher's or Anderson's) Iris data set, which gives the measurements in centimeters of the sepal length and width and petal length and width, respectively, for 50 flowers from each of 3 species of Iris. The species are Iris setosa, versicolor, and virginica.

```
In [34]: np.random.seed(2)
```

```
iris_data, iris_labels = load_iris(return_X_y=True)
iris_names = ['setosa', 'versicolor', 'virginica']
```

first we split a separate test set of relative size 30%

```
In [35]: X_train, X_test, y_train, y_test = train_test_split(iris_data,
                                                            iris_labels,
                                                            test_size=0.3)
```

setup a kNN model with 3 neighbours Notice there is no "learning" ... the data is the model (just test!)

```
In [36]: myknn = KNeighborsClassifier(n_neighbors=3).fit(X_train, y_train)
confusion(y_test, myknn.predict(X_test), classes=iris_names)
```

```
Out [36]: Predicted   setosa  versicolor  virginica
Actual
setosa           17           0           0
versicolor        0          15           0
virginica         0           0          13
```

rows are predictions, columns are true test targets  
How do we optimize k? One way is by using LOOCV

```
In [37]: cm, err = loocv(X_train,y_train,myknn, classes=iris_names)
```

```
cm
```

```
err*100
```

```
Out [37]: Predicted   setosa  versicolor  virginica
Actual
setosa           33           0           0
versicolor        0          32           3
virginica         0           3          34
```

```
Out [37]: 5.714285714285716
```

aha! now you see that previous training error (0%) was a little bit optimistic  
Let's loop over k

```
In [38]: np.random.seed(4321)
```

```
errors = []
for i in range(1,int(np.sqrt(X_train.shape[0]))+1):
    myknn_cv = KNeighborsClassifier(n_neighbors=i)
    _, error = loocv(X_train,y_train,myknn_cv, classes=iris_names)
    errors.append(error)

pd.DataFrame({'K':range(1,int(np.sqrt(X_train.shape[0]))+1),
              'LOOCV error':errors})
```

```
Out[38]:
```

	K	LOOCV error
0	1	0.067
1	2	0.086
2	3	0.057
3	4	0.076
4	5	0.076
5	6	0.057
6	7	0.048
7	8	0.048
8	9	0.029
9	10	0.029

It seems that k=9 is the best value.

Now we *refit* with k=9 and predict the test set

```
In [39]: myknn = KNeighborsClassifier(n_neighbors=9).fit(X_train, y_train)
confusion(y_test, myknn.predict(X_test), classes=iris_names)
(1-accuracy_score(y_test, myknn.predict(X_test)))*100
```

```
Out[39]:
```

Predicted	setosa	versicolor	virginica
Actual			
setosa	17	0	0
versicolor	0	14	1
virginica	0	0	13

```
Out[39]: 2.222222222222254
```

so our error is 2.2%

---

## 1.4 Example 4: Logistic Regression using artificial data

The goal of this example is to get acquainted with the call to `glm()` `glm()` is used to fit generalized linear models (of which both linear and logistic regression are particular cases)

You may need to recall at this point the logistic regression model ...

Let  $x$  represent a single continuous predictor

Let  $y$  represent a class ('0' or '1'), with a probability of being 1 that is related linearly to the predictor via the logit function, that is  $\text{logit}(p) = a * x + b$  (or  $\beta_1 * x + \beta_0$  if you prefer)

```

In [40]: np.random.seed(1968)

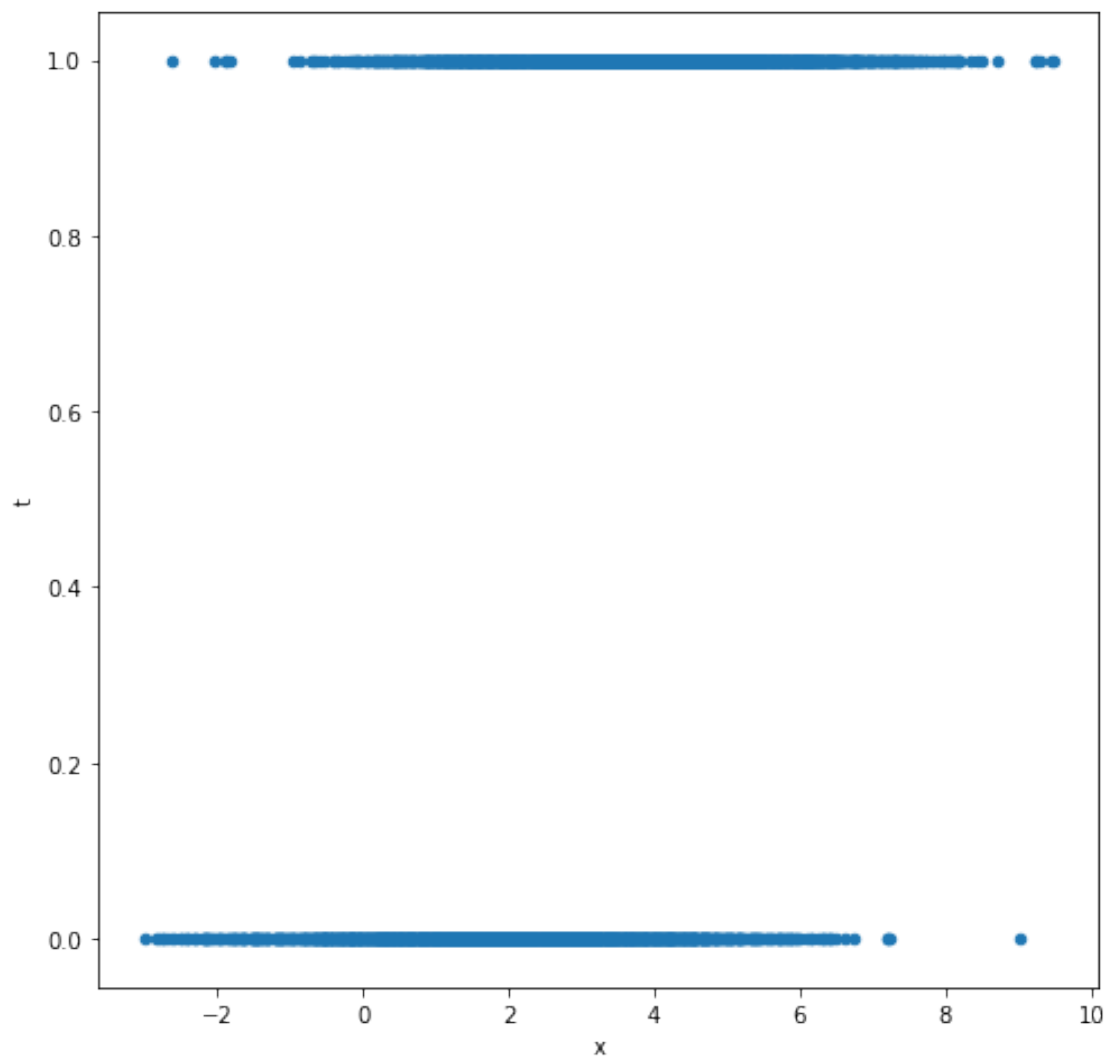
N = 4000
x = normal (3,2,N)  # generate the x_n

a = 0.6
b = -1.5 # this is the ground truth, which is unknown

p = 1/(1+np.exp( -(a*x + b) )) # generate the p_n
t = binomial(1,p, N) # generate the targets according to p
data = pd.DataFrame({'x':x, 't':t})

data.plot.scatter('x','t',figsize=(8,8));

```



```

In [41]: model = GLM.from_formula('t ~ x', data, family=Binomial())

```

```
result = model.fit()
result.summary()
```

```
Out[41]: <class 'statsmodels.iolib.summary.Summary'>
"""
                        Generalized Linear Model Regression Results
=====
Dep. Variable:          t      No. Observations:          4000
Model:                  GLM    Df Residuals:              3998
Model Family:           Binomial  Df Model:              1
Link Function:          logit    Scale:                1.0000
Method:                 IRLS     Log-Likelihood:      -2248.1
Date:                   Thu, 06 Sep 2018  Deviance:         4496.1
Time:                   14:32:49  Pearson chi2:       3.99e+03
No. Iterations:         4        Covariance Type:      nonrobust
=====
                        coef      std err          z      P>|z|      [0.025      0.975]
-----
Intercept      -1.5217      0.075     -20.278      0.000     -1.669     -1.375
x               0.6040      0.023     26.463      0.000      0.559      0.649
=====
"""
```

Obviously x is very significant (and the Intercept is always significant)

Therefore, our estimated model is  $\text{logit}(p_n) = \{\text{result.params}[1]\}x_n + \{\text{result.params}[0]\}$   
quite close to the ground truth

In general you get this as:

`result.params`

Interpretation of the coefficients:

- For a 1 unit increase in x, there is an increase in the odds for t by a factor of ...

```
In [42]: result.params
```

```
Out[42]: Intercept    -1.522
x                   0.604
dtype: float64
```

```
In [43]: np.exp(result.params[1])
```

```
Out[43]: 1.8293462220130607
```

that is almost doubling the odds (~82% more)

## 1.5 Example 5: Logistic regression for classifying spam mail

This example will also illustrate how to change the 'cut point' for prediction, when there is an interest in minimizing a particular source of errors

```
In [44]: spam = read_csv("spambase.data", delimiter=',', header=None)
         file = open('spambase.names', 'r')
         spam.columns = [n.strip() for n in file.readlines()]

         spam.head()
```

```
Out[44]:
```

	word_freq_make	word_freq_address	word_freq_all	word_freq_3d	\
0	0.00	0.64	0.64	0.0	
1	0.21	0.28	0.50	0.0	
2	0.06	0.00	0.71	0.0	
3	0.00	0.00	0.00	0.0	
4	0.00	0.00	0.00	0.0	

	word_freq_our	word_freq_over	word_freq_remove	word_freq_internet	\
0	0.32	0.00	0.00	0.00	
1	0.14	0.28	0.21	0.07	
2	1.23	0.19	0.19	0.12	
3	0.63	0.00	0.31	0.63	
4	0.63	0.00	0.31	0.63	

	word_freq_order	word_freq_mail	...	char_freq_;	char_freq(	\
0	0.00	0.00	...	0.00	0.000	
1	0.00	0.94	...	0.00	0.132	
2	0.64	0.25	...	0.01	0.143	
3	0.31	0.63	...	0.00	0.137	
4	0.31	0.63	...	0.00	0.135	

	char_freq_[	char_freq_!	char_freq_\$	char_freq_#	\
0	0.0	0.778	0.000	0.000	
1	0.0	0.372	0.180	0.048	
2	0.0	0.276	0.184	0.010	
3	0.0	0.137	0.000	0.000	
4	0.0	0.135	0.000	0.000	

	capital_run_length_average	capital_run_length_longest	\
0	3.756	61	
1	5.114	101	
2	9.821	485	
3	3.537	40	
4	3.537	40	

	capital_run_length_total	Class
0	278	1
1	1028	1

2	2259	1
3	191	1
4	191	1

[5 rows x 58 columns]

We do some basic pre-processing

```
In [45]: spam.loc[:, 'capital_run_length_average': 'capital_run_length_total'] = \
        spam.loc[:, 'capital_run_length_average': 'capital_run_length_total']. \
            apply(lambda x: np.log10(x+1))
spam = spam[spam.word_freq_george==0]
spam = spam[spam.word_freq_650==0]
spam = spam[spam.word_freq_hp==0]
spam = spam[spam.word_freq_hpl==0]
spam = spam.drop(columns=['word_freq_george', 'word_freq_650',
                        'word_freq_hp', 'word_freq_hpl'])
spam['about_money'] = spam.word_freq_free + spam.word_freq_business + \
    spam.word_freq_credit + spam.word_freq_money
spam = spam.drop(columns=['word_freq_free', 'word_freq_business',
                        'word_freq_credit', 'word_freq_money'])
Class = spam.Class    # move the Class column to the last position
spam = spam.drop(columns=['Class'])
spam['Class'] = Class

spam.shape
```

Out[45]: (2999, 51)

```
In [46]: np.random.seed(4321)
        train, test = train_test_split(spam, test_size=0.33)
```

Fit a GLM in the learning data

```
In [47]: spamM1 = GLM(train.Class,
                    add_constant(train.loc[:, 'about_money']),
                    family=Binomial())
resultM1 = spamM1.fit()
resultM1.summary()
```

```
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:880: RuntimeWarni
n_endog_mu = self._clean((1. - endog) / (1. - mu))
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/links.py:167: RuntimeWarnin
t = np.exp(-z)
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:879: RuntimeWarni
endog_mu = self._clean(endog / mu)
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:932: RuntimeWarni
special.gammaln(n - y + 1) + y * np.log(mu / (1 - mu)) +
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:932: RuntimeWarni
```



```

special.gammaln(n - y + 1) + y * np.log(mu / (1 - mu)) +
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:932: RuntimeWarni
special.gammaln(n - y + 1) + y * np.log(mu / (1 - mu)) +
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:933: RuntimeWarni
n * np.log(1 - mu)) * var_weights
/usr/local/lib64/python3.6/site-packages/statsmodels/genmod/families/family.py:933: RuntimeWarni
n * np.log(1 - mu)) * var_weights

```

```

Out[47]: <class 'statsmodels.iolib.summary.Summary'>
"""

```

# Generalized Linear Model Regression Results

```

=====
Dep. Variable:          Class    No. Observations:          2009
Model:                  GLM      Df Residuals:          1958
Model Family:           Binomial Df Model:              50
Link Function:          logit    Scale:                1.0000
Method:                 IRLS     Log-Likelihood:        nan
Date:                   Thu, 06 Sep 2018    Deviance:              nan
Time:                   14:32:54    Pearson chi2:          2.24e+04
No. Iterations:         100        Covariance Type:       nonrobust
=====

```

	coef	std err	z	P> z	[0.025
const	-5.0624	0.464	-10.904	0.000	-5.972
word_freq_make	-0.1299	0.293	-0.443	0.658	-0.705
word_freq_address	-0.1619	0.131	-1.239	0.216	-0.418
word_freq_all	-0.0884	0.167	-0.531	0.595	-0.415
word_freq_3d	350.6953	8.8e+05	0.000	1.000	-1.73e+06
word_freq_our	0.5344	0.120	4.464	0.000	0.300
word_freq_over	0.6943	0.353	1.965	0.049	0.002
word_freq_remove	1.6118	0.329	4.895	0.000	0.966
word_freq_internet	0.5349	0.148	3.621	0.000	0.245
word_freq_order	-0.1082	0.304	-0.356	0.722	-0.703
word_freq_mail	0.0561	0.090	0.625	0.532	-0.120
word_freq_receive	-0.5076	0.380	-1.334	0.182	-1.253
word_freq_will	-0.1398	0.107	-1.308	0.191	-0.349
word_freq_people	-0.0531	0.334	-0.159	0.874	-0.708
word_freq_report	0.0120	0.307	0.039	0.969	-0.590
word_freq_addresses	0.8487	0.802	1.058	0.290	-0.724
word_freq_email	0.1439	0.147	0.978	0.328	-0.144
word_freq_you	0.1415	0.051	2.782	0.005	0.042
word_freq_your	0.1381	0.074	1.870	0.061	-0.007
word_freq_font	0.1757	0.279	0.630	0.529	-0.371
word_freq_000	1.0192	0.468	2.176	0.030	0.101
word_freq_lab	-12.4533	10.458	-1.191	0.234	-32.950
word_freq_labs	-1.6974	1.046	-1.623	0.105	-3.748
word_freq_telnet	-0.2793	1.874	-0.149	0.882	-3.952

word_freq_857	-0.5357	3.463	-0.155	0.877	-7.323	
word_freq_data	-0.4946	0.292	-1.696	0.090	-1.066	
word_freq_415	424.0571	4.52e+07	9.38e-06	1.000	-8.86e+07	8
word_freq_85	-1.8880	1.518	-1.244	0.214	-4.863	
word_freq_technology	0.8304	0.547	1.519	0.129	-0.241	
word_freq_1999	0.2841	0.237	1.197	0.231	-0.181	
word_freq_parts	0.3967	2.682	0.148	0.882	-4.861	
word_freq_pm	-1.1317	0.562	-2.015	0.044	-2.233	
word_freq_direct	-0.3583	0.445	-0.806	0.421	-1.230	
word_freq_cs	-354.6653	3.85e+06	-9.21e-05	1.000	-7.55e+06	7
word_freq_meeting	-6.4294	2.328	-2.762	0.006	-10.993	
word_freq_original	-1.4478	1.044	-1.387	0.165	-3.493	
word_freq_project	-1.7884	0.827	-2.164	0.030	-3.408	
word_freq_re	-0.7498	0.195	-3.842	0.000	-1.132	
word_freq_edu	-4.7957	0.948	-5.059	0.000	-6.654	
word_freq_table	-1.4227	2.442	-0.583	0.560	-6.210	
word_freq_conference	-4.2235	1.523	-2.773	0.006	-7.209	
char_freq_;	-1.5962	0.696	-2.295	0.022	-2.960	
char_freq_(	-0.6276	0.413	-1.519	0.129	-1.437	
char_freq_[	-0.0500	0.979	-0.051	0.959	-1.969	
char_freq_!	0.2124	0.061	3.471	0.001	0.092	
char_freq_\$	5.7022	1.098	5.193	0.000	3.550	
char_freq_#	1.2748	1.654	0.771	0.441	-1.966	
capital_run_length_average	0.7272	0.786	0.925	0.355	-0.813	
capital_run_length_longest	0.5299	0.448	1.182	0.237	-0.349	
capital_run_length_total	1.8490	0.311	5.948	0.000	1.240	
about_money	0.6960	0.117	5.926	0.000	0.466	

=====

"""

We can see that there are some variables that have small weights and are probably not very relevant. The R notebook uses stepwise variable selection to simplify the model.

Statsmodels does not have stepwise variable selection, but we can use crossvalidated Recursive Forward Elimination (RFE) with the implementation of logistic regression from scikit learn. RFE does the same thing as stepwise variable selection but uses accuracy to select the best model using cross validation. The implementation of logistic regression in scikit-learn is more sophisticated and uses regularization so the results will be different than in the R notebook.

```
In [48]: # we use L1 regularization to make 0 a large number
# of the weights, the lower the C the more attributes will be discarded
logreg = LogisticRegression(solver='liblinear',penalty='l1',C=1)
#njobs = -1 means that all the cores from the CPU are used
rfe = RFECV(estimator=logreg,cv=10,n_jobs=-1)
rfe.fit(train.loc[:, 'about_money'],train.Class);

In [49]: print('Features Selected:',rfe.n_features_)
print('\n Ranking of features')
sel = pd.DataFrame({'features': train.columns[: -1],
```

```

        'ranking': rfe.ranking_,
        'selected': rfe.support_})
sel.sort_values(by='ranking')

```

Features Selected: 38

Ranking of features

```

Out[49]:

```

	features	ranking	selected
24	word_freq_data	1	True
27	word_freq_technology	1	True
28	word_freq_1999	1	True
30	word_freq_pm	1	True
31	word_freq_direct	1	True
32	word_freq_cs	1	True
33	word_freq_meeting	1	True
34	word_freq_original	1	True
35	word_freq_project	1	True
26	word_freq_85	1	True
36	word_freq_re	1	True
39	word_freq_conference	1	True
40	char_freq_;	1	True
41	char_freq_(	1	True
43	char_freq_!	1	True
44	char_freq_\$	1	True
45	char_freq_#	1	True
46	capital_run_length_average	1	True
47	capital_run_length_longest	1	True
37	word_freq_edu	1	True
48	capital_run_length_total	1	True
49	about_money	1	True
11	word_freq_will	1	True
10	word_freq_receive	1	True
21	word_freq_labs	1	True
5	word_freq_over	1	True
4	word_freq_our	1	True
14	word_freq_addresses	1	True
15	word_freq_email	1	True
6	word_freq_remove	1	True
16	word_freq_you	1	True
17	word_freq_your	1	True
18	word_freq_font	1	True
19	word_freq_000	1	True
20	word_freq_lab	1	True
1	word_freq_address	1	True
3	word_freq_3d	1	True
7	word_freq_internet	1	True

9	word_freq_mail	2	False
2	word_freq_all	3	False
0	word_freq_make	4	False
42	char_freq_	5	False
29	word_freq_parts	6	False
8	word_freq_order	7	False
38	word_freq_table	8	False
13	word_freq_report	9	False
23	word_freq_857	10	False
22	word_freq_telnet	11	False
25	word_freq_415	12	False
12	word_freq_people	13	False

We get the estimator from the RFE and the list of selected variable to slice the data matrix

```
In [50]: resultM1 = rfe.estimator_
sel_features = list(sel.features[sel.selected])
sel_features
```

```
Out[50]: ['word_freq_address',
'word_freq_3d',
'word_freq_our',
'word_freq_over',
'word_freq_remove',
'word_freq_internet',
'word_freq_receive',
'word_freq_will',
'word_freq_addresses',
'word_freq_email',
'word_freq_you',
'word_freq_your',
'word_freq_font',
'word_freq_000',
'word_freq_lab',
'word_freq_labs',
'word_freq_data',
'word_freq_85',
'word_freq_technology',
'word_freq_1999',
'word_freq_pm',
'word_freq_direct',
'word_freq_cs',
'word_freq_meeting',
'word_freq_original',
'word_freq_project',
'word_freq_re',
'word_freq_edu',
'word_freq_conference',
```

```
'char_freq_',
'char_freq(',
'char_freq!',
'char_freq$',
'char_freq#',
'capital_run_length_average',
'capital_run_length_longest',
'capital_run_length_total',
'about_money']
```

We define now a convenience function:

'P' is a parameter; whenever our filter assigns spam with probability at least P then we predict spam

```
In [51]: def spam_acc(P=0.5):
    # We use predict_proba instead of prediction to obtain
    # the probabilities of the classes and
    # we select only the probability for class 1 as
    # the other is just the complementary

    # Accuracy in training
    pred = resultM1.predict_proba(train.loc[:,sel_features])[:,1]
    lab_tr = [1 if i>=P else 0 for i in pred]
    df_tr=confusion(train.Class,lab_tr, classes=['nospam','spam'])

    # Accuracy in test
    pred = resultM1.predict_proba(test.loc[:,sel_features])[:,1]
    lab_ts = [1 if i>=P else 0 for i in pred]
    df_ts=confusion(test.Class,lab_ts, classes=['nospam','spam'])

    return df_tr, (1-accuracy_score(train.Class,lab_tr))*100,\
           df_ts, (1-accuracy_score(test.Class,lab_ts))*100
```

```
In [52]: c_tr,e_tr,c_ts,e_ts= spam_acc()
c_tr
print(f'Training error: {e_tr}%')
c_ts
print(f'Test error: {e_ts}%')
```

```
Out[52]: Predicted nospam spam
Actual
nospam          759    90
spam             67  1093
```

Training error: 7.81483325037332%

```
Out[52]: Predicted nospam spam
Actual
```

nospam	382	41
spam	32	535

Test error: 7.373737373737377%

Although the errors are quite low still one could argue that we should try to lower the probability of predicting spam when it is not. We can do this (at the expense of increasing the converse probability) by:

```
In [53]: c_tr,e_tr,c_ts,e_ts= spam_acc(0.7)
         c_tr
         print(f'Training error: {e_tr}%')
         c_ts
         print(f'Test error: {e_ts}%')
```

```
Out[53]: Predicted  nospam  spam
         Actual
         nospam      801    48
         spam       156   1004
```

Training error: 10.154305624688897%

```
Out[53]: Predicted  nospam  spam
         Actual
         nospam      403    20
         spam       78   489
```

Test error: 9.898989898989896%

So we get a much better spam filter; notice that the filter has a very low probability of predicting spam when it is not (which is the delicate case), of about

```
In [54]: c_ts.loc['nospam','spam'] /c_ts.loc['nospam'].sum()*100
```

```
Out[54]: 4.7281323877068555
```