Phase_1

May 14, 2020

[9]:
```
import pandas as pd
import pylab as pl
import numpy as np
import scipy.optimize as opt
from sklearn import preprocessing
from sklearn.model_selection import train_test_split
%matplotlib inline
import matplotlib.pyplot as plt
```

[6]:
```
#!pip install ibm_watson wget
!wget -O italy_normal2.csv https://drive.google.com/open?
 ↪id=1CX6txmOBM4flZITxNCa_YVDdb76nFrGu
```

```
--2020-05-05 18:12:19--
https://drive.google.com/open?id=1CX6txmOBM4flZITxNCa_YVDdb76nFrGu
Resolving drive.google.com (drive.google.com)… 172.217.1.174,
2607:f8b0:400b:809::200e
Connecting to drive.google.com (drive.google.com)|172.217.1.174|:443…
connected.
HTTP request sent, awaiting response… 307 Temporary Redirect
Location: https://drive.google.com/file/d/1CX6txmOBM4flZITxNCa_YVDdb76nFrGu/view
?usp=drive_open [following]
--2020-05-05 18:12:20--  https://drive.google.com/file/d/1CX6txmOBM4flZITxNCa_YV
Ddb76nFrGu/view?usp=drive_open
Reusing existing connection to drive.google.com:443.
HTTP request sent, awaiting response… 200 OK
Length: unspecified [text/html]
Saving to: 'italy_normal2.csv'

italy_normal2.csv       [ <=>                ]  68.79K  --.-KB/s    in 0.02s

2020-05-05 18:12:21 (4.20 MB/s) - 'italy_normal2.csv' saved [70445]
```

[10]:
```
covid19 = pd.read_csv("italy_normal2.csv")
covid19.head()
```

[10]:

	age	gender	employmentstatus	tested	Construction	Deliveringtohomes	\
0	57	2	1	4.0	0	0	
1	63	2	1	4.0	0	0	
2	57	2	1	4.0	0	0	
3	23	2	1	4.0	0	0	
4	60	1	1	4.0	0	0	

	Foodretail	Healthcare	Logisticsothertransportation	Manufacturing	…	\
0	0	0	0	0	…	
1	0	0	0	0	…	
2	0	0	0	0	…	
3	0	0	0	0	…	
4	0	0	0	0	…	

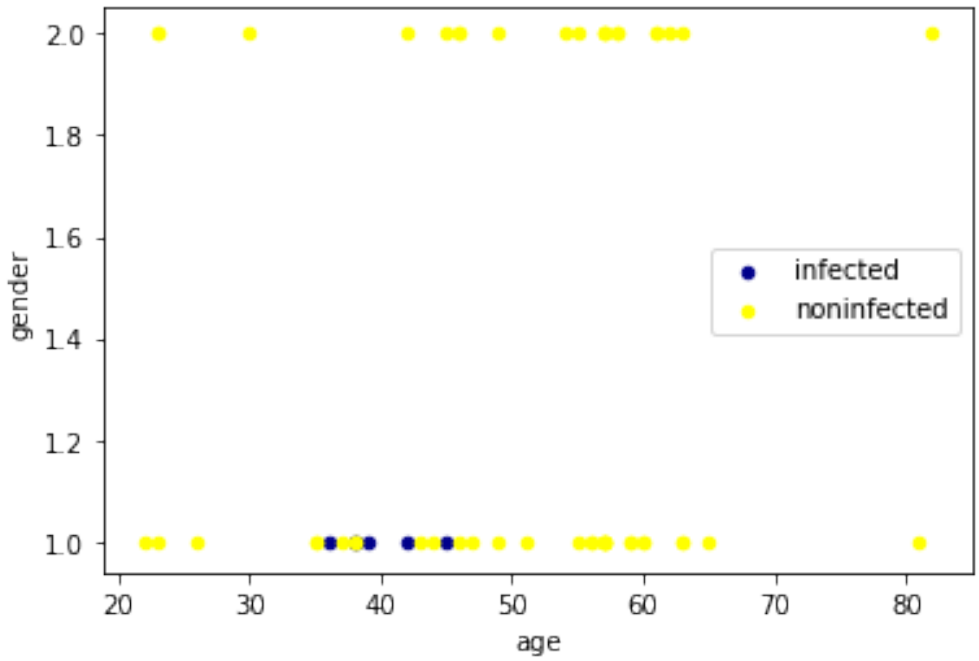
	Chronicobstructivepulmonary disease(COPD)	Diabetes	Epilepsy	\
0	0.0	0.0	0.0	
1	0.0	0.0	0.0	
2	0.0	0.0	0.0	
3	0.0	0.0	0.0	
4	0.0	0.0	0.0	

	Heartdisease	Highbloodpressure	Highcholesterol	HIVAids	\
0	0.0	0.0	0.0	0.0	
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	1.0	0.0	

	Mentalhealthcondition	MultipleSclerosis	another
0	0.0	0.0	1.0
1	0.0	0.0	0.0
2	0.0	0.0	1.0
3	0.0	0.0	1.0
4	0.0	0.0	0.0

[5 rows x 29 columns]

[11]:
```
ax = covid19[covid19['tested'] == 1][0:50].plot(kind='scatter', x='age',␣
 ↪y='gender', color='DarkBlue', label='infected');
covid19[covid19['tested'] == 4][0:50].plot(kind='scatter', x='age', y='gender',␣
 ↪color='Yellow', label='noninfected', ax=ax);
plt.show()
```



[13]:
```
#to fix the data from any nan or infinte value (very important pre process lone)
covid = covid19.replace(r'^\s*$', np.NaN, regex=True)
covid = covid.fillna(0)
np.any(np.isnan(covid))
np.all(np.isfinite(covid))
covid.dtypes
```

[13]:
```
age                              int64
gender                           int64
employmentstatus                 int64
tested                         float64
Construction                     int64
Deliveringtohomes                int64
Foodretail                       int64
Healthcare                       int64
Logisticsothertransportation     int64
Manufacturing                    int64
Policingorprisons                int64
Publictransport                  int64
School                           int64
Socialcare                       int64
notworking                       int64
Arthritis                      float64
Asthma                         float64
Cancer                         float64
```

```
Cysticfibrosis                                  float64
Chronicobstructivepulmonary disease(COPD)       float64
Diabetes                                        float64
Epilepsy                                        float64
Heartdisease                                    float64
Highbloodpressure                               float64
Highcholesterol                                 float64
HIVAids                                         float64
Mentalhealthcondition                           float64
MultipleSclerosis                               float64
another                                         float64
dtype: object
```

[127]:
```
feature_df = covid[['age', 'gender', 'employmentstatus', 'tested',␣
↪'Construction',
       'Deliveringtohomes', 'Foodretail', 'Healthcare',
       'Logisticsothertransportation', 'Manufacturing', 'Policingorprisons',
       'Publictransport', 'School', 'Socialcare', 'notworking', 'Arthritis',
       'Asthma', 'Cancer', 'Cysticfibrosis',
       'Chronicobstructivepulmonary disease(COPD)', 'Diabetes', 'Epilepsy',
       'Heartdisease', 'Highbloodpressure', 'Highcholesterol', 'HIVAids',
       'Mentalhealthcondition', 'MultipleSclerosis', 'another']]
X = np.asarray(feature_df)
X[0:5]
```

[127]:
```
array([[57.,  2.,  1.,  4.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  1.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  0.,  1.],
       [63.,  2.,  1.,  4.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  0.,  1.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  0.,  0.],
       [57.,  2.,  1.,  4.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  0.,  1.],
       [23.,  2.,  1.,  4.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  1.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  0.,  1.],
       [60.,  1.,  1.,  4.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,
         0.,  1.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  1.,  0.,  0.,
         0.,  0.,  0.]])
```

[128]:
```
y = np.asarray(covid['tested'])
y [0:100]
```

[128]:
```
array([4, 4, 4, 4, 4, 4, 4, 4, 2, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4,
       4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4,
       4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4,
```

```
       4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 4, 3, 4, 4, 4, 4, 4, 4, 4, 4, 4,
       4, 4, 4, 4, 2, 4, 4, 4, 4, 4, 4, 4])
```

[129]:
```
X_train, X_test, y_train, y_test = train_test_split( X, y, test_size=0.2,␣
 ↪random_state=4)
print ('Train set:', X_train.shape,  y_train.shape)
print ('Test set:', X_test.shape,  y_test.shape)
```

```
Train set: (1672, 29) (1672,)
Test set: (419, 29) (419,)
```

[130]:
```
from sklearn import svm
clf = svm.SVC(kernel='rbf')
clf.fit(X_train, y_train)
```

```
/home/jupyterlab/conda/envs/python/lib/python3.6/site-
packages/sklearn/svm/base.py:196: FutureWarning: The default value of gamma will
change from 'auto' to 'scale' in version 0.22 to account better for unscaled
features. Set gamma explicitly to 'auto' or 'scale' to avoid this warning.
  "avoid this warning.", FutureWarning)
```

[130]:
```
SVC(C=1.0, cache_size=200, class_weight=None, coef0=0.0,
  decision_function_shape='ovr', degree=3, gamma='auto_deprecated',
  kernel='rbf', max_iter=-1, probability=False, random_state=None,
  shrinking=True, tol=0.001, verbose=False)
```

[131]:
```
yhat = clf.predict(X_test)
yhat [0:5]
```

[131]:
```
array([4, 4, 4, 4, 4])
```

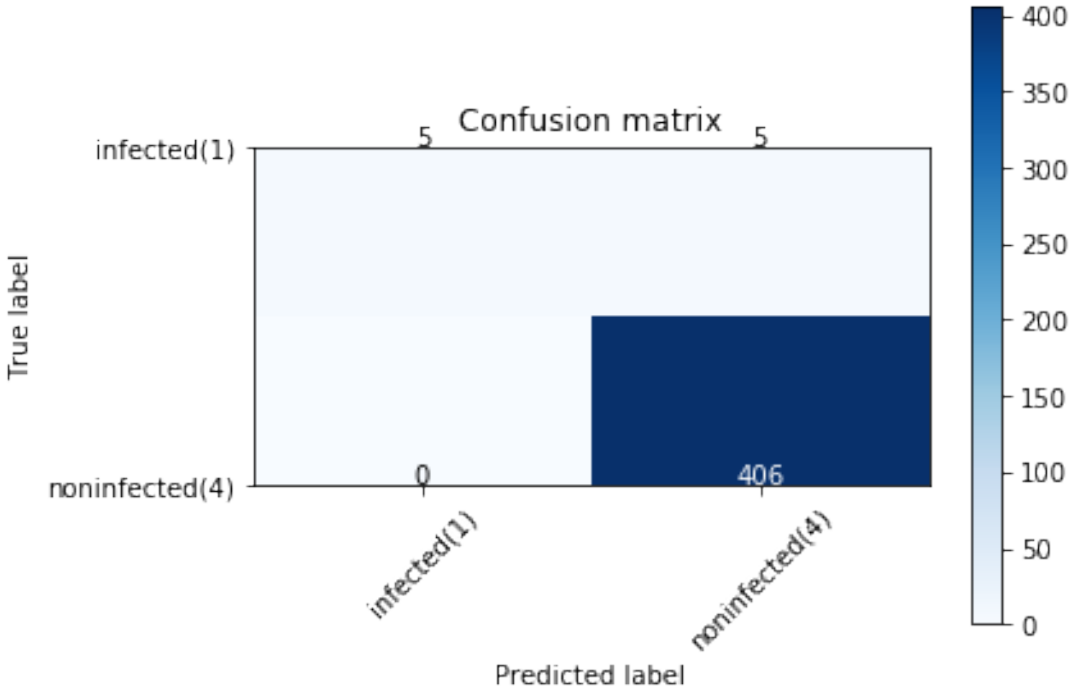
[132]:
```
from sklearn.metrics import classification_report, confusion_matrix
import itertools
```

[133]:
```
def plot_confusion_matrix(cm, classes,
                          normalize=False,
                          title='Confusion matrix',
                          cmap=plt.cm.Blues):
    """
    This function prints and plots the confusion matrix.
    Normalization can be applied by setting `normalize=True`.
    """
    if normalize:
        cm = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
        print("Normalized confusion matrix")
    else:
        print('Confusion matrix, without normalization')
```

```
    print(cm)

    plt.imshow(cm, interpolation='nearest', cmap=cmap)
    plt.title(title)
    plt.colorbar()
    tick_marks = np.arange(len(classes))
    plt.xticks(tick_marks, classes, rotation=45)
    plt.yticks(tick_marks, classes)

    fmt = '.2f' if normalize else 'd'
    thresh = cm.max() / 2.
    for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
        plt.text(j, i, format(cm[i, j], fmt),
                 horizontalalignment="center",
                 color="white" if cm[i, j] > thresh else "black")

    plt.tight_layout()
    plt.ylabel('True label')
    plt.xlabel('Predicted label')
```

[135]:
```
# Compute confusion matrix
cnf_matrix = confusion_matrix(y_test, yhat, labels=[2,4])
np.set_printoptions(precision=2)

print (classification_report(y_test, yhat))

# Plot non-normalized confusion matrix
plt.figure()
plot_confusion_matrix(cnf_matrix,␣
 ↪classes=['infected(1)','noninfected(4)'],normalize= False,  title='Confusion␣
 ↪matrix')
```

```
              precision    recall  f1-score   support

           1       0.00      0.00      0.00         1
           2       0.83      0.50      0.62        10
           3       0.00      0.00      0.00         2
           4       0.98      1.00      0.99       406

   micro avg       0.98      0.98      0.98       419
   macro avg       0.45      0.38      0.40       419
weighted avg       0.97      0.98      0.98       419

Confusion matrix, without normalization
[[  5   5]
 [  0 406]]
```



[136]:
```
#f1_score from sklearn library
from sklearn.metrics import f1_score
f1_score(y_test, yhat, average='weighted')
```

```
/home/jupyterlab/conda/envs/python/lib/python3.6/site-
packages/sklearn/metrics/classification.py:1143: UndefinedMetricWarning: F-score
is ill-defined and being set to 0.0 in labels with no predicted samples.
  'precision', 'predicted', average, warn_for)
```

[136]: 0.9756083878995574

[137]:
```
#jaccard index for accuracy
from sklearn.metrics import jaccard_similarity_score
jaccard_similarity_score(y_test, yhat)
```

[137]: 0.9809069212410502

[ ]:

[ ]: