

```
In [1]: ### BREAST CANCER CASES ###
        ##### SVM WITH RBF KERNEL CODE IN JUPYTER NOTEBOOK #####
```

```
In [2]: ## Modules required
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
import math
import keras
import tensorflow as tf
import warnings
warnings.filterwarnings("ignore")
```

```
In [3]: # Code
BC = (pd.read_excel('cancer.xlsx'))
```

```
In [4]: BC.head()
```

```
Out[4]:
```

	PAT_ID	Race	MarST	Gender	PatStatus	AgeDiag	Grade	Stability	No.Visits	Lstay	...	LyNode	Amorx
0	32400010	3	1	0	1	52	3	0	5	1	...	1	
1	32400023	3	1	0	1	48	3	0	4	3	...	1	
2	32400073	3	0	1	1	61	3	0	9	1	...	1	
3	32400073	3	1	1	1	63	2	0	3	5	...	1	
4	32400396	3	0	0	0	69	2	0	7	9	...	1	

5 rows × 26 columns

```
In [5]: #Import 'train_test_split' from 'sklearn.model_selection'
        from sklearn.model_selection import train_test_split

        #Import numpy#
import numpy as np
```

```
In [6]: y = BC.PatStatus
        x = BC.drop(['PatStatus'], axis = 1)
```

```
In [7]: #Split the data into train and test sets #
        #Import 'train_test_split' from 'sklearn.model_selection'
        from sklearn.model_selection import train_test_split
        x_train, x_test, y_train, y_test=train_test_split(x,y, test_size=0.2, random_state=
        123)

        ## Scaling the data
        from sklearn.preprocessing import MinMaxScaler
        from sklearn import preprocessing
        import numpy as np

        min_max_scaler = preprocessing.MinMaxScaler()
        x_train_minmax = min_max_scaler.fit_transform(x_train)
        x_test_minmax = min_max_scaler.fit_transform(x_test)
```

```
In [8]: x_train = x_train_minmax  
x_test = x_test_minmax
```

```
In [9]: x_train.shape
```

```
Out[9]: (80001, 24)
```

```
In [10]: x_test.shape
```

```
Out[10]: (20001, 24)
```

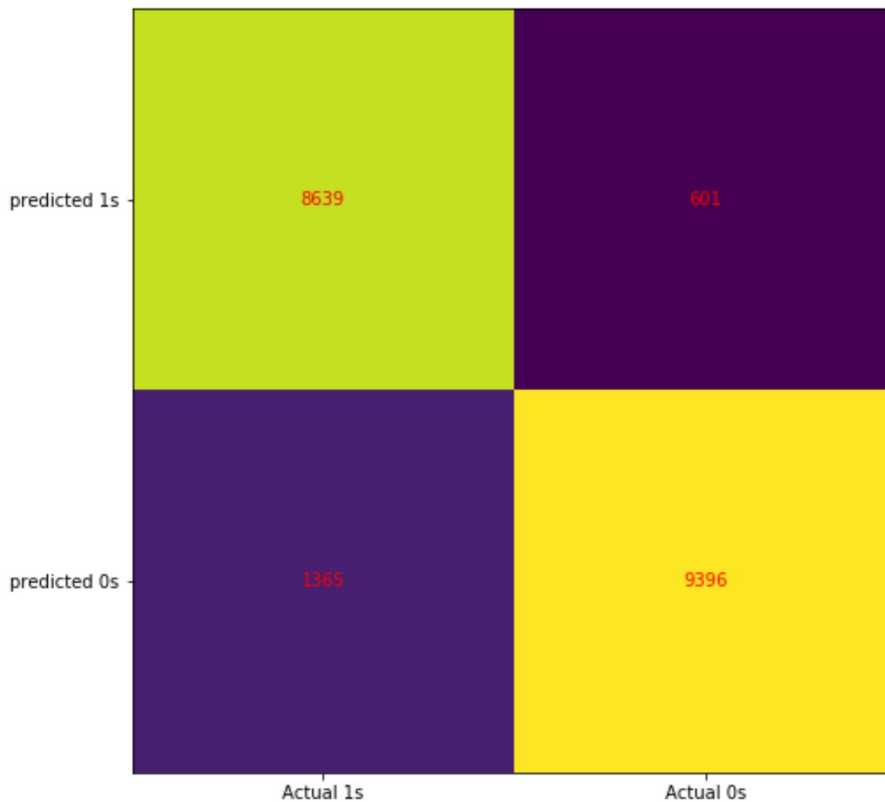
```
In [11]: ## Fitting the model  
## Models required  
from keras.preprocessing import image  
from keras.preprocessing.image import load_img  
from keras.preprocessing.image import img_to_array  
from keras.applications.imagenet_utils import decode_predictions  
from keras.utils import layer_utils, np_utils  
from sklearn.metrics import confusion_matrix, classification_report  
import tensorflow as tf  
from hyperas import optim  
from hyperas.distributions import choice, uniform  
import warnings  
warnings.filterwarnings("ignore")  
from sklearn.datasets import make_classification  
from sklearn.metrics import confusion_matrix, roc_curve, roc_auc_score, plot_roc_curve  
from sklearn.model_selection import cross_val_score, cross_validate  
from sklearn.svm import SVC  
  
# SVM classifier with Gaussian RBF kernel  
classifier = SVC(kernel='rbf', random_state=0)  
classifier.fit(x_train, y_train)
```

```
Out[11]: SVC(C=1.0, break_ties=False, cache_size=200, class_weight=None, coef0=0.0,  
decision_function_shape='ovr', degree=3, gamma='scale', kernel='rbf',  
max_iter=-1, probability=False, random_state=0, shrinking=True, tol=0.001,  
verbose=False)
```

```
In [12]: # predict with splitted test data  
y_pred = classifier.predict(x_test)
```

```
In [13]: ##Fitting the neural network model using training dataset  
tns_probs=[0 for _ in range(len(y_test))]
```

```
In [14]: ## CONFUSION MATRIX FOR BOTH SEX DATA
test_cm = confusion_matrix(y_test, y_pred)
fig, ax = plt.subplots(figsize = (8, 8))
ax.imshow(test_cm)
ax.grid(False)
ax.xaxis.set(ticks=(0,1), ticklabels=('Actual 1s', 'Actual 0s'))
ax.yaxis.set(ticks=(0,1), ticklabels=('predicted 1s', 'predicted 0s'))
ax.set_ylim(1.5, -0.5)
for i in range(2):
    for j in range(2):
        ax.text(j, i, test_cm[i, j], ha= 'center', va= 'center', color= 'red')
plt.show()
```



```
In [15]: ## Error for the prediction for test dataset outcomes
test_error = (test_cm[0,1] + test_cm[1,0])/np.sum(test_cm)
print(test_error)
```

0.09829508524573771

```
In [16]: ## Accuracy of prediction
1-test_error
```

Out[16]: 0.9017049147542623

```
In [17]: ## Sensitivity Analysis
test_sens = test_cm[1, 1]/(test_cm[1, 1] + test_cm[0, 1])
print(test_sens)
```

0.9398819645893768

```
In [18]: ## Specificity Analysis
test_spec = test_cm[0, 0]/(test_cm[0, 0]+test_cm[1, 0])
print(test_spec)

0.8635545781687325
```

```
In [19]: ## PPV Analysis
test_npv = test_cm[1, 1]/(test_cm[1, 1] + test_cm[1, 0])
print(test_npv)

0.8731530526902704
```

```
In [20]: ## NPV Analysis
test_npv = test_cm[0, 0]/(test_cm[0, 0]+test_cm[0, 1])
print(test_npv)

0.93495670995671
```

```
In [21]: ## The AUC Score
test_auc = roc_auc_score(y_test, tns_probs)
y_pred_auc = np.round(roc_auc_score(y_test, y_pred), decimals = 2)
```

```
In [22]: ## calculate ROC Curves
test_fpr, test_tpr, _ = roc_curve(y_test, tns_probs)
y_pred_fpr, y_pred_tpr, _ = roc_curve(y_test, y_pred)
```

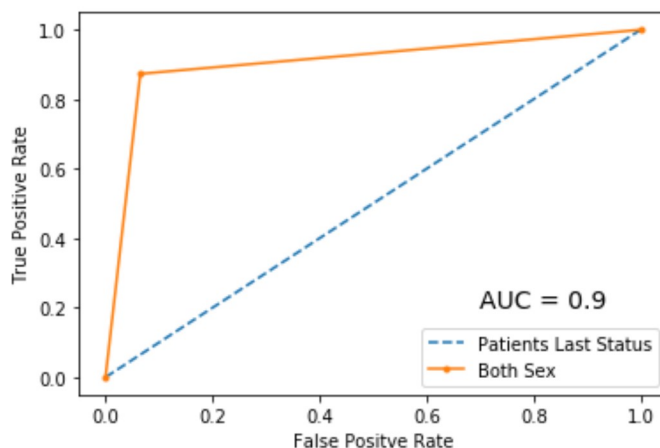
```
In [23]: ## Plot Curve for the model
import numpy as np
import matplotlib.pyplot as plt

plt.plot(test_fpr, test_tpr, linestyle = '--', label = 'Patients Last Status')
plt.plot(y_pred_fpr, y_pred_tpr, marker = '.', label = 'Both Sex')
plt.text(0.7, 0.2, "AUC = " + str(y_pred_auc), fontsize = 14)

## Axis lable
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")

## Show Legend
plt.legend()
```

Out[23]: <matplotlib.legend.Legend at 0x15ac9843788>



In []:

In []:

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```
In [1]: ## CONSIDER THE NEURAL NETWORK FOR EACH GENDER SEPARATELY
## Modules required
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
```

```
In [2]: # Code
MBC = (pd.read_excel('MBC.xlsx'))
```

```
In [3]: #Import 'train_test_split' from 'sklearn.model_selection'
from sklearn.model_selection import train_test_split
#Import numpy#
import numpy as np
```

```
In [4]: ##### THE MALE DATASET
my=MBC.PatStatus
mx=MBC.drop(['PatStatus', 'Gender'], axis=1)
```

```
In [5]: ## CONSIDER RBF FITTING FOR THE MALE GENDER
```

```
In [6]: #Split the Male data into train and test sets #
#Import 'train_test_split' from 'sklearn.model_selection'
from sklearn.model_selection import train_test_split
#Import numpy#
import numpy as np
mx_train, mx_test, my_train, my_test=train_test_split(mx,my, test_size=0.2, random_
state=124)
```

```
In [7]: mx_train.head()
```

Out[7]:

	Race	MarST	AgeDiag	Grade	Stability	No.Visits	Lstay	Laterality	FamHist	PrioBSurg	...	LyNode
12630	1	1	77	3	0	10	9	5	1	0	...	1
6329	1	0	54	2	0	15	1	3	1	0	...	1
14089	10	0	63	3	0	16	9	8	1	0	...	1
15211	1	1	61	1	0	16	12	4	1	0	...	1
8895	1	1	69	3	0	16	9	9	1	0	...	1

5 rows × 23 columns

```
In [8]: mx_test.head()
```

```
Out[8]:
```

	Race	MarST	AgeDiag	Grade	Stability	No.Visits	Lstay	Laterality	FamHist	PrioBSurg	...	LyNode
1541	1	1	70	2	0	10	9	8	1	0	...	1
8170	1	0	59	2	0	15	1	1	1	0	...	1
6776	1	0	46	3	0	14	9	3	1	0	...	1
4929	2	0	48	1	0	14	9	9	1	0	...	1
13849	1	0	58	1	0	15	9	8	1	0	...	1

5 rows × 23 columns

```
In [9]: mx_train.shape
```

```
Out[9]: (12479, 23)
```

```
In [10]: mx_test.shape
```

```
Out[10]: (3120, 23)
```

```
In [11]: ## Scaling the male data set
from sklearn.preprocessing import MinMaxScaler
from sklearn import preprocessing
import numpy as np

min_max_scaler = preprocessing.MinMaxScaler()
mx_train_minmax = min_max_scaler.fit_transform(mx_train)
mx_test_minmax = min_max_scaler.fit_transform(mx_test)
```

```
In [12]: mx_train = mx_train_minmax
mx_test = mx_test_minmax
```

```
In [13]: ## FITTING NEURAL NETWORK FOR MALE DATA
```

```
In [14]: from sklearn.neural_network import MLPClassifier
from sklearn.datasets import make_classification
from sklearn.metrics import confusion_matrix, roc_curve, roc_auc_score, plot_roc_curve
from sklearn.model_selection import cross_val_score, cross_validate
```

```
In [15]: ##Fitting the neural network model using training dataset
tns_probs=[0 for _ in range(len(my_test))]
```

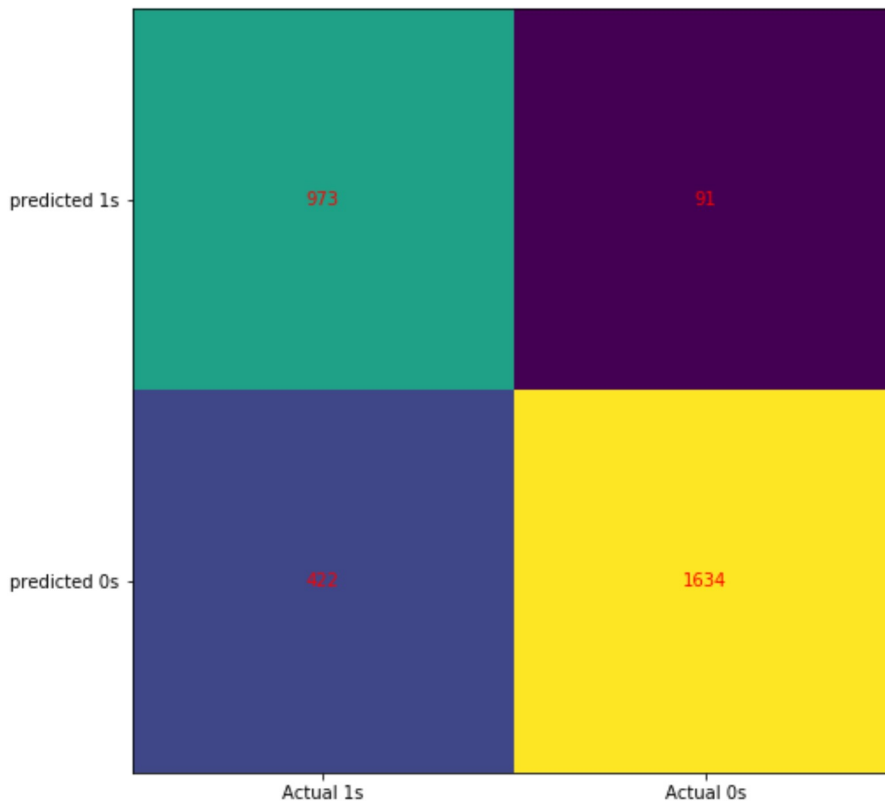
```
In [16]: ## Fitting the model
# SVM classifier with Gaussian RBF kernel
from sklearn.svm import SVC
classifier = SVC(kernel='rbf', random_state=0)
classifier.fit(mx_train, my_train)
```

```
Out[16]: SVC(C=1.0, break_ties=False, cache_size=200, class_weight=None, coef0=0.0,
decision_function_shape='ovr', degree=3, gamma='scale', kernel='rbf',
max_iter=-1, probability=False, random_state=0, shrinking=True, tol=0.001,
verbose=False)
```

```
In [17]: ### PREDICTION USING THE TEST DATASET
```

```
In [18]: ### Getting the prediction for the Testing dataset
my_pred = classifier.predict(mx_test)
```

```
In [19]: ## CONFUSION MATRIX FOR MALE DATA
mtest_cm = confusion_matrix(my_test, my_pred)
fig, ax = plt.subplots(figsize = (8, 8))
ax.imshow(mtest_cm)
ax.grid(False)
ax.xaxis.set(ticks=(0,1), ticklabels=('Actual 1s', 'Actual 0s'))
ax.yaxis.set(ticks=(0,1), ticklabels=('predicted 1s', 'predicted 0s'))
ax.set_ylim(1.5, -0.5)
for i in range(2):
    for j in range(2):
        ax.text(j, i, mtest_cm[i, j], ha= 'center', va= 'center', color= 'red')
plt.show()
```



```
In [20]: ## Error for the prediction for test dataset outcomes
mtest_error = (mtest_cm[0,1] + mtest_cm[1,0])/np.sum(mtest_cm)
print(mtest_error)

0.16442307692307692
```

```
In [21]: ## Accuracy of prediction
1-mtest_error
```

```
Out[21]: 0.8355769230769231
```

```
In [22]: ## Sensitivity Analysis
mtest_sens = mtest_cm[1, 1]/(mtest_cm[1, 1] + mtest_cm[0, 1])
print(mtest_sens)

0.9472463768115942
```

```
In [23]: ## Specificity Analysis
mtest_spec = mtest_cm[0, 0]/(mtest_cm[0, 0]+ mtest_cm[1, 0])
print(mtest_spec)

0.6974910394265234
```

```
In [24]: ## PPV Analysis
mtest_npv = mtest_cm[1, 1]/(mtest_cm[1, 1] + mtest_cm[1, 0])
print(mtest_npv)

0.7947470817120622
```

```
In [25]: ## NPV Analysis
mtest_npv = mtest_cm[0, 0]/(mtest_cm[0, 0] + mtest_cm[0, 1])
print(mtest_npv)

0.9144736842105263
```

```
In [26]: ## The AUC Score
tns_probs=[0 for _ in range(len(my_test))]
mtest_auc = roc_auc_score(my_test, tns_probs)
my_pred_auc = np.round(roc_auc_score(my_test, my_pred), decimals = 2)
```

```
In [27]: ## calculate ROC Curves
mtest_fpr, mtest_tpr, _ = roc_curve(my_test, tns_probs)
my_pred_fpr, my_pred_tpr, _ = roc_curve(my_test, my_pred)
```

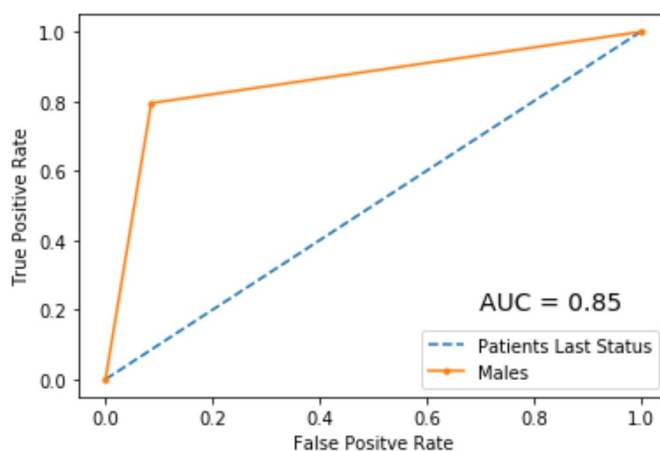
```
In [28]: ## Plot Curve for the model
import numpy as np
import matplotlib.pyplot as plt

plt.plot(mtest_fpr, mtest_tpr, linestyle = '--', label = 'Patients Last Status')
plt.plot(my_pred_fpr, my_pred_tpr, marker = '.', label = 'Males')
plt.text(0.7, 0.2, "AUC = " + str(my_pred_auc), fontsize = 14)

## Axis lable
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")

## Show Legend
plt.legend()
```

Out[28]: <matplotlib.legend.Legend at 0x2781ec16308>



In []:

In []:

In []:

```
In [1]: ## CONSIDERING THE FEMALE DATA
## Modules required
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
FBC = (pd.read_excel('FBC.xlsx'))

fy=FBC.PatStatus
fx=FBC.drop(['PatStatus','Gender'],axis=1)
```

```
In [2]: #Import 'train_test_split' from 'sklearn.model_selection'
from sklearn.model_selection import train_test_split

#Import numpy#
import numpy as np
#Split the Male data into train and test sets #
fx_train, fx_test, fy_train, fy_test=train_test_split(fx,fy, test_size=0.2, random_state=125)
```

```
In [3]: # Scaling the female data
from sklearn.preprocessing import MinMaxScaler
from sklearn import preprocessing
import numpy as np

min_max_scaler = preprocessing.MinMaxScaler()
fx_train_minmax = min_max_scaler.fit_transform(fx_train)
fx_test_minmax = min_max_scaler.fit_transform(fx_test)
```

```
In [4]: fx_train = fx_train_minmax
fx_test = fx_test_minmax
```

```
In [5]: ### FITTING THE NEURAL NETWORK USING THE FEMALE TRAINING DATASET
from sklearn.neural_network import MLPClassifier
from sklearn.datasets import make_classification
from sklearn.metrics import confusion_matrix, roc_curve, roc_auc_score, plot_roc_curve
from sklearn.model_selection import cross_val_score, cross_validate

tns_probs=[0 for _ in range(len(fy_test))]
```

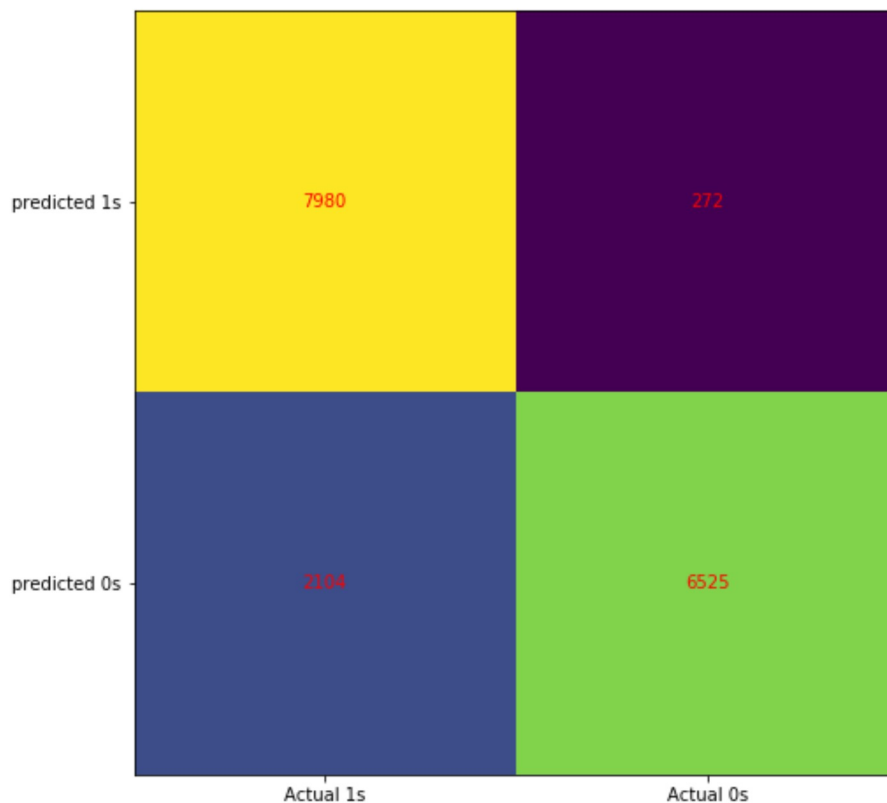
```
In [6]: ## Fitting the model
# SVM classifier with Gaussian RBF kernel
from sklearn.svm import SVC
classifier = SVC(kernel='rbf',random_state=0)
classifier.fit(fx_train,fy_train)
```

```
Out[6]: SVC(C=1.0, break_ties=False, cache_size=200, class_weight=None, coef0=0.0,
decision_function_shape='ovr', degree=3, gamma='scale', kernel='rbf',
max_iter=-1, probability=False, random_state=0, shrinking=True, tol=0.001,
verbose=False)
```

```
In [7]: ## PREDICTION USING THE TEST DATASET
```

```
In [8]: ### Getting the prediction for the Testing dataset  
fy_pred = classifier.predict(fx_test)
```

```
In [9]: ## CONFUSION MATRIX FOR MALE DATA  
ftest_cm = confusion_matrix(fy_test, fy_pred)  
fig, ax = plt.subplots(figsize = (8, 8))  
ax.imshow(ftest_cm)  
ax.grid(False)  
ax.xaxis.set(ticks=(0,1), ticklabels=('Actual 1s', 'Actual 0s'))  
ax.yaxis.set(ticks=(0,1), ticklabels=('predicted 1s', 'predicted 0s'))  
ax.set_ylim(1.5, -0.5)  
for i in range(2):  
    for j in range(2):  
        ax.text(j, i, ftest_cm[i, j], ha= 'center', va= 'center', color= 'red')  
plt.show()
```



```
In [10]: ## Error for the prediction for test dataset outcomes  
ftest_error = (ftest_cm[0,1] + ftest_cm[1,0])/np.sum(ftest_cm)  
print(ftest_error)  
  
0.14074995557135242
```

```
In [11]: ## Accuracy of prediction  
1-ftest_error
```

```
Out[11]: 0.8592500444286476
```

```
In [12]: ## Sensitivity Analysis  
ftest_sens = ftest_cm[1, 1]/(ftest_cm[1, 1] + ftest_cm[0, 1])  
print(ftest_sens)  
  
0.9599823451522731
```

```
In [13]: ## Specificity Analysis
ftest_spec = ftest_cm[0, 0]/(ftest_cm[0, 0]+ ftest_cm[1, 0])
print(ftest_spec)

0.7913526378421262
```

```
In [14]: ## PPV Analysis
ftest_npv = ftest_cm[1, 1]/(ftest_cm[1, 1] + ftest_cm[1, 0])
print(ftest_npv)

0.7561710511067331
```

```
In [15]: ## NPV Analysis
ftest_npv = ftest_cm[0, 0]/(ftest_cm[0, 0] + ftest_cm[0, 1])
print(ftest_npv)

0.9670382937469705
```

```
In [16]: ## The AUC Score
ftest_auc = roc_auc_score(fy_test, tns_probs)
fy_pred_auc = np.round(roc_auc_score(fy_test, fy_pred), decimals = 2)
```

```
In [17]: ## calculate ROC Curves
ftest_fpr, ftest_tpr, _ = roc_curve(fy_test, tns_probs)
fy_pred_fpr, fy_pred_tpr, _ = roc_curve(fy_test, fy_pred)
```

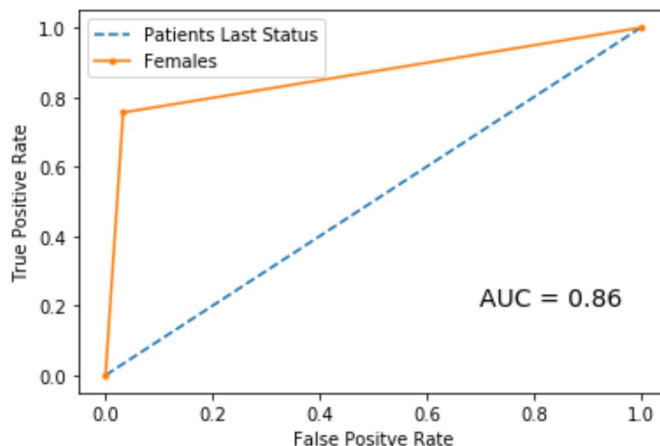
```
In [18]: ## Plot Curve for the model
import numpy as np
import matplotlib.pyplot as plt

plt.plot(ftest_fpr, ftest_tpr, linestyle = '--', label = 'Patients Last Status')
plt.plot(fy_pred_fpr, fy_pred_tpr, marker = '.', label = 'Females')
plt.text(0.7, 0.2, "AUC = " + str(fy_pred_auc), fontsize = 14)

## Axis lable
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")

## Show Legend
plt.legend()
```

Out[18]: <matplotlib.legend.Legend at 0x2e3411fc5c8>



In []:

In []:

In []: