

National University of Modern Languages

Artificial Intelligence - Lab Assignment # 2

BSSE - 5 - Morning

Submitted By:

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Submitted To:

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TASK 1

Implement the decision Tree classifier to detect breast cancer.

Code:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
data = pd.read csv(r'E:\NUML\Semester Data\Semester 5\AI\AI Lab\12093 Muhammad Um
air A2\dataR2.csv')
print(data.head())
print("\nCancer data set dimensions : {}".format(data.shape))
x = data.drop('Classification', 'columns')
y = data['Classification']
x train, x test, y train, y test = train test split(x, y, test size=0.25, random state=5)
scaler = StandardScaler()
scaler.fit(x train)
x train=scaler.transform(x train)
x test=scaler.transform(x test)
from sklearn.metrics import confusion matrix, classification report, accuracy score
from sklearn.tree import DecisionTreeClassifier
classifier = DecisionTreeClassifier(criterion= 'entropy', random state=0)
classifier.fit(x train, y train)
y pred dt = classifier.predict(x test)
print('Accuracy: ',accuracy score(y test, y pred dt))
cm = confusion matrix(y test, y pred dt)
print('Confusion Matrix: \n',cm)
```

```
plt.title("Heatmap of Confusion Matrix", fontsize = 15)
sns.heatmap(cm, annot= True)
plt.show()
print(classification_report(y_test, y_pred_dt))
```

Output:

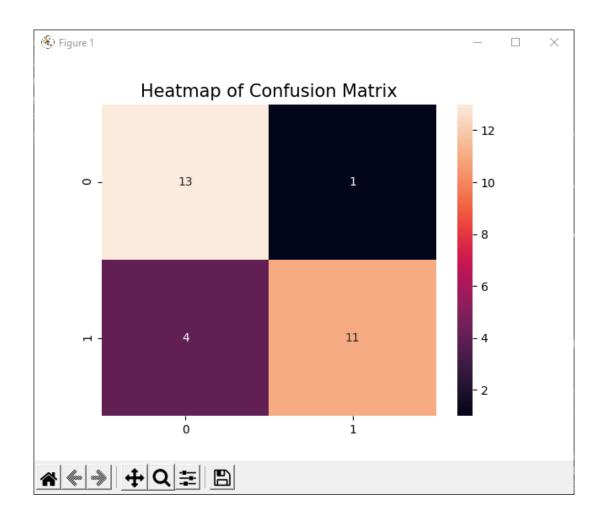
```
PS E:\NUML\Semester Data\Semester 5\AI\AI Lab\12093_Muhammad Umair_A2> & C:/Users/muham/AppData/Local/Programs/Python/Python39/python.exe "e:/NUML/Se
                                                                     Leptin Adiponectin Resistin
8.8071 9.702400 7.99585
8.8438 5.429285 4.06405
                                                           HOMA
                                                                                                                     MCP.1 Classification
                    BMI Glucose Insulin
                                           3.115 0.766897 8.8438
4.498 1.009651 17.9393
3.226 0.612725 9.8827
3.549 0.805386 6.6994
                                  70
92
91
77
92
                                                                                                                    468.786

    22.432040
    9.27715
    554.697

    7.169560
    12.76600
    928.220

    4.819240
    10.57635
    773.920

          21.367521
21.111111
Cancer data set dimensions : (116, 10)
Accuracy: 0.8275862068965517
Confusion Matrix:
 [[13 1]
[ 4 11]]
                    precision
                                       recall f1-score
                                                         0.81
                                                         0.83
0.83
      accuracy
weighted avg
                            0.84
                                          0.83
                                                         0.83
PS E:\NUML\Semester Data\Semester 5\AI\AI Lab\12093_Muhammad Umair_A2> []
```



TASK 2

Implement the logistic regression to detect breast cancer. Note: Standardize the data before training the model

Code: import numpy as np import pandas as pd import matplotlib.pyplot as plt import seaborn as sns from sklearn.model_selection import train_test_split from sklearn.preprocessing import StandardScaler data = pd.read csv(r'E:\NUML\Semester Data\Semester 5\AI\AI Lab\12093 Muhammad Um air A2\dataR2.csv') print(data.head()) print("\nCancer data set dimensions : {}".format(data.shape)) x = data.drop('Classification', 'columns') y = data['Classification'] x train, x test, y train, y test = train test split(x, y, test size=0.25, random state=5) scaler = StandardScaler() scaler.fit(x train) x train=scaler.transform(x train) x test=scaler.transform(x test) from sklearn.metrics import confusion matrix, classification report, accuracy score from sklearn.linear model import LogisticRegression classifier = LogisticRegression(random state = 0) classifier.fit(x train, y train) y pred lr = classifier.predict(x test) print('Accuracy: ',accuracy score(y test, y pred lr))

cm = confusion matrix(y test, y pred lr)

```
print('Confusion Matrix: \n',cm)
plt.title("Heatmap of Confusion Matrix", fontsize = 15)
sns.heatmap(cm, annot= True)
plt.show()
print(classification_report(y_test, y_pred_lr))
```

Output:

```
PS E:\NUML\Semester Data\Semester 5\AI\AI Lab\12093_Muhammad Umair_A2> & C:/Users/muham/AppData/Local/Programs/Python/Python39/python.exe "e:/NUML/Sem
                BMI Glucose Insulin
00000 70 2.707
90495 92 3.115
24670 91 4.498
                                     Leptin Adiponectin Resistin
8.8071 9.702400 7.99585
                                                                                                        MCP.1 Classification
417.114 1
                                                                             5.429285
22.432040
                                                                                            4.06405
9.27715
        20.690495
23.124670
                                                                                                         468.786
554.697
                                       3.226 0.612725
3.549 0.805386
                                                               9.8827
6.6994
                                                                              7.169560 12.76600
4.819240 10.57635
                                                                                                         928.220
773.920
Cancer data set dimensions : (116, 10)
Accuracy: 0.7931034482758621
Confusion Matrix:
[[10 4]
[ 2 13]]
                  precision
                                   recall f1-score
                                                                    14
15
                                                                    29
                                                                    29
29
PS E:\NUML\Semester Data\Semester 5\AI\AI Lab\12093_Muhammad Umair_A2> []
```

