## practical-06

## April 26, 2024

```
[23]: import pandas as pd
      import matplotlib.pyplot as plt
      from sklearn.preprocessing import LabelEncoder
      from sklearn.model_selection import train_test_split
      from sklearn.naive_bayes import GaussianNB
      from sklearn import metrics
      from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay
 [2]: data = pd.read_csv("C:/Users/gugal/Desktop/THIRD 2/PRACTICALS/DS/CODES/DATASETS/
       ⇔iris.csv")
      data
 [2]:
           sepal_length sepal_width petal_length petal_width
                                                                     species
      0
                    5.1
                                  3.5
                                                1.4
                                                              0.2
                                                                      setosa
      1
                    4.9
                                  3.0
                                                1.4
                                                              0.2
                                                                      setosa
                    4.7
                                                              0.2
      2
                                  3.2
                                                1.3
                                                                      setosa
      3
                    4.6
                                  3.1
                                                1.5
                                                              0.2
                                                                      setosa
      4
                    5.0
                                  3.6
                                                1.4
                                                              0.2
                                                                      setosa
      . .
      145
                    6.7
                                  3.0
                                                5.2
                                                              2.3 virginica
                    6.3
                                  2.5
                                                5.0
      146
                                                              1.9 virginica
      147
                    6.5
                                  3.0
                                                5.2
                                                              2.0 virginica
      148
                                  3.4
                                                5.4
                    6.2
                                                              2.3 virginica
      149
                    5.9
                                  3.0
                                                5.1
                                                              1.8 virginica
      [150 rows x 5 columns]
 [3]: data.isnull().sum()
 [3]: sepal_length
                      0
      sepal_width
                      0
      petal_length
      petal_width
                      0
      species
                      0
      dtype: int64
 [4]: data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
     RangeIndex: 150 entries, 0 to 149
     Data columns (total 5 columns):
          Column
                         Non-Null Count Dtype
                         _____
          sepal_length 150 non-null
                                          float64
      0
      1
          sepal width
                         150 non-null
                                          float64
                                          float64
          petal_length 150 non-null
      3
          petal width
                         150 non-null
                                          float64
                         150 non-null
      4
          species
                                          object
     dtypes: float64(4), object(1)
     memory usage: 6.0+ KB
 [5]: data.columns
 [5]: Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
             'species'],
            dtype='object')
 [6]: data.describe()
 [6]:
             sepal_length
                            sepal_width petal_length petal_width
               150.000000
                             150.000000
                                           150.000000
                                                         150.000000
      count
                 5.843333
                               3.054000
                                             3.758667
                                                           1.198667
      mean
      std
                 0.828066
                               0.433594
                                              1.764420
                                                           0.763161
      min
                 4.300000
                               2.000000
                                             1.000000
                                                           0.100000
      25%
                 5.100000
                               2.800000
                                              1.600000
                                                           0.300000
      50%
                 5.800000
                               3.000000
                                             4.350000
                                                           1.300000
      75%
                 6.400000
                               3.300000
                                             5.100000
                                                           1.800000
                 7.900000
                               4.400000
                                             6.900000
      max
                                                           2.500000
 [9]: le = LabelEncoder()
      data['species'] = le.fit_transform(data['species'])
      data['species'].unique()
 [9]: array([0, 1, 2])
[10]:
     data
[10]:
           sepal_length
                         sepal_width petal_length petal_width species
      0
                    5.1
                                  3.5
                                                 1.4
                                                              0.2
                                                                          0
      1
                    4.9
                                  3.0
                                                 1.4
                                                              0.2
                                                                          0
                                  3.2
                                                              0.2
      2
                    4.7
                                                 1.3
                                                                          0
                                                              0.2
      3
                    4.6
                                  3.1
                                                 1.5
                                                                          0
                    5.0
                                                              0.2
                                                                          0
      4
                                  3.6
                                                 1.4
      145
                    6.7
                                  3.0
                                                 5.2
                                                              2.3
                                                                          2
```

```
2.5
146
                6.3
                                                5.0
                                                               1.9
                                                                            2
                                                                            2
147
                6.5
                               3.0
                                                5.2
                                                               2.0
148
                6.2
                               3.4
                                                5.4
                                                               2.3
                                                                            2
                                                                            2
149
                5.9
                               3.0
                                                5.1
                                                               1.8
```

[150 rows x 5 columns]

```
[11]: X = data[['sepal_length', 'sepal_width', 'petal_length', 'petal_width']].values
X
```

```
[11]: array([[5.1, 3.5, 1.4, 0.2],
             [4.9, 3., 1.4, 0.2],
             [4.7, 3.2, 1.3, 0.2],
             [4.6, 3.1, 1.5, 0.2],
             [5., 3.6, 1.4, 0.2],
             [5.4, 3.9, 1.7, 0.4],
             [4.6, 3.4, 1.4, 0.3],
             [5., 3.4, 1.5, 0.2],
             [4.4, 2.9, 1.4, 0.2],
             [4.9, 3.1, 1.5, 0.1],
             [5.4, 3.7, 1.5, 0.2],
             [4.8, 3.4, 1.6, 0.2],
             [4.8, 3., 1.4, 0.1],
             [4.3, 3., 1.1, 0.1],
             [5.8, 4., 1.2, 0.2],
             [5.7, 4.4, 1.5, 0.4],
             [5.4, 3.9, 1.3, 0.4],
             [5.1, 3.5, 1.4, 0.3],
             [5.7, 3.8, 1.7, 0.3],
             [5.1, 3.8, 1.5, 0.3],
             [5.4, 3.4, 1.7, 0.2],
             [5.1, 3.7, 1.5, 0.4],
             [4.6, 3.6, 1., 0.2],
             [5.1, 3.3, 1.7, 0.5],
             [4.8, 3.4, 1.9, 0.2],
             [5., 3., 1.6, 0.2],
             [5., 3.4, 1.6, 0.4],
             [5.2, 3.5, 1.5, 0.2],
             [5.2, 3.4, 1.4, 0.2],
             [4.7, 3.2, 1.6, 0.2],
             [4.8, 3.1, 1.6, 0.2],
             [5.4, 3.4, 1.5, 0.4],
             [5.2, 4.1, 1.5, 0.1],
             [5.5, 4.2, 1.4, 0.2],
             [4.9, 3.1, 1.5, 0.1],
             [5., 3.2, 1.2, 0.2],
             [5.5, 3.5, 1.3, 0.2],
```

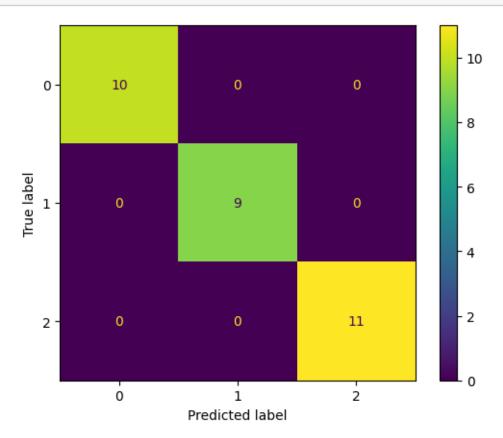
```
[4.9, 3.1, 1.5, 0.1],
[4.4, 3., 1.3, 0.2],
[5.1, 3.4, 1.5, 0.2],
[5., 3.5, 1.3, 0.3],
[4.5, 2.3, 1.3, 0.3],
[4.4, 3.2, 1.3, 0.2],
[5., 3.5, 1.6, 0.6],
[5.1, 3.8, 1.9, 0.4],
[4.8, 3., 1.4, 0.3],
[5.1, 3.8, 1.6, 0.2],
[4.6, 3.2, 1.4, 0.2],
[5.3, 3.7, 1.5, 0.2],
[5., 3.3, 1.4, 0.2],
[7., 3.2, 4.7, 1.4],
[6.4, 3.2, 4.5, 1.5],
[6.9, 3.1, 4.9, 1.5],
[5.5, 2.3, 4., 1.3],
[6.5, 2.8, 4.6, 1.5],
[5.7, 2.8, 4.5, 1.3],
[6.3, 3.3, 4.7, 1.6],
[4.9, 2.4, 3.3, 1.],
[6.6, 2.9, 4.6, 1.3],
[5.2, 2.7, 3.9, 1.4],
[5., 2., 3.5, 1.],
[5.9, 3., 4.2, 1.5],
[6., 2.2, 4., 1.],
[6.1, 2.9, 4.7, 1.4],
[5.6, 2.9, 3.6, 1.3],
[6.7, 3.1, 4.4, 1.4],
[5.6, 3., 4.5, 1.5],
[5.8, 2.7, 4.1, 1.],
[6.2, 2.2, 4.5, 1.5],
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[6.1, 2.8, 4., 1.3],
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[6.4, 2.9, 4.3, 1.3],
[6.6, 3., 4.4, 1.4],
[6.8, 2.8, 4.8, 1.4],
[6.7, 3., 5., 1.7],
[6., 2.9, 4.5, 1.5],
[5.7, 2.6, 3.5, 1.],
[5.5, 2.4, 3.8, 1.1],
[5.5, 2.4, 3.7, 1.],
[5.8, 2.7, 3.9, 1.2],
[6., 2.7, 5.1, 1.6],
```

[5.4, 3., 4.5, 1.5],[6., 3.4, 4.5, 1.6],[6.7, 3.1, 4.7, 1.5],[6.3, 2.3, 4.4, 1.3],[5.6, 3., 4.1, 1.3],[5.5, 2.5, 4., 1.3],[5.5, 2.6, 4.4, 1.2],[6.1, 3., 4.6, 1.4],[5.8, 2.6, 4., 1.2],[5., 2.3, 3.3, 1.],[5.6, 2.7, 4.2, 1.3], [5.7, 3., 4.2, 1.2],[5.7, 2.9, 4.2, 1.3], [6.2, 2.9, 4.3, 1.3],[5.1, 2.5, 3., 1.1],[5.7, 2.8, 4.1, 1.3],[6.3, 3.3, 6., 2.5],[5.8, 2.7, 5.1, 1.9], [7.1, 3., 5.9, 2.1],[6.3, 2.9, 5.6, 1.8],[6.5, 3., 5.8, 2.2],[7.6, 3., 6.6, 2.1],[4.9, 2.5, 4.5, 1.7],[7.3, 2.9, 6.3, 1.8],[6.7, 2.5, 5.8, 1.8],[7.2, 3.6, 6.1, 2.5],[6.5, 3.2, 5.1, 2.],[6.4, 2.7, 5.3, 1.9],[6.8, 3., 5.5, 2.1],[5.7, 2.5, 5., 2.],[5.8, 2.8, 5.1, 2.4],[6.4, 3.2, 5.3, 2.3],[6.5, 3., 5.5, 1.8],[7.7, 3.8, 6.7, 2.2],[7.7, 2.6, 6.9, 2.3],[6., 2.2, 5., 1.5],[6.9, 3.2, 5.7, 2.3],[5.6, 2.8, 4.9, 2.], [7.7, 2.8, 6.7, 2.],[6.3, 2.7, 4.9, 1.8],[6.7, 3.3, 5.7, 2.1],[7.2, 3.2, 6., 1.8],[6.2, 2.8, 4.8, 1.8],[6.1, 3., 4.9, 1.8],[6.4, 2.8, 5.6, 2.1],[7.2, 3., 5.8, 1.6],[7.4, 2.8, 6.1, 1.9],

```
[6.4, 2.8, 5.6, 2.2],
         [6.3, 2.8, 5.1, 1.5],
         [6.1, 2.6, 5.6, 1.4],
         [7.7, 3., 6.1, 2.3],
         [6.3, 3.4, 5.6, 2.4],
         [6.4, 3.1, 5.5, 1.8],
         [6., 3., 4.8, 1.8],
         [6.9, 3.1, 5.4, 2.1],
         [6.7, 3.1, 5.6, 2.4],
         [6.9, 3.1, 5.1, 2.3],
         [5.8, 2.7, 5.1, 1.9],
         [6.8, 3.2, 5.9, 2.3],
         [6.7, 3.3, 5.7, 2.5],
         [6.7, 3., 5.2, 2.3],
         [6.3, 2.5, 5., 1.9],
         [6.5, 3., 5.2, 2.],
         [6.2, 3.4, 5.4, 2.3],
         [5.9, 3., 5.1, 1.8]
[12]: Y = data['species'].values
    Y
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
        [15]: X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.2,__
     →random_state=42)
[17]: print(len(Y_test))
   30
[18]: model = GaussianNB()
    model.fit(X_train,Y_train)
[18]: GaussianNB()
[20]: prediction = model.predict(X_test)
    print('Accuracy is', metrics.accuracy_score(Y_test,prediction))
   Accuracy is 1.0
```

[7.9, 3.8, 6.4, 2.],

```
[24]: disp = ConfusionMatrixDisplay(confusion_matrix=CM)
    disp.plot()
    plt.show()
```



```
[25]: TN = CM[0, 0]
FP = CM[0, 1]
FN = CM[1, 0]
TP = CM[1, 1]

print("True Negative (TN):", TN)
print("False Positive (FP):", FP)
print("False Negative (FN):", FN)
print("True Positive (TP):", TP)
```

```
True Negative (TN): 10
     False Positive (FP): 0
     False Negative (FN): 0
     True Positive (TP): 9
[26]: acc= (TP + TN)/(TP+FP+TN+FN)
      acc
[26]: 1.0
[27]: # Error Rate
      Error_Rate = (FP + FN)/(TP+FP+TN+FN)
      Error_Rate
[27]: 0.0
[28]: # Precision
      Precision = (TP)/(TP+FP)
      Precision
[28]: 1.0
[29]: # Recall
      Recall = (TP)/(TP+FN)
      Recall
```