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April 15, 2025

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[1]: import numpy as np
     import matplotlib.pyplot as plt
     import matplotlib.image as mpimg
     import pandas as pd
[2]: kd = pd.read_csv('iris.csv')
[3]: kd.head()
[3]:
        sepal_length sepal_width petal_length petal_width species
     0
                 5.1
                              3.5
                                            1.4
                                                         0.2 setosa
                 4.9
     1
                              3.0
                                            1.4
                                                         0.2 setosa
                 4.7
                                            1.3
     2
                              3.2
                                                         0.2 setosa
                                                         0.2 setosa
     3
                 4.6
                              3.1
                                            1.5
                                                         0.2 setosa
                 5.0
                              3.6
                                            1.4
[4]: %matplotlib inline
[5]: img=mpimg.imread('iris_types.jpg')
[6]: plt.figure(figsize=(20,40))
     plt.axis('off')
     plt.imshow(img)
```

[6]: <matplotlib.image.AxesImage at 0x136bbfed050>



```
[7]: X = kd.iloc[:,:4].values
      y = kd['species'].values
 [8]: from sklearn.model_selection import train_test_split
      X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20, __
       →random_state = 82)
 [9]: from sklearn.preprocessing import StandardScaler
      sc = StandardScaler()
      X_train = sc.fit_transform(X_train)
      X_test = sc.transform(X_test)
[10]: from sklearn.naive_bayes import GaussianNB
      nvclassifier = GaussianNB()
      nvclassifier.fit(X_train, y_train)
[10]: GaussianNB()
[11]: y_pred = nvclassifier.predict(X_test)
      print(y_pred)
     ['Iris-virginica' 'Iris-virginica' 'setosa' 'setosa' 'setosa'
      'Iris-virginica' 'Iris-versicolor' 'Iris-versicolor' 'Iris-versicolor'
      'Iris-versicolor' 'Iris-versicolor' 'Iris-virginica' 'setosa' 'setosa'
      'setosa' 'setosa' 'Iris-virginica' 'Iris-versicolor' 'setosa'
      'Iris-versicolor' 'setosa' 'Iris-virginica' 'setosa' 'Iris-virginica'
      'Iris-virginica' 'Iris-versicolor' 'Iris-virginica' 'setosa'
      'Iris-virginica' 'Iris-versicolor']
[12]: y_compare = np.vstack((y_test,y_pred)).T
[13]: y_compare[:5,:]
[13]: array([['Iris-virginica', 'Iris-virginica'],
             ['Iris-virginica', 'Iris-virginica'],
             ['setosa', 'setosa'],
             ['setosa', 'setosa'],
             ['setosa', 'setosa']], dtype=object)
[14]: from sklearn.metrics import confusion_matrix
      cm = confusion_matrix(y_test, y_pred)
      print(cm)
     [[8 1 0]
      [1 9 0]
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[ 0 0 11]]
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[15]: a = cm.shape
     corrPred = 0
     falsePred = 0
[21]: corrPred = 0
     falsePred = 0
     for row in range(a[0]):
        for c in range(a[1]):
            if row == c:
               corrPred += cm[row, c]
            else:
               falsePred += cm[row, c]
     print('Correct predictions:', corrPred)
     print('False predictions:', falsePred)
     print('\n\nAccuracy of the Naive Bayes classification is:', corrPred / cm.sum())
    Correct predictions: 28
    False predictions: 2
    []:
[]:
```