Practical 4 PART A

B-76

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```
import numpy as np
import pandas as pd
import scipy.stats as stats
import matplotlib.pyplot as plt
import seaborn as sns
sns.set(style="whitegrid")
# i. Plot the density plot for each feature
df = pd.read csv("data/iris.csv")
df = df.sample(frac=1, random_state=1).reset_index(drop=True)
print(df.shape)
df1 = pd.read_csv("data/iris_train.csv")
df2 = pd.read_csv("data/iris_test.csv")
X1, y1 = df1.iloc[:, :-1], df1.iloc[:, -1]
X2, y2 = df2.iloc[:, :-1], df2.iloc[:, -1]
X_train, y_train = X1, y1
X_{\text{test}}, y_{\text{test}} = X2, y2
# print(X train.shape, y train.shape)
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si=d+i| Sepai.Lengtn |
sl.plot.density(color='green')
plt.title('Density plot for Sepal.Length')
plt.show()
sl=df1['Sepal.Width']
sl.plot.density(color='green')
plt.title('Density plot for Sepal.width')
plt.show()
sl=df1['Petal.Length']
sl.plot.density(color='green')
plt.title('Density plot for Petal.Length')
plt.show()
sl=df1['Petal.Width']
sl.plot.density(color='green')
plt.title('Density plot for Petal.Width')
plt.show()
```

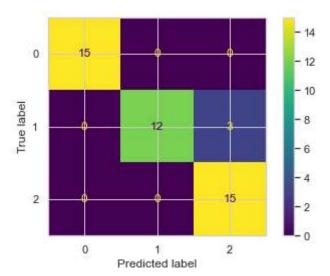
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(150, 5)
                     Density plot for Sepal.Length
# ii. Plot the confusion matrix and compute sensitivity, specificity, accuracy for your pr
         X train.shape
     (105, 4)
         x = NaiveBayesClassifier()
                            # FUNCTION CREATED IN LAST CELL
                              # - OF THIS NOTEBOOK
x.fit(X_train, y_train)
                                                                                  x.classes, x.feature nums, x.rows, x.count
     (array(['setosa', 'versicolor', 'virginica'], dtype=object), 4, 105, 3)
                  import matplotlib.pyplot as plt
import numpy
from sklearn import metrics
import copy
confusion_matrix = metrics.confusion_matrix(list(y_test), list(predictions))
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix)
cm display.plot()
plt.show()
a=copy.deepcopy(confusion_matrix)
print("Accuracy: ",metrics.accuracy_score(y_test, predictions))
FN=[a[0][1]+a[0][2],a[1][0]+a[1][2],a[2][0]+a[2][1]]
                                 a[0][2]+a[1][2]]
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Specificity1 = (a[1][1]+a[2][2])/(a[1][0]+a[2][0]+a[1][1]+a[2][2])
Specificity2 = (a[0][0]+a[2][2])/(a[0][1]+a[2][1]+a[0][0]+a[2][2])
Specificity3 = (a[0][0]+a[1][1])/(a[0][2]+a[1][2]+a[0][0]+a[1][1])
print(f"\nSpecificity:\nClass 01 {Specificity1*100}% \nClass 02 {Specificity2*100}% \nClas
# Sensitivity
Sensitivity1 = a[0][0]/(a[0][0]+a[0][1]+a[0][2])
Sensitivity2 = a[1][1]/(a[1][0]+a[1][1]+a[1][2])
Sensitivity3 = a[2][2]/(a[2][0]+a[2][1]+a[2][2])
print(f"\nSensitivity: \nClass 01 {Sensitivity1*100} \nClass 02 {Sensitivity2*100}% \nClas
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Specificity: Class 01 100.0% Class 02 100.0% Class 03 90.0%

iii) Compute class wise probabilities and plot the histogram of predicted results.

array([0.33333333, 0.33333333, 0.33333333])

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x.accuracy(y_test, predictions)
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0.9333333333333333

y_test.value_counts(normalize=True)

setosa 0.333333 versicolor 0.333333 virginica 0.333333

Name: Species, dtype: float64

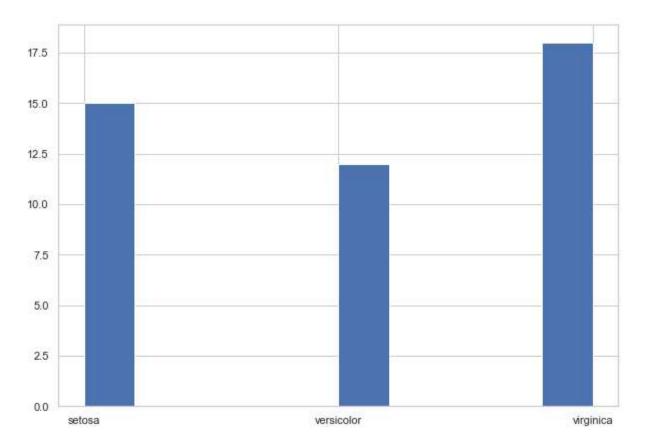
from matplotlib import pyplot as plt
import numpy as np

Creating histogram
fig, ax = plt.subplots(figsize =(10, 7))

ax.hist(predictions)

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```
# Show plot
plt.show()
```



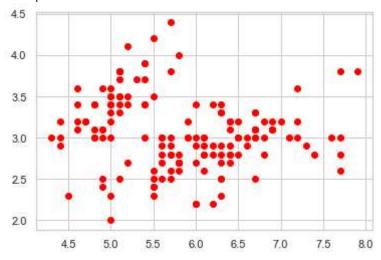
iv. Plot scatter plot showing the decision boundary for Sepal length vs. Sepal width for # all species.

```
class NaiveBayesClassifier():
    def calc_prior(self, features, target):
        self.prior = (features.groupby(target).apply(lambda x: len(x)) / self.rows).to num
        return self.prior
                                    es, target):
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                                    (target).apply(np.mean).to_numpy()
        self.var = features.groupby(target).apply(np.var).to_numpy()
        return self.mean, self.var
   def gaussian_density(self, class_idx, x):
        mean = self.mean[class_idx]
        var = self.var[class idx]
        numerator = np.exp((-1/2)*((x-mean)**2) / (2 * var))
        denominator = np.sqrt(2 * np.pi * var)
        prob = numerator / denominator
        return prob
   def calc_posterior(self, x):
        posteriors = []
        for i in range(self.count):
            prior = np.log(self.prior[i])
            conditional = np.sum(np.log(self.gaussian_density(i, x)))
            posterior = prior + conditional
            posteriors.append(posterior)
        return self.classes[np.argmax(posteriors)]
```

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```
def fit(self, features, target):
        self.classes = np.unique(target)
        self.count = len(self.classes)
        self.feature nums = features.shape[1]
        self.rows = features.shape[0]
        self.calc_statistics(features, target)
        self.calc prior(features, target)
   def predict(self, features):
        preds = [self.calc posterior(f) for f in features.to numpy()]
        return preds
   def accuracy(self, y_test, y_pred):
        accuracy = np.sum(y_test == y_pred) / len(y_test)
        return accuracy
   def visualize(self, y_true, y_pred, target):
        tr = pd.DataFrame(data=y_true, columns=[target])
        pr = pd.DataFrame(data=y_pred, columns=[target])
        fig, ax = plt.subplots(1, 2, sharex='col', sharey='row', figsize=(15,6))
        sns.countplot(x=target, data=tr, ax=ax[0], palette='viridis', alpha=0.7, hue=targe
        sns.countplot(x=target, data=pr, ax=ax[1], palette='viridis', alpha=0.7, hue=targe
        fig.suptitle('True vs Predicted Comparison', fontsize=20)
        ax[0].tick_params(labelsize=12)
        ax[1].tick_params(labelsize=12)
        ax[0].set_title("True values", fontsize=18)
        ax[1].set_title("Predicted values", fontsize=18)
        plt.show()
df['variety'] = pd.Series(df.variety)
df.head()
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<matplotlib.collections.PathCollection at 0x1a1ed6fc580>



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