Assignment2_NaiveBayes

June 10, 2021

0.1 Part B

1 Libraries

```
[1]: from sklearn.datasets import load_iris
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
#Import Gaussian Naive Bayes model
from sklearn.naive_bayes import GaussianNB
```

2 1) Load the Iris dataset

[5 rows x 5 columns]

```
[2]: # Load the Iris dataset
    iris = load_iris()
    df = pd.DataFrame(iris['data'], columns=iris['feature_names'])
    df['Species'] = iris['target']
    df.head()
       sepal length (cm) sepal width (cm) ... petal width (cm)
                                                                   Species
[2]:
    0
                     5.1
                                       3.5 ...
                                                              0.2
                                       3.0 ...
                     4.9
                                                              0.2
                                                                         0
    1
                                       3.2 ...
    2
                     4.7
                                                              0.2
                                                                         0
    3
                                       3.1 ...
                                                              0.2
                     4.6
                                                                         0
                                       3.6 ...
                                                              0.2
                     5.0
                                                                         0
```

3 2) Drop the petal length and petal width features to form a 2D Iris dataset

```
[3]: iris_2d = df.drop(['petal length (cm)', 'petal width (cm)'] , axis = 1) iris_2d.head()
```

```
[3]:
       sepal length (cm) sepal width (cm) Species
    0
                                          3.5
                                          3.0
                      4.9
                                                      0
    1
    2
                      4.7
                                          3.2
                                                      0
    3
                      4.6
                                          3.1
                                                      0
    4
                      5.0
                                          3.6
```

```
[4]: #function for calculating 1D dataset likelihoods
def likelihood_1d(mean , var , x):
    mean = mean
    var = var
    numerator = np.exp((-1/2)*((x-mean)**2) / (2 * var))
    denominator = np.sqrt(2 * np.pi * var)
    prob = numerator / denominator
    return prob
```

4 3) Model I of iris dataset

- 1. applying gaussian naive bayes on 2D iris dataset.
- 2. plotting likelihood of sepal length feature only.
- 3. plotting posterior of 2D dataset.
- 4. Calculating the Accuracy.

1. Applying Gaussian Naive Bayes

```
[5]: #Create a Gaussian Classifier
model1 = GaussianNB()

# Train the model using the training sets
model1.fit(iris_2d.iloc[:,:2],iris_2d.iloc[:,-1])
```

[5]: GaussianNB(priors=None, var_smoothing=1e-09)

plt.ylabel('Naive Bayes Likelihood')

2. Plotting the likelihood of sepal length only.

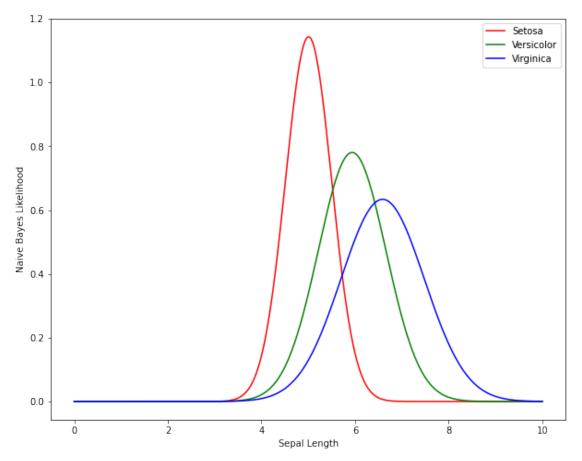
```
[6]: #taking the variance and standard deviation of first feature only.
var_1d = model1.sigma_[:,0]
mean_1d = model1.theta_[:,0]

[7]: x = np.arange(0 , 10 ,0.001)
CO_likelihood = likelihood_1d(mean_1d[0],var_1d[0],x)
C1_likelihood = likelihood_1d(mean_1d[1],var_1d[1],x)
C2_likelihood = likelihood_1d(mean_1d[2],var_1d[2],x)

plt.figure(figsize=(10,8))
plt.plot(x , C0_likelihood , label = 'Setosa' , color = 'r')
plt.plot(x , C1_likelihood , label = 'Versicolor' , color = 'g')
```

plt.plot(x , C2_likelihood , label = 'Virginica' , color = 'b')

```
plt.xlabel('Sepal Length')
plt.legend()
plt.show()
```



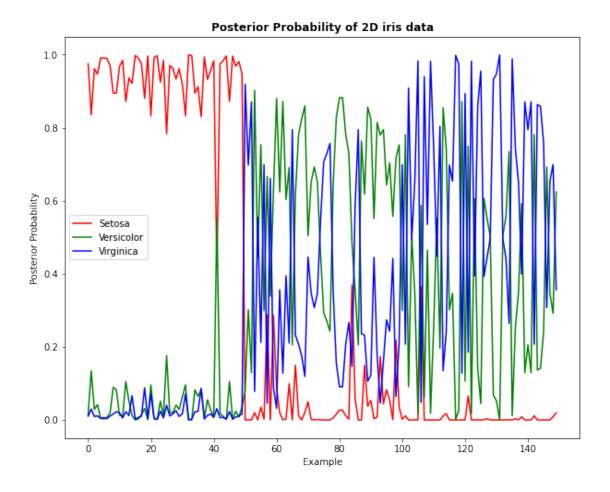
3. plotting posterior of 2D dataset.

```
[8]: prob2 = model1.predict_proba(iris_2d.iloc[:,:2])

x = np.arange(0,iris_2d.shape[0])

plt.figure(figsize=(10,8))
plt.plot(x , prob2[:,0] , label = 'Setosa' , color = 'r')
plt.plot(x , prob2[:,1] , label = 'Versicolor' , color = 'g')
plt.plot(x , prob2[:,2] , label = 'Virginica' , color = 'b')
plt.title('Posterior Probability of 2D iris data' , fontweight ="bold")
plt.xlabel('Example')
plt.ylabel('Posterior Probability')

plt.legend()
plt.show()
```



4. Calculating the Accuracy.

```
[9]: accuracy = model1.score(iris_2d.iloc[:,:2],iris_2d.iloc[:,-1])
print(f'Accuracy of the model = {accuracy * 100} %\n')
```

Accuracy of the model = 78.0 %

5 4)1) Model II of iris dataset

mu1 = mu2 = mu3 = 5.5 and keep the actual values of sigma1, sigma2, and sigma3 for the first feature.

- 1. applying gaussian naive bayes on 2D iris dataset.
- 2. plotting likelihood of sepal length feature only.
- 3. plotting posterior of 2D dataset.
- 4. Calculating the Accuracy.

1. Applying Gaussian Naive Bayes

```
[10]: #Create a Gaussian Classifier
model2 = GaussianNB()

# Train the model using the training sets
model2.fit(iris_2d.iloc[:,:2],iris_2d.iloc[:,-1])
model2.theta_[:,0] = [5.5,5.5,5.5]
print(f'New means of the model = \n {model2.theta_}')
```

```
New means of the model = [[5.5 3.428] [5.5 2.77] [5.5 2.974]]
```

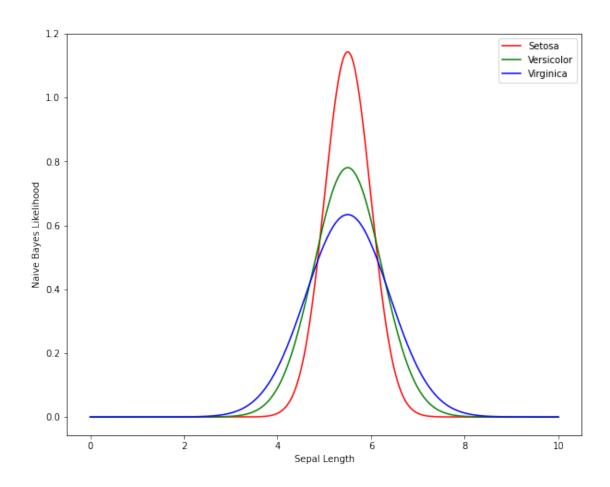
2. Plotting the likelihood of sepal length only.

```
[11]: var_1d = model2.sigma_[:,0]
    mean_1d = model2.theta_[:,0]

[12]: x = np.arange(0 , 10 ,0.001)
    C0_likelihood = likelihood_1d(mean_1d[0],var_1d[0],x)
    C1_likelihood = likelihood_1d(mean_1d[1],var_1d[1],x)
    C2_likelihood = likelihood_1d(mean_1d[2],var_1d[2],x)

plt.figure(figsize=(10,8))
    plt.plot(x , C0_likelihood , label = 'Setosa' , color = 'r')
    plt.plot(x , C1_likelihood , label = 'Versicolor' , color = 'g')
    plt.plot(x , C2_likelihood , label = 'Virginica' , color = 'b')
    plt.ylabel('Naive Bayes Likelihood')
    plt.xlabel('Sepal Length')

plt.legend()
    plt.show()
```



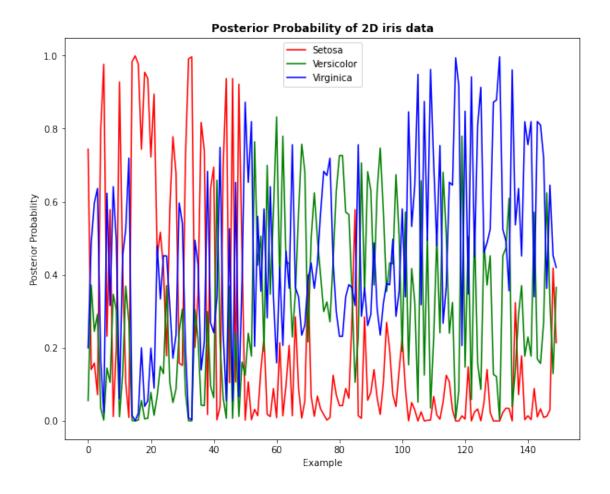
3. Plotting posterior of 2D dataset.

```
[13]: prob2 = model2.predict_proba(iris_2d.iloc[:,:2])

import matplotlib.pyplot as plt
    x = np.arange(0,iris_2d.shape[0])

plt.figure(figsize=(10,8))
    plt.plot(x , prob2[:,0] , label = 'Setosa' , color = 'r')
    plt.plot(x , prob2[:,1] , label = 'Versicolor' , color = 'g')
    plt.plot(x , prob2[:,2] , label = 'Virginica' , color = 'b')
    plt.title('Posterior Probability of 2D iris data' , fontweight ="bold")
    plt.xlabel('Example')
    plt.ylabel('Posterior Probability')

plt.legend()
    plt.show()
```



4. Calculating the Accuracy.

```
[14]: accuracy = model2.score(iris_2d.iloc[:,:2],iris_2d.iloc[:,-1])
print(f'Accuracy of the model = {accuracy * 100} %\n')
```

6 4)2) Model III of iris dataset

sigma1 = sigma2 = sigma3 = 0.26 and keep the actual values of mu1, mu2, and mu3 for the first feature.

- 1. applying gaussian naive bayes on 2D iris dataset.
- 2. plotting likelihood of sepal length feature only.
- 3. plotting posterior of 2D dataset.
- 4. Calculating the Accuracy.

1. Applying Gaussian Naive Bayes

```
[15]: #Create a Gaussian Classifier
model3 = GaussianNB()

# Train the model using the training sets
model3.fit(iris_2d.iloc[:,:2],iris_2d.iloc[:,-1])
model3.sigma_[:,0] = [0.26,0.26,0.26]
print(model3.sigma_)
```

```
[[0.26 0.140816]
[0.26 0.0965]
[0.26 0.101924]]
```

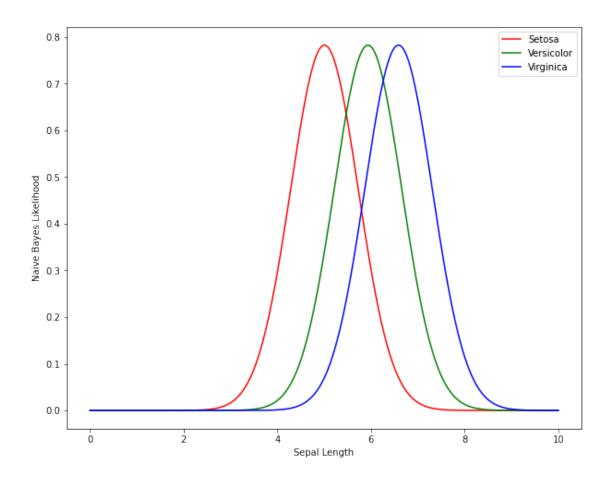
2. Plotting the likelihood of sepal length only.

```
[16]: var_1d = model3.sigma_[:,0]
    mean_1d = model3.theta_[:,0]

[17]: x = np.arange(0 , 10 ,0.001)
    C0_likelihood = likelihood_1d(mean_1d[0],var_1d[0],x)
    C1_likelihood = likelihood_1d(mean_1d[1],var_1d[1],x)
    C2_likelihood = likelihood_1d(mean_1d[2],var_1d[2],x)

plt.figure(figsize=(10,8))
    plt.plot(x , C0_likelihood , label = 'Setosa' , color = 'r')
    plt.plot(x , C1_likelihood , label = 'Versicolor' , color = 'g')
    plt.plot(x , C2_likelihood , label = 'Virginica' , color = 'b')
    plt.ylabel('Naive Bayes Likelihood')
    plt.xlabel('Sepal Length')

plt.legend()
    plt.show()
```



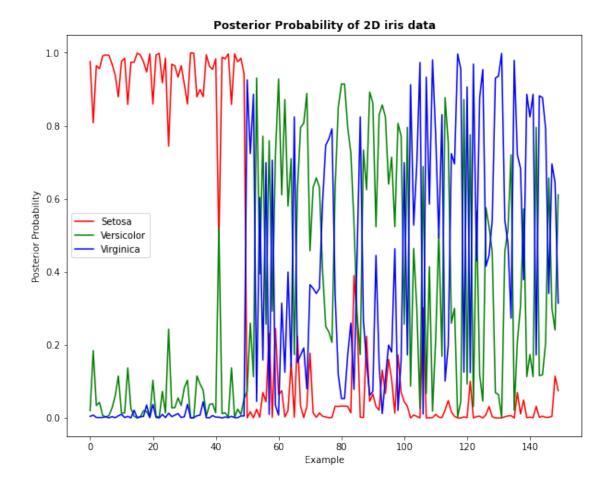
3. Plotting posterior of 2D dataset.

```
[18]: prob3 = model3.predict_proba(iris_2d.iloc[:,:2])

x = np.arange(0,iris_2d.shape[0])

plt.figure(figsize=(10,8))
 plt.plot(x , prob3[:,0] , label = 'Setosa' , color = 'r')
 plt.plot(x , prob3[:,1] , label = 'Versicolor' , color = 'g')
 plt.plot(x , prob3[:,2] , label = 'Virginica' , color = 'b')
 plt.title('Posterior Probability of 2D iris data' , fontweight ="bold")
 plt.xlabel('Example')
 plt.ylabel('Posterior Probability')

plt.legend()
 plt.show()
```



4. Calculating the Accuracy.

```
[19]: accuracy = model3.score(iris_2d.iloc[:,:2],iris_2d.iloc[:,-1])
print(f'Accuracy of the model = {accuracy * 100} %\n')
```

Accuracy of the model = 80.0 %

7 4)3) Comparing Accuracy & Comment

Model 1 accuracy = 78.0 % Model 2 accuracy = 62.67 % Model 3 accuracy = 80.0 % Conclusion:

1. when we fix the mean, the accuracy decrease because the data is centered around the same mean of each class, and the probability of setosa is the largest, so a large amount of data are misclassified due to the overlapping in the wide range around the mean of each class.

