```
In [ ]:
           import numpy as np
           import pandas as pd
           import matplotlib.pyplot as plt
In [ ]:
           import seaborn as sns
           df=sns.load_dataset("iris")
          df.head()
Out[ ]:
             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
          2
                     4.7
                                  3.2
                                               1.3
                                                            0.2
                                                                 setosa
         3
                     4.6
                                  3.1
                                               1.5
                                                            0.2
                                                                 setosa
                     5.0
                                                            0.2
          4
                                  3.6
                                               1.4
                                                                 setosa
In [ ]:
          \#design x and y
          X= df.iloc[:, :-1] #features
          y= df.iloc[ : , -1:] #labels
In [ ]:
          X.head()
Out[]:
             sepal_length sepal_width petal_length petal_width
         0
                     5.1
                                  3.5
                                               1.4
                                                            0.2
          1
                     4.9
                                  3.0
                                               1.4
                                                            0.2
         2
                     4.7
                                  3.2
                                               1.3
                                                            0.2
          3
                                                            0.2
                     4.6
                                  3.1
                                               1.5
                                                            0.2
          4
                     5.0
                                  3.6
                                               1.4
In [ ]:
          y.head()
Out[]:
             species
         0
              setosa
              setosa
              setosa
              setosa
              setosa
In [ ]:
          #train and fir the model
```

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```
from sklearn.naive bayes import GaussianNB
         model= GaussianNB()
         model.fit(X, y)
        C:\Users\Javeria\AppData\Local\Programs\Python\Python310\lib\site-packages\sklearn\utils
        \validation.py:993: DataConversionWarning: A column-vector y was passed when a 1d array
        was expected. Please change the shape of y to (n_samples, ), for example using ravel().
          y = column or 1d(y, warn=True)
        GaussianNB()
Out[]:
In [ ]:
         from sklearn.model_selection import train_test_split
         X_train, X_test, y_train, y_test= train_test_split(X, y, test_size= 0.3, random_state=0
In [ ]:
         # Predicting the Test set results
         y prediction = model.predict(X test)
         y prediction
        array(['virginica', 'versicolor', 'setosa', 'virginica', 'setosa',
Out[ ]:
                'virginica', 'setosa', 'versicolor', 'versicolor', 'versicolor',
               'virginica', 'versicolor', 'versicolor', 'versicolor',
               'versicolor', 'setosa', 'versicolor', 'versicolor', 'setosa',
               'setosa', 'virginica', 'versicolor', 'setosa', 'setosa',
               'virginica', 'setosa', 'setosa', 'versicolor', 'versicolor',
               'setosa', 'virginica', 'versicolor', 'setosa', 'virginica',
               'virginica', 'versicolor', 'setosa', 'versicolor', 'versicolor',
               'versicolor', 'virginica', 'setosa', 'virginica', 'setosa',
               'setosa'], dtype='<U10')
In [ ]:
         from sklearn import metrics
         score = metrics.accuracy_score(y_test,y_prediction)
         print("Gaussian Naive Bayes model accuracy (in %):", metrics.accuracy_score(y_test,y_pr
        Gaussian Naive Bayes model accuracy (in %): 100.0 %
In [ ]:
         from sklearn import metrics
         cm= metrics.confusion matrix(y test, y prediction)
        array([[16, 0, 0],
Out[ ]:
               [ 0, 18, 0],
               [ 0, 0, 11]], dtype=int64)
In [ ]:
         import seaborn as sns
         plt.figure(figsize= (12,12))
         sns.heatmap(cm, annot = True, fmt= ".3f", linewidths=.5, square = True, cmap = 'Spectra'
         plt.ylabel("Actual label")
         plt.xlabel("predicted label")
         sample title= ("Gaussian Naive Bayes model accuracy in (%): {0}" .format(score*100))
         plt.title(sample title, size= 15)
Out[]: Text(0.5, 1.0, 'Gaussian Naive Bayes model accuracy in (%): 100.0')
```

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