

```
In [1]: # Data:
        # artificially created dataset with two feature columns and a label column
        # with 0 and 1
        # data could be interpreted as a group of people who exhibit or not certain
        # medical symptoms based on Age and Weight
        # Classification:
        # apply KNN and Naive Bayes and compare results
        # emphasis is on visual representation of the models goodness of fit
```

```
In [1]: # import libraries

import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
sns.set(style = "whitegrid", font_scale = 1.5)
```

```
In [2]: # artificial data generation using normal distributions

np.random.seed(0)
```

```
In [3]: # generating dataset with 600 points separated into two classes

# labels
class_1 = np.transpose([np.zeros(100)]) # 0 - do not exhibit symptoms
class_2 = np.transpose([np.ones(100)]) # 1 - exhibit symptoms

# features
A1 = np.transpose([np.random.normal(30, 5, 100)])
A2 = np.transpose([np.random.normal(200, 10, 100)])
A = np.concatenate((A1, A2, class_1), axis = 1)

B1 = np.transpose([np.random.normal(45, 5, 100)])
B2 = np.transpose([np.random.normal(210, 10, 100)])
B = np.concatenate((B1, B2, class_1), axis = 1)

C1 = np.transpose([np.random.normal(40, 5, 100)])
C2 = np.transpose([np.random.normal(175, 10, 100)])
C = np.concatenate((C1, C2, class_1), axis = 1)

D1 = np.transpose([np.random.normal(40, 8, 100)])
D2 = np.transpose([np.random.normal(245, 10, 100)])
D = np.concatenate((D1, D2, class_2), axis = 1)

E1 = np.transpose([np.random.normal(57, 4, 100)])
E2 = np.transpose([np.random.normal(230, 15, 100)])
E = np.concatenate((E1, E2, class_2), axis = 1)

F1 = np.transpose([np.random.normal(60, 3, 100)])
F2 = np.transpose([np.random.normal(190, 15, 100)])
F = np.concatenate((F1, F2, class_2), axis = 1)

data = np.concatenate((A, B, C, D, E, F), axis = 0)
```

```
In [4]: # plot data

# set plot x and y limits
x_min = 10
x_max = 80

d_x = 10

y_min = 120
y_max = 300

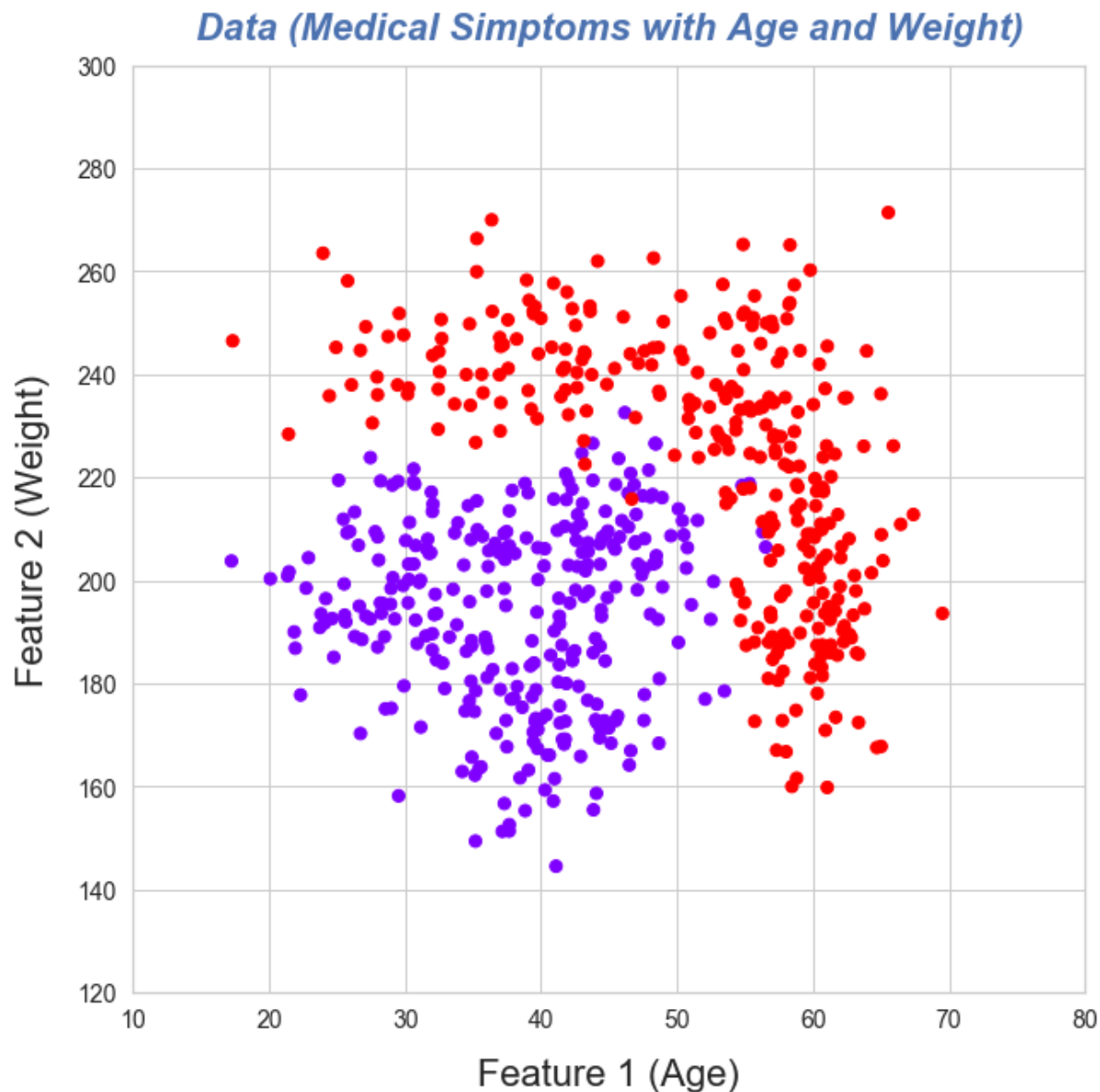
d_y = 20

plt.figure(figsize = (10, 10))

ax = plt.axes()
ax.set_xlim(x_min, x_max)
ax.set_xticks(np.arange(x_min, x_max + d_x, d_x))
ax.set_ylim(y_min, y_max)
ax.set_yticks(np.arange(y_min, y_max + d_y, d_y))

plt.scatter(data[:, 0], data[:, 1], c = data[:, 2], cmap = 'rainbow', s = 50)
plt.xlabel('Feature 1 (Age)', fontsize = 22, labelpad = 15)
plt.ylabel('Feature 2 (Weight)', fontsize = 22, labelpad = 15)
plt.title('Data (Medical Symptoms with Age and Weight)', fontsize = 22, fontwe
ight = 'bold',
         fontstyle = 'italic', pad = 15, c = 'b')
plt.tick_params(labelsize = 14)

plt.show()
```



```
In [6]: # as shown above data has been artificially generated, yet it is quite realistic
# the problem setup: use ML model with this data to predict whether somebody is likely to exhibit the symptoms
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In [5]: # separate data into features, X, and target, y
```

```
X = data[:, :-1]
y = data[:, -1]
```

```
In [6]: # since KNN uses distance as a measure of separation, we need to scale the features
```

```
from sklearn.preprocessing import StandardScaler

scaler = StandardScaler()

X = scaler.fit_transform(X)
```

```
In [9]: # build KNN model and apply it to the data set
```

```
In [7]: # first: find the optimal n_neighbors using the elbow method

# split data in train/test sets
from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, random_state=0)

# import class KNeighborsClassifier
from sklearn.neighbors import KNeighborsClassifier
```

```
In [8]: # use elbow method to find optimal n_neighbors

error_rate = [] # initialize array error_rate

# create loop to run through different n_neighbors
for i in range(1,30):
    knc = KNeighborsClassifier(n_neighbors = i)
    knc.fit(X_train, y_train)
    pred_i = knc.predict(X_test)
    error_rate.append(np.mean(pred_i != y_test)) # != is non-equal - this determines the error rate
```

In [9]: *# plot error rate*

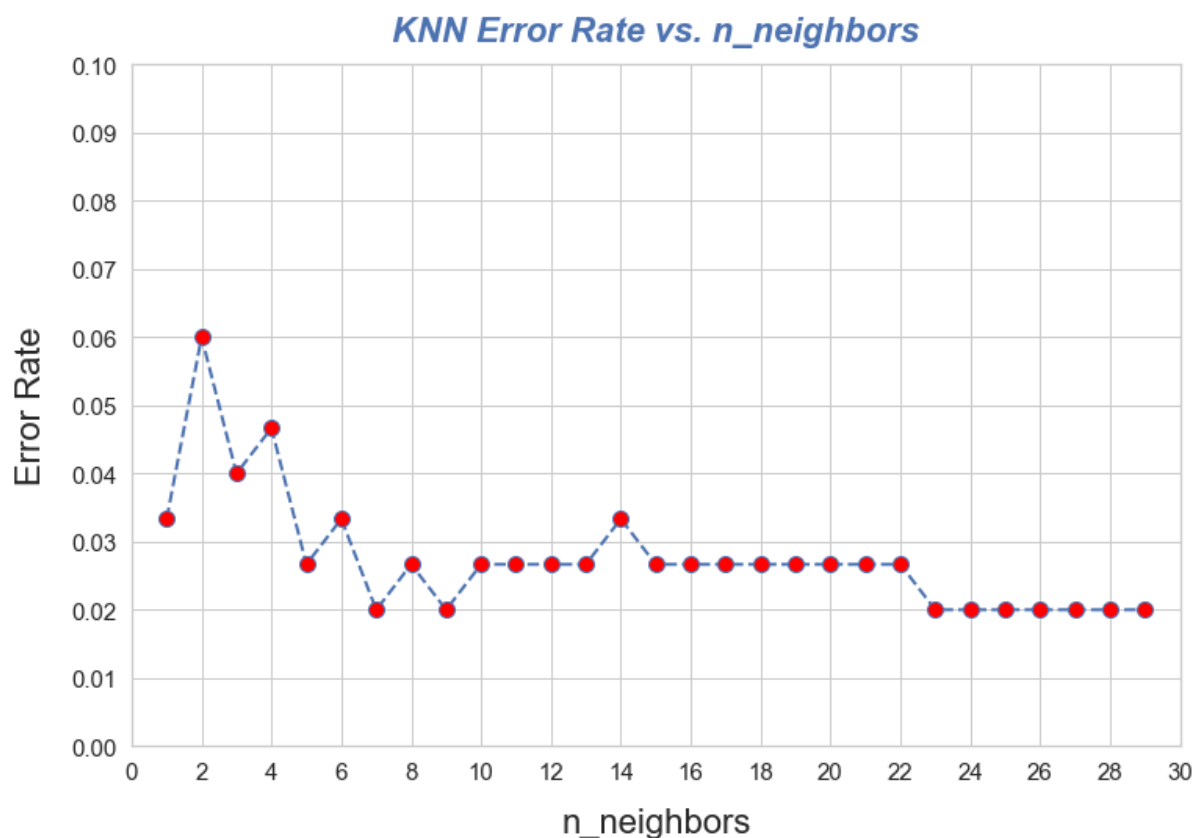
```
plt.figure(figsize=(12,8))

axes = plt.axes()
axes.set_xlim(0, 30)
axes.set_xticks(np.arange(0, 31, 2))
axes.set_ylim(0.0, 0.1)
axes.set_yticks(np.arange(0.0, 0.11, 0.01))

plt.plot(range(1,30), error_rate, 'b--', lw=2, marker='o', ms = 10, markerface
color='red')

# Add labels to your graph
plt.xlabel('n_neighbors', fontsize = 22, labelpad = 15)
plt.ylabel('Error Rate', fontsize = 22, labelpad = 15)
plt.title('KNN Error Rate vs. n_neighbors', fontsize = 22, fontweight = 'bold'
, fontstyle = 'italic', pad = 15, c = 'b')
plt.tick_params(labelsize = 15)

plt.show()
```



In [13]: *# results show that after n\_neighbors = 7 error rate flattens out --> use n\_neighbors = 7 in model*

In [10]: *# create KNN model with n\_neighbors = 7*

```
knc = KNeighborsClassifier(n_neighbors = 7)

# fit and predict
knc.fit(X_train, y_train)
y_pred_knc = knc.predict(X_test)
```

In [11]: *# compare predictions with true test data, y\_test*

```
from sklearn.metrics import confusion_matrix, classification_report

print('Confusion Matrix - KNeighborsClassifier with n_neighbors = 7:')
print(confusion_matrix(y_test, y_pred_knc))
print('\n')
print('Classification Report - - KNeighborsClassifier with n_neighbors = 7:')
print(classification_report(y_test, y_pred_knc))
```

Confusion Matrix - KNeighborsClassifier with n\_neighbors = 7:

```
[[67  3]
 [ 0 80]]
```

Classification Report - - KNeighborsClassifier with n\_neighbors = 7:

	precision	recall	f1-score	support
0.0	1.00	0.96	0.98	70
1.0	0.96	1.00	0.98	80
accuracy			0.98	150
macro avg	0.98	0.98	0.98	150
weighted avg	0.98	0.98	0.98	150

In [16]: *# excellent prediction accuracy*

In [17]: *# visualize how well the model captured both the training and test data*

```
In [12]: # define mapping function

def mapData(clf):
    # Create a dense grid of points to sample
    xx, yy = np.meshgrid(np.arange(-ax_min, ax_max, .005),
                          np.arange(-ax_min, ax_min, .005))

    # Convert to Numpy arrays
    npx = xx.ravel()
    npy = yy.ravel()

    # Convert to a List of 2D points
    samplePoints = np.c_[npx, npy]

    # Generate predicted labels (cluster numbers) for each point
    Z = clf.predict(samplePoints)

    plt.figure(figsize=(10, 10))

    Z = Z.reshape(xx.shape) # Reshape results to match xx dimension
    plt.contourf(xx, yy, Z, cmap=plt.cm.Paired, alpha=0.6) # Draw the contour
    plt.scatter(X_p[:,0], X_p[:,1], s = 50, c=y_p, cmap = 'rainbow') # data points
    plt.xlabel('Feature 1 (scaled)', fontsize = 20, labelpad = 15)
    plt.ylabel('Feature 2 (scaled)', fontsize = 20, labelpad = 15)
    plt.title(title_str, fontsize = 22, c = 'blue', pad = 20)
    plt.tick_params(labelsize= 18)
    plt.show()
```

```
In [13]: # mapping of training data

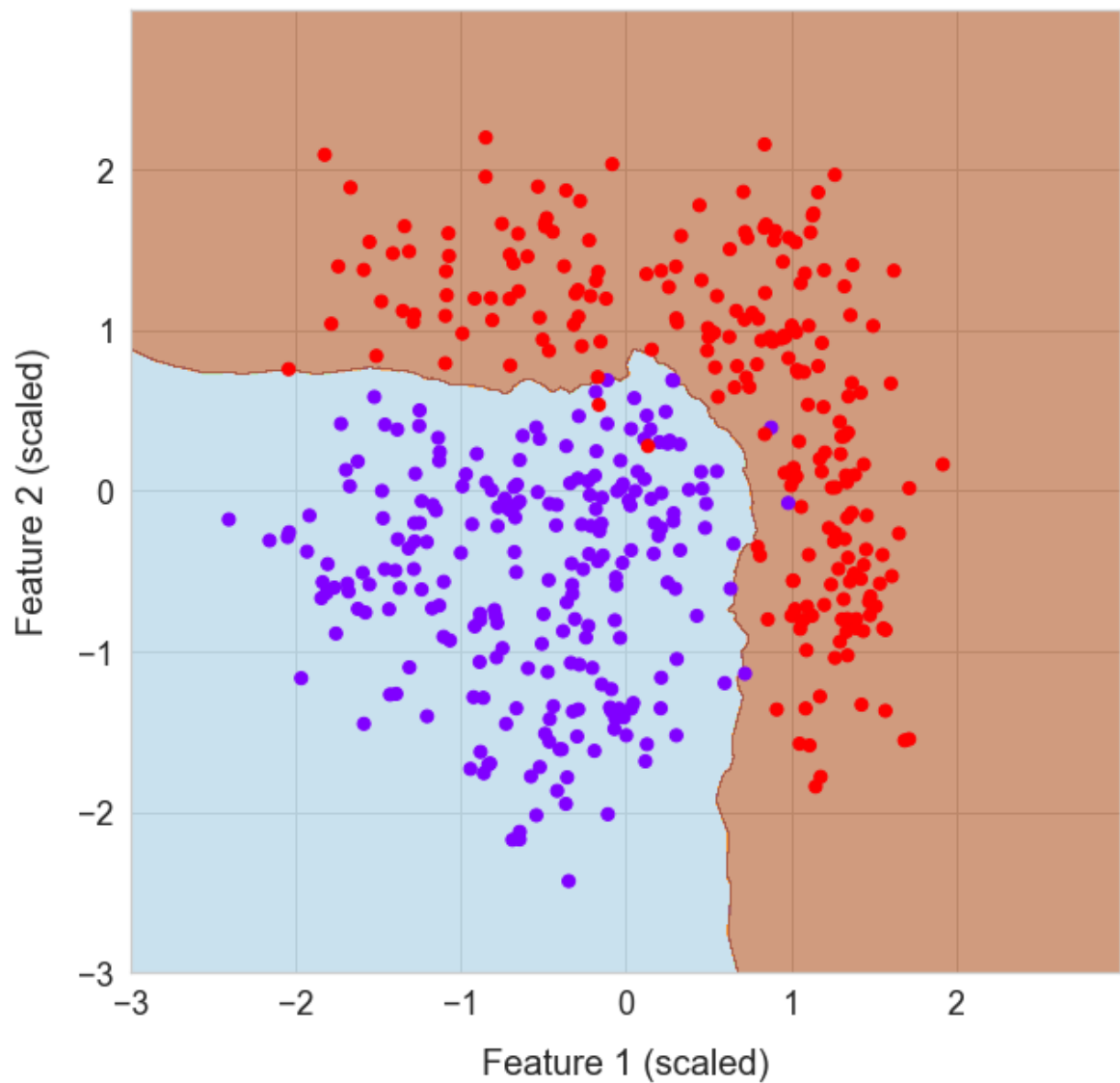
ax_min = 3.0
ax_max = 3.0

X_p = X_train
y_p = y_train

title_str = 'Mapping of Training Data: KNClassifier with n_neighbors = 7'

mapData(knc)
```

Mapping of Training Data: KNClassifier with n\_neighbors = 7



```
In [20]: # excellent classes separation!
```



```
In [14]: # mapping of test data

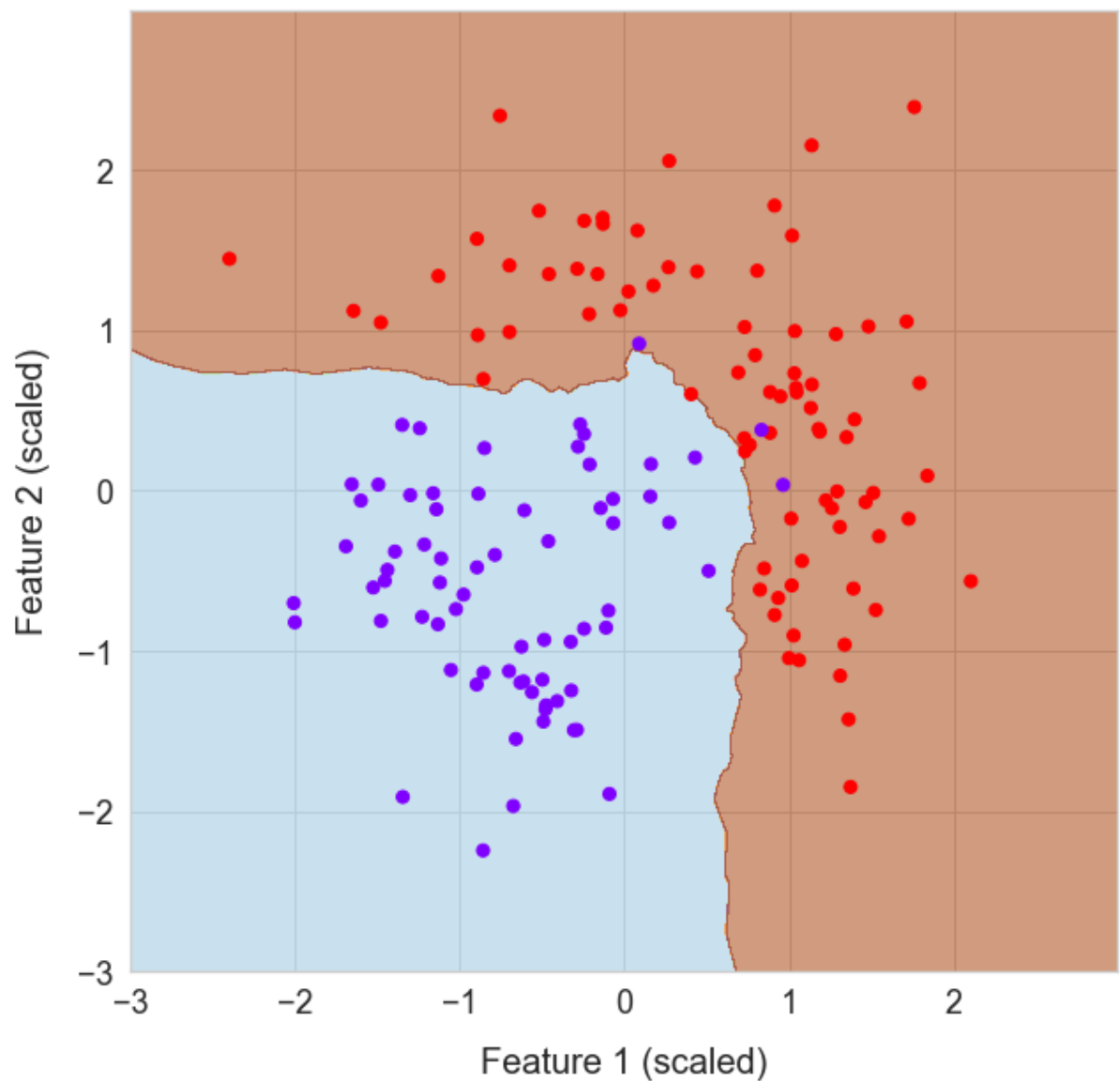
ax_min = 3.0
ax_max = 3.0

X_p = X_test
y_p = y_test

title_str = 'Mapping of Test Data: KNClassifier with n_neighbors = 7'

mapData(knc)
```

Mapping of Test Data: KNClassifier with n\_neighbors = 7



```
In [22]: # excellent classes separation here, too
```

```
In [23]: # use NaiveBayes model with same data and compare results to KNClassifier results
```

In [15]: *# create Naive Bayes model, fit, predict*

```
from sklearn.naive_bayes import GaussianNB
nbc = GaussianNB()

nbc.fit(X_train, y_train)
y_pred_nbc = nbc.predict(X_test)
```

In [16]: *# compare predictions with true test data, y\_test*

```
print('Confusion Matrix - Naive Bayes:')
print(confusion_matrix(y_test, y_pred_nbc))
print('\n')
print('Classification Report - - Naive Bayes:')
print(classification_report(y_test, y_pred_nbc))
```

Confusion Matrix - Naive Bayes:

```
[[65  5]
 [ 1 79]]
```

Classification Report - - Naive Bayes:

	precision	recall	f1-score	support
0.0	0.98	0.93	0.96	70
1.0	0.94	0.99	0.96	80
accuracy			0.96	150
macro avg	0.96	0.96	0.96	150
weighted avg	0.96	0.96	0.96	150

In [17]: *# for comparison print again knn results*

```
print('Confusion Matrix - KNeighborsClassifier with n_neighbors = 7:')
print(confusion_matrix(y_test, y_pred_knn))
print('\n')
print('Classification Report - - KNeighborsClassifier with n_neighbors = 7:')
print(classification_report(y_test, y_pred_knn))
```

Confusion Matrix - KNeighborsClassifier with n\_neighbors = 7:

```
[[67  3]
 [ 0 80]]
```

Classification Report - - KNeighborsClassifier with n\_neighbors = 7:

	precision	recall	f1-score	support
0.0	1.00	0.96	0.98	70
1.0	0.96	1.00	0.98	80
accuracy			0.98	150
macro avg	0.98	0.98	0.98	150
weighted avg	0.98	0.98	0.98	150

```
In [27]: # KNNClassifier does a little bit better, but both models have excellent prediction scores
```

```
In [28]: # visualize data separation with Naive Bayes
```

```
In [18]: # mapping of training data
```

```
ax_min = 3.0
```

```
ax_max = 3.0
```

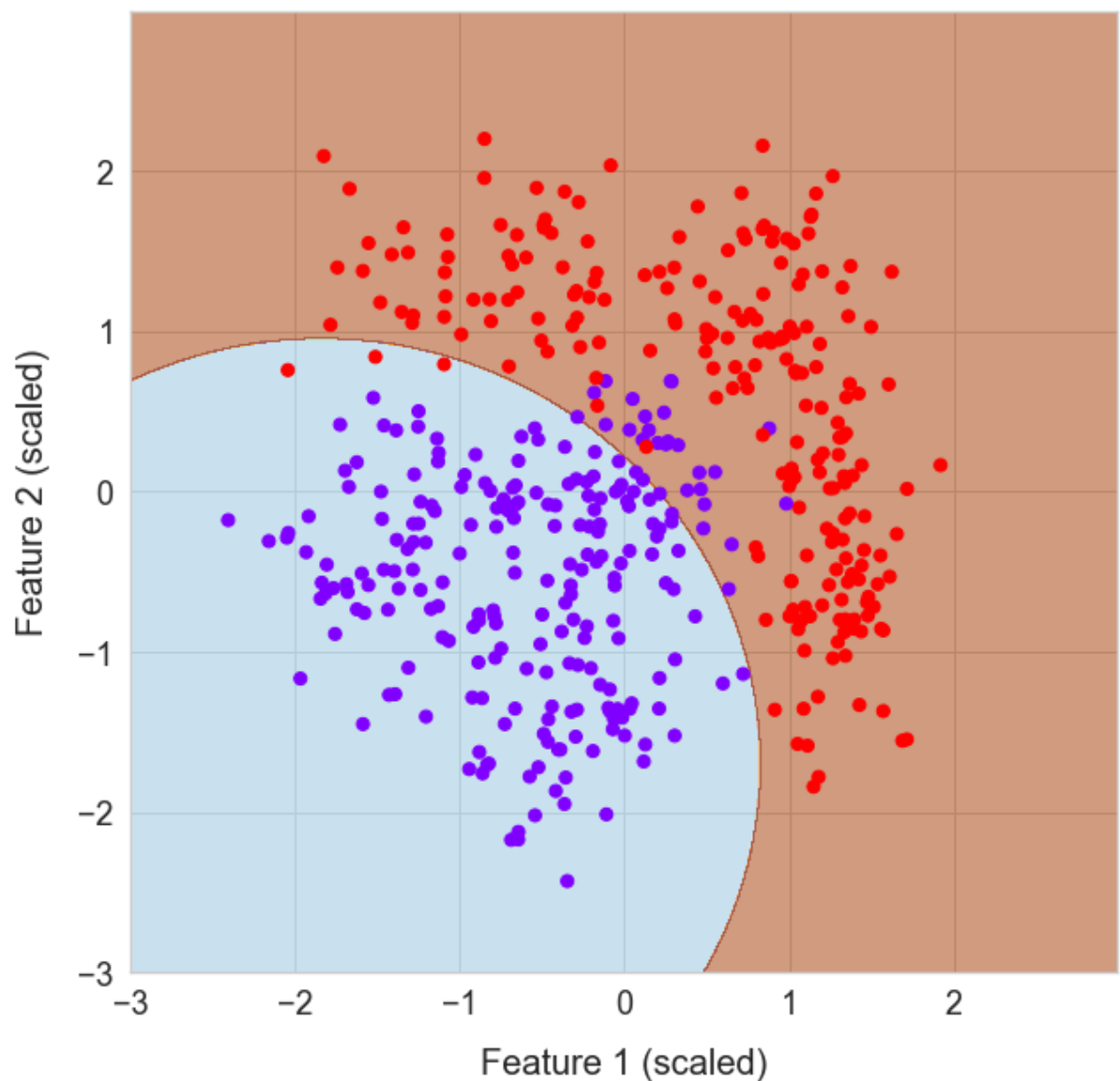
```
X_p = X_train
```

```
y_p = y_train
```

```
title_str = 'Mapping of Training Data: Naive Bayes'
```

```
mapData(nbc)
```

### Mapping of Training Data: Naive Bayes



```
In [30]: # now we can see the difference between the two models more clearly  
# Naive Bayes experiences problems with classifying the 'blue' data points which propagate deeper into the 'red' region  
# we would not have seen these issues if we were gauging only by the confusion matrix and the classification report  
# thus, it is important (if possible) to have a visual or some other representation of the goodness of fit
```

```
In [19]: # mapping of test data

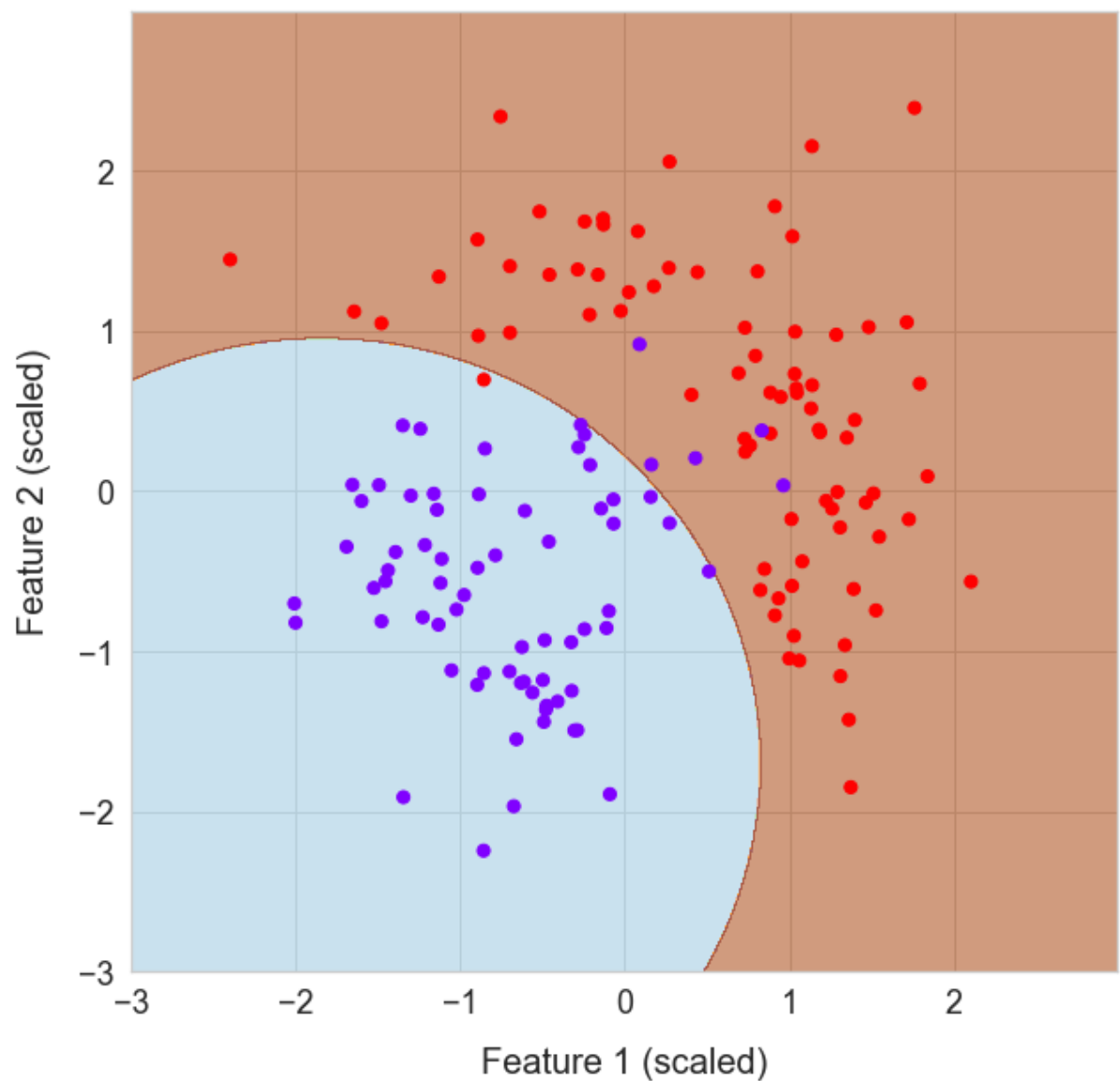
ax_min = 3.0
ax_max = 3.0

X_p = X_test
y_p = y_test

title_str = 'Mapping of Test Data: Naive Bayes'

mapData(nbc)
```

Mapping of Test Data: Naive Bayes



```
In [32]: # same issue is illustrated here, too
```

```
In [33]: # Summary:
          # KNN and NaiveBayes were applied for classification of hypothetical medical dataset
          # Confusion matrix and Classification Report show high accuracy scores for both models
          # Visual representation of the fit, however, reveals that KNN performs better than NaiveBayes with this dataset
```