November 8, 2020

This problem was adapted from Professor Farimani's paper. If you are interested in learning more, you can read it here.

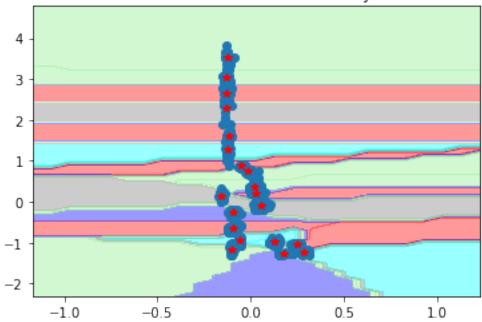
```
[15]: import numpy as np
  import pandas as pd
  import matplotlib.pyplot as plt
  from sklearn.model_selection import train_test_split
  from sklearn.cluster import KMeans
  from sklearn.ensemble import RandomForestClassifier
  from matplotlib.colors import ListedColormap
```

```
[16]: # (a)
      # data preprocessing
      #import data
      dataset=pd.read_csv('data.csv')
      #get string labels
      hd = list(dataset.columns)
      check = ~dataset.columns.str.contains('~Unnamed') #get boolean of named and_
      \rightarrowunnamed string label
      header = []
      for i in range(len(hd)):
          if check[i]:
              header.append(hd[i])
      #map string label to numeric values
      header = pd.factorize(header)[0]
      #verify range of numeric label
      assert(len(header) == 20)
      #import raw feature values
      x = dataset.iloc[:,:].values
      r = np.shape(x)[0] #number of data per label
      #initialize rearranged data and label
      X = x[:,[0,1]]
      y = np.ones(r)*header[0]
      #iterate to rearrange data and label
      for i in range(2,len(header)*2, 2):
```

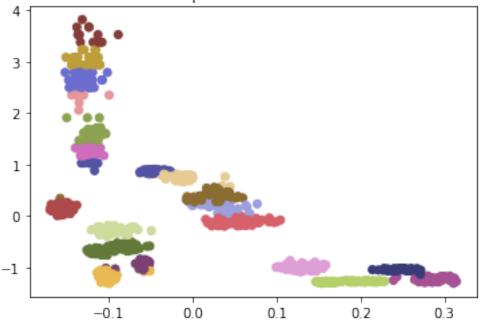
```
arr = x[:, [i, i+1]]
          X = np.vstack((X, arr))
          y = np.hstack((y, np.ones(r)*header[int(i/2)]))
      #verify shape of rearranged data and label
      assert(X.shape[0] == 2000)
      assert(y.shape[0] == 2000)
      assert(X.shape[1] == 2)
      # Save the preprocessed data as a new csv file as you may need that for problem,
      ⊶3
      data = np.hstack((X, y.reshape(-1,1)))
      dict = {'x1': data[:,0], 'x2': data[:,1], 'label': data[:,2]}
      df = pd.DataFrame(dict)
      df.to_csv('output_data.csv')
[21]: # (b)
      # k-means
      #The train-test split to be used for the dataset is 80%-20%
      X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.30)
      #Mean centering and normalization
      mean = np.mean(X_train, axis=0)
      std = np.std(X_train)
      X_train = (X_train - mean)/std
      X_{test} = (X_{test} - mean)/std
      #K-Means -> pretending it is unsupervised
      kmeans = KMeans(n clusters=20, max iter=1000, random state=200)
      kmeans.fit(X_train)
      center = kmeans.cluster centers
      y_pred_tr = kmeans.predict(X_train)
      y_pred_ts = kmeans.predict(X_test)
      # print accuracy on training and testing datasets
      count_pred = np.sum(y_train == y_pred_tr, axis=0)
      accuracy_train = (count_pred/y_train.shape[0])*100
      print("Accuracy of training dataset is",accuracy_train, '%')
      count_pred = np.sum(y_test == y_pred_ts, axis=0)
      accuracy_test = (count_pred/y_test.shape[0])*100
      print("Accuracy of test dataset is",accuracy_test, '%')
      #adapted from sklearn examples#
      colors = ('red', 'blue', 'lightgreen', 'gray', 'cyan')
      cmap = ListedColormap(colors[:len(np.unique(y_pred_tr))])
      x1_low, x1_high = X_train[:, 0].min() - 1, X_train[:, 0].max() + 1
      x2_{low}, x2_{high} = X_{train}[:, 1].min() - 1, <math>X_{train}[:, 1].max() + 1
```

Accuracy of training dataset is 10.285714285714285% Accuracy of test dataset is 9.166666666666666%

Data with decision boundary

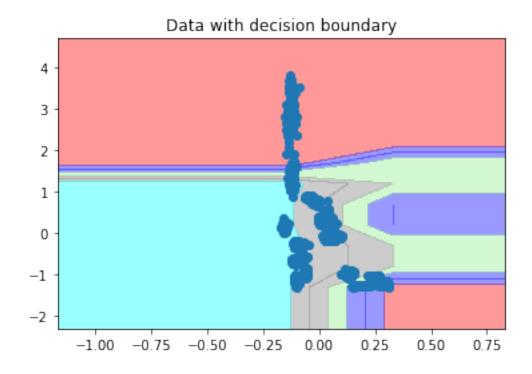


Scatter plot with centroid stars

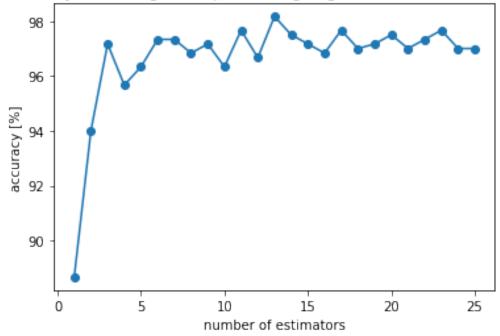


```
[24]: # (c)
      # random forest
      rfc = RandomForestClassifier()
      rfc.fit(X_train, y_train)
      y_pred_tr = rfc.predict(X_train)
      y_pred_ts = rfc.predict(X_test)
      count_pred = np.sum(y_train == y_pred_tr, axis=0)
      accuracy_train = (count_pred/y_train.shape[0])*100
      print("Accuracy of training dataset is",accuracy_train, '%')
      count_pred = np.sum(y_test == y_pred_ts, axis=0)
      accuracy_test = (count_pred/y_test.shape[0])*100
      print("Accuracy of test dataset is",accuracy_test, '%')
      #adapted from sklearn examples#
      colors = ('red', 'blue', 'lightgreen', 'gray', 'cyan')
      cmap = ListedColormap(colors[:len(np.unique(y_pred_tr))])
      x1_low, x1_high = X_train[:, 0].min() - 1, X_train[:, 0].max() + 1
      x2_low, x2_high = X_train[:, 1].min() - 1, X_train[:, 1].max() + 1
      xx1, xx2 = np.meshgrid(np.arange(x1_low, x1_high, 0.5), np.arange(x2_low,__
       \rightarrowx2_high, 0.5))
```

```
z = rfc.predict(np.array([xx1.ravel(), xx2.ravel()]).T).reshape(xx1.shape)
plt.figure()
plt.contourf(xx1, xx2, z, alpha=0.4, cmap=cmap)
plt.xlim(xx1.min(), xx1.max())
plt.ylim(xx2.min(), xx2.max())
plt.plot(X_train[:,0], X_train[:,1], 'o')
plt.title("Data with decision boundary")
num = np.arange(1, 26)
accuracy_num = []
for k in range(1, len(num)+1):
    rfc = RandomForestClassifier(n_estimators=k)
    rfc.fit(X_train, y_train)
    y_pred_nm = rfc.predict(X_test)
    count_pred = np.sum(y_test == y_pred_nm, axis=0)
    accuracy_num.append((count_pred/y_test.shape[0])*100)
plt.figure()
plt.plot(num, accuracy_num,'-o')
plt.xlabel('number of estimators')
plt.ylabel('accuracy [%]')
plt.title('Accurcay on testing set in percentage against the no. of \Box
⇔estimators')
plt.show()
```







1 (d)

2 Analysis

When comparing the plots of kmeans versus random forest (RF), we can see that the decision boundary is significantly different. It's difficult to understand the decision boundary of the random forest despite it's high accuracy. Although the kmeans has a much lower accuracy because it performs an unsupervised learning algorithm where the labels are unknown, the decision boundary is much clearer. The decision boundary is much different for the random forest because is performs a supervised learning algorithm where the labels are given, thus increasing it's accuracy. If possible, it is clear that one should train with labels to increase accuracy in predicting labels.

[]: