## **A4Q4**

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### 1 CMPT 423/820

### 1.1 Assignment 4 Question 4

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The a4q4.csv file contains a 6D dataset (each sample has 6 numeric features, one class label, and an index column). It's too many dimensions to visualize directly.

We'll try to visualize the data, and build a classifier from it. Then we'll apply PCA, use the first two principle components for visualization and to train a classifier.

To complete this question: 1. Produce a scatter plot of any 2 dimensions. Do you see any natural clusters in 2D? 1. Build a classifier (your choice, e.g., Naive Bayes, KNN, etc) using the 6 features and the label. Get an estimate of the classifier performance using cross validation. 1. Use Principle Component Analysis to reduce the data to 2 principle components. (look at the docs: sklearn.decomposition.PCA) 1. Produce a scatter plot of the 2 principle components. Do you see any natural clusters? 1. Build a classifier (the same type you used above) using the 2 principle components and the label. Get an estimate of the classifier performance using cross validation. 1. Report on the difference in classifier accuracy.

First some libraries...

```
[0]: import matplotlib as mpl
  import matplotlib.pyplot as plt
  import pandas as pd
  import numpy as np

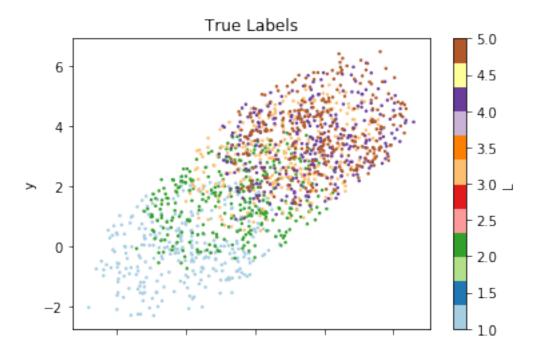
from sklearn import decomposition

[0]: df = pd.read_csv('data/a4q4.csv', index_col=0)

#Here we can choose a global colormap, to help visualization
# see https://matplotlib.org/tutorials/colors/colormaps.html
  cmapstr = 'Paired'

# here we choose how big each point in the plot will be
  ptsize = 2

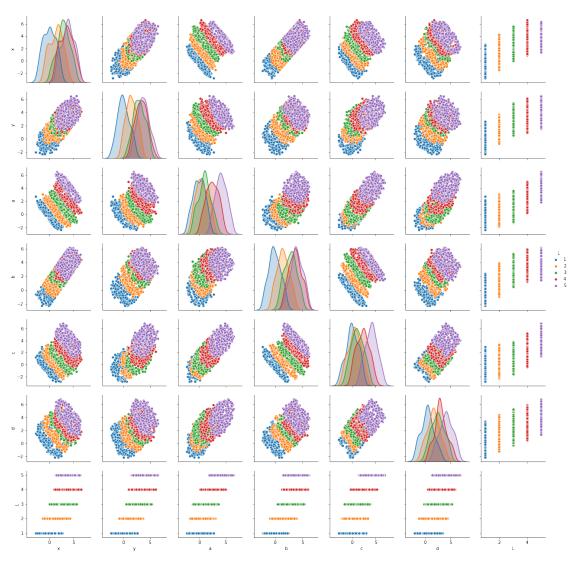
# plot any 2 of the 6 features, using the given label
```



### 1.1.1 1. Produce a scatter plot of any 2 dimensions.

[0]: <seaborn.axisgrid.PairGrid at 0x1a2dcb5310>





Do you see any natural clusters in 2D?

Yes, some features are separable in the plots. Here is the list: \* x and a \* b and c

# 1.1.2 2. Build a classifier (your choice, e.g., Naive Bayes, KNN, etc) using the 6 features and the label. Get an estimate of the classifier performance using cross validation.

I choose *Naive Bayes* classifier for this purpose because we have thoroughly investigated the application of NB on the Iris dataset. The following function implements a *Gaussian Naive Bayes* classifier and calculated 10-fold cross-validation on the given dataset.

[0]: from sklearn.naive\_bayes import GaussianNB from sklearn.model\_selection import cross\_val\_score

```
KFOLDS = 10
   def nb_classifier(data_in, label_in):
        :purpose: This function creates an NB classifier and calculates
        its cross validated accuracy
        :param data_in: dataset
        :param label_in: true answers for the dataset
        :return: the accuracy of the model with 5 point accuracy
       clf_nb = GaussianNB()
       # fitting the model to the data
       clf_nb.fit(data_in, label_in)
       result = cross_val_score(clf_nb, data_in,
                        label_in,
                        cv=KFOLDS,
                        scoring='accuracy')
       return round(result.mean(),5)
[0]: labels = df.values[:, 6]
   data = df.values[:, :6]
   result = nb_classifier(data, labels)
   print('Naive Bayes accuracy using all features:', result)
```

Naive Bayes accuracy using all features: 0.91533

## 1.1.3 3. Use Principle Component Analysis to reduce the data to 2 principle components. (look at the docs: sklearn.decomposition.PCA)

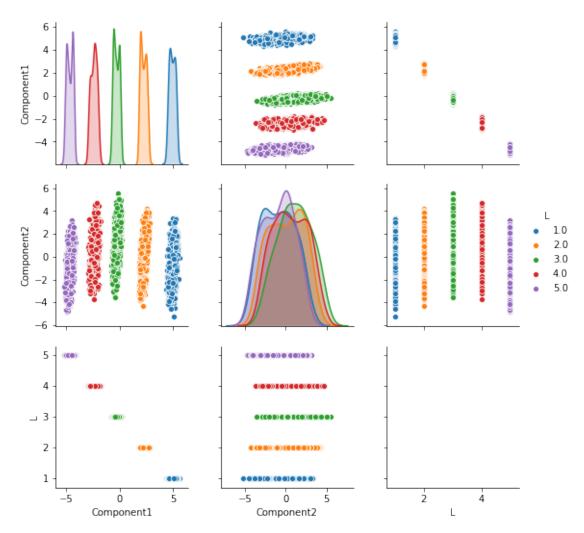
```
pca_data['L'] = df.values[:, 6]
   pca_data
[0]:
          Component1 Component2
                                    L
            4.620505
                      -1.270596
                                 1.0
    0
    1
            4.887661
                       0.586271
                                  1.0
    2
            4.570505
                      -1.464535 1.0
    3
            4.933026
                       -1.594514
                                 1.0
            4.563424
                      -3.746621
                                 1.0
                 . . .
                             . . .
                                 . . .
          -4.696963
    1495
                      -4.116303 5.0
    1496
          -4.950297
                      -2.309152 5.0
    1497
          -4.479358
                       -1.594562 5.0
    1498
          -4.828170
                       1.153164 5.0
    1499
           -4.429090
                       -0.327566 5.0
```

### 1.1.4 4. Produce a scatter plot of the 2 principle components. Do you see any natural clusters?

[0]: <seaborn.axisgrid.PairGrid at 0x1a28e74dd0>

[1500 rows x 3 columns]

### Scatter Plot of 2 principle components



# 1.1.5 5. Build a classifier (the same type you used above) using the 2 principle components and the label. Get an estimate of the classifier performance using cross validation.

```
[0]: labels = pca_data.values[:, 2]
data = pca_data.values[:, :2]

result = nb_classifier(data, labels)

print('Naive Bayes accuracy using principle components:', result)
```

Naive Bayes accuracy using principle components: 1.0

### 1.1.6 6. Report on the difference in classifier accuracy.

My classifier obtains an accuracy of 1.0 using principle components and an accuracy of 0.91 using the original dataset.

The scatter plot of principle components shows distinctive clusters compared to the scatter plot of all features with overlapping clusters. That's why *Naive Bayes* obtain better classification accuracy using principle components compared to the original dataset.

### 1.2 Grading

- 1. 3 marks: You built a classifier using the whole dataset, and reported an accuracy value.
- 2. 3 marks: You applied PCA, and visualized the first two principle components.
- 3. 3 marks: You built a classifier using the first two principle components, and reported an accuracy value.
- 4. 3 marks: You reported on the difference in accuracy between your two classifiers.