# Applied Data Mining: Homework #3

Due on Fill-in this please

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In this homework, you will work with Ionosphere Data Set to answer some questions regarding Principal Component Analysis (PCA), exploratory data analysis and k-means clustering. Here is the beginning of an R session that allows us to read this data from the web into our local R session:

# Problem 1

For the Ionosphere Data Set, answer the following questions:

### Discussion of Data

Briefly describe this data set—what is its purpose? How should it be used? What are the kinds of data it's using?

#### R Code

Using R, show code that answers the following questions:

1. How many entries are in the data set? Answer here ...

```
Listing 1: Sample R Script With Highlighting
```

```
%% You provide code here %%
```

2. How many unknown or missing data are in the data set? Answer here ...

Listing 2: Sample R Script With Highlighting

```
7% You provide code here 7%
```

3. Create a bar plot of 1st, 2nd, 35th variables. Label the plots properly. Discuss the distribution of values e.g., are uniform, skewed, normal. Place images of these bar plots into the document. Show the R code that you used below and discussion below that.

Listing 3: Sample R Script With Highlighting

```
%% You provide code here %%
```

#### Discussion of Bar Plots

Answer here...

### **Bar Plots**

Place images here with suitable captions.

4. Make a scatter plots of [V22, V20] and [V1, V2] variables and color the data points with the class variable [V35]. Discuss the plots, i.e., do you observe any relationships between variables?

Listing 4: Sample R Script With Highlighting

%% You **provide** code here %%

#### Discussion of Scatter Plots

Answer here...

# **Scatter Plots**

Place images here with suitable captions.

## Problem 2

In this question, you will run k-means clustering algorithm against Ionosphere data set. The input data for k-means is mydata[,-35] – removing the class variable since this is a clustering task.

## R Code

Using R, show code that answers the following questions:

1. Run "Lloyd, Forgy and Hartigan-Wong's" heuristic algorithms for k-means and report total within sum of squared error (SSE) for k=2 and nstart=50. Compare the results?i.e., which/why is better? Discuss nstart parameter. Show the R code that you used below and discussion and results below that.

Listing 5: Sample R Script With Highlighting

%% You provide code here %%

# Total SSE

Answer here...

### Discussion of nstart and Results

Answer here...

2. Elbow method is a technique used to decide optimal cluster number. The code below gives a plot of total SSE for k = 1, ..., 10. Discuss the elbow technique, i.e., what would be the optimal k based on the plot, can optimal k always be identified by elbow method?

#### Discussion of Results

Answer here...

## Problem 3

Use Principal Component Analysis (PCA) over Ionosphere Data Set to answer the below questions. You may want to use either "princomp()" or "prcomp()" functions in R. In this question, remove the 2nd (all 0s) and 35th variable (class variable) before using PCA.

```
> mydata <- mydata[,-35]
> mydata <- mydata[,-2]
> dim(mydata)
[1] 351 33
> mydata.pca <- prcomp(mydata, scale =TRUE)</pre>
```

#### R Code

Using R, show code that answers the following questions:

1. Make a scatter plot of PC1 and PC2 (the first and second principal components). Discuss principal components? What is PC1 and PC2? Show the R code that you used below and the scatter plot and discussion below that

Listing 6: Sample R Script With Highlighting

```
%% You provide code here %%
```

#### Scatter Plot

Place images here with suitable captions. Answer here...

# **Discussion of Principal Components**

Answer here...

2. You can observe the loadings as follows (using prcomp() function):

```
>mydata.pca$rotation
```

Discuss loadings in PCA?i.e., how are principal components and original variables of the data (mydata) related? (loadings(mydata.pca) if princomp() is used)

3. Scree plot is among the most popular methods to decide optimal dimension number.

```
> plot(mydata.pca, type = "1")
> screeplot(mydata.pca)
```

What is the optimal dimension number (d) for this data set? How much of the variation is kept with your optimal d? Discuss the results.