# Notebook 3 Clustering

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The dataset used for this project is an online retail set pulled from UCI. It contains tuples containing data on Customer ID, quantity purchased, price purchased at, description of the data, and country of origin of purchaser alongside some other columns.

Importing required packages into the project.

```
if(!require('tidyverse')){
  install.packages('tidyverse')
}
```

```
## Loading required package: tidyverse
```

```
## - Attaching packages -
                                                              - tidyverse 1.3.2 —
## √ ggplot2 3.4.1
                      ✓ purrr
                                1.0.1

√ dplyr

## √ tibble 3.1.8
                                1.1.0
## √ tidyr 1.3.0
                      ✓ stringr 1.5.0
           2.1.4

√ forcats 1.0.0

## √ readr
## -- Conflicts -
                                                        - tidyverse_conflicts() -
## X dplyr::filter() masks stats::filter()
## X dplyr::lag()
                    masks stats::lag()
```

```
if(!require('e1071')){
  install.packages('e1071')
}
```

```
## Loading required package: e1071
```

```
if(!require('caret')){
  install.packages('caret')
}
```

```
## Loading required package: caret
## Loading required package: lattice
##
## Attaching package: 'caret'
##
## The following object is masked from 'package:purrr':
##
## lift
```

```
if(!require('word2vec')){
  install.packages('word2vec')
}
## Loading required package: word2vec
## Warning: package 'word2vec' was built under R version 4.2.3
if(!require('cluster')){
  install.packages('cluster')
}
## Loading required package: cluster
if(!require('mclust')){
  install.packages('mclust')
}
## Loading required package: mclust
## Warning: package 'mclust' was built under R version 4.2.3
## Package 'mclust' version 6.0.0
## Type 'citation("mclust")' for citing this R package in publications.
##
## Attaching package: 'mclust'
##
## The following object is masked from 'package:purrr':
##
##
       map
library('word2vec')
library('caret')
library('tidyverse')
library('e1071')
library('cluster')
library('mclust')
data <- read.csv("data/online_retail.csv", header=TRUE)</pre>
set.seed(1234)
str(data)
```

```
## 'data.frame':
                   541909 obs. of 8 variables:
  $ InvoiceNo : chr "536365" "536365" "536365" ...
##
   $ StockCode : chr "85123A" "71053" "84406B" "84029G" ...
##
   $ Description: chr "WHITE HANGING HEART T-LIGHT HOLDER" "WHITE METAL LANTERN" "CREAM CUP
ID HEARTS COAT HANGER" "KNITTED UNION FLAG HOT WATER BOTTLE" ...
   $ Quantity : int 6 6 8 6 6 2 6 6 6 32 ...
##
##
   $ InvoiceDate: chr "12/1/2010 8:26" "12/1/2010 8:26" "12/1/2010 8:26" "12/1/2010 8:26"
   $ UnitPrice : num 2.55 3.39 2.75 3.39 3.39 7.65 4.25 1.85 1.85 1.69 ...
##
   $ CustomerID : int 17850 17850 17850 17850 17850 17850 17850 17850 17850 13047 ...
   $ Country
##
                : chr "United Kingdom" "United Kingdom" "United Kingdom" "United Kingdom"
```

```
normalize <- function(x) {
  return((x - min(x)) / (max(x) - min(x)))
}</pre>
```

I'm taking the Customer ID, Price, Quantity, and Country of Origin as the values to evaluate on. I'm omitting NA data instead of replacing it, as there are enough values already in the dataset to where removing some will not affect the data in a major way. Additionally, I get rid of some values that are not possible, i.e. removing negative bought items or items with a negative purchase price.

```
zero <- 0.0
data <- data[, c(4,6,7,8)]
data <- na.omit(data)
data <- data[data$Quantity > 0,]
data <- data[c(data$UnitPrice > as.numeric(zero)),]
```

I'm leaving this in, even though it failed. What I had attempted was to use the Word to Vec library to attempt to turn the character strings of Descriptions into a format that could be used in clustering. I wanted to create prediction matrices that could be used to help determine the likelyhood of a user ID purchasing something based on similar words. This was extremely time intensive and in the end, I couldn't get something to work in the time I had allotted.

```
##temp <- data$Description
#word_vector <- word2vec(temp, min_count = 1L, split=c(" \t", "\n."))
#emb <- as.matrix(word_vector)</pre>
```

Here is checking to make sure that the data cleaning steps worked.

```
i <- which.min(data$Quantity)
print(data[i,])</pre>
```

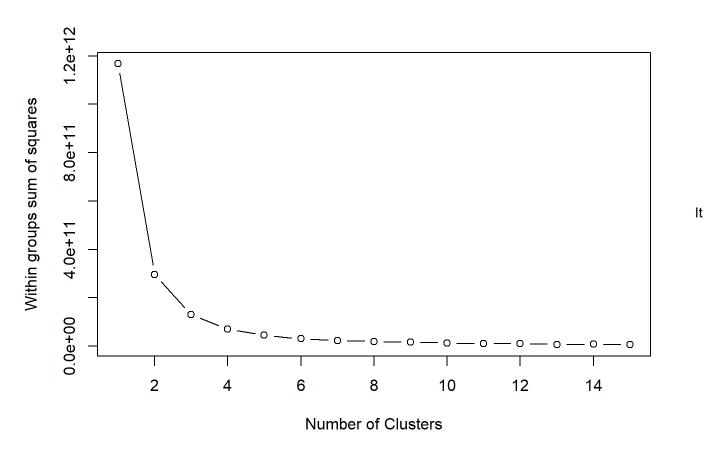
```
## Quantity UnitPrice CustomerID Country
## 114     1     1.25     15311 United Kingdom
```

```
i <- which.max(data$Quantity)
print(data[i,])</pre>
```

```
## Quantity UnitPrice CustomerID Country
## 540422 80995 2.08 16446 United Kingdom
```

```
data_norm <- as.data.frame(lapply(data[,1:2], normalize))</pre>
data norm$CustomerID <- data$CustomerID</pre>
#data_norm$Country <- factor(data$Country)
##Attempted to convert character vectors into a usuable format for clustering
##Unable to find a good solution
#for(x in data$Description){
# split <- unlist(lapply(sapply(x, strsplit, "\\s+|\\.+", USE.NAMES=FALSE), function(Z) {Z</pre>
[Z!=""]\}))
# vector <- rep(0, 50)
# for(y in split){
     predictions = tryCatch({
       vector <- vector + emb[y, ]</pre>
#
     }, error = function(e) "")
  }
#
  predictions <- rbind(predict(word_vector, vector, type='embedding'))</pre>
#}
```

With clustering, it is good to know how many clusters to push everything into. Plotting by the sum of squares will give a good approximation of the number of clusters to form. Of course, this isn't the absolute value to follow so I will try with some other values.



would seem that around 2 - 4 clusters is a good idea. The main change in the slope occurs between 2 and 3 clusters but doesn't really level off until around 4 to 5.

```
retail2KCluster <- kmeans(data_norm, 2, nstart=20)</pre>
retail3KCluster <- kmeans(data_norm, 3, nstart=20)</pre>
retail4KCluster <- kmeans(data_norm, 4, nstart=20)</pre>
aggregate(data norm,by=list(retail2KCluster$cluster),FUN=mean)
                              UnitPrice CustomerID
##
     Group.1
                 Quantity
           1 0.0001372414 0.0003591180
                                           16874.32
## 1
## 2
           2 0.0001574700 0.0004032289
                                           13907.60
aggregate(data_norm,by=list(retail3KCluster$cluster),FUN=mean)
##
     Group.1
                 Quantity
                              UnitPrice CustomerID
## 1
           1 0.0001376345 0.0003766579
                                           15148.45
## 2
           2 0.0001408467 0.0003495202
                                           17275.62
           3 0.0001676361 0.0004258166
## 3
                                           13274.43
aggregate(data_norm,by=list(retail4KCluster$cluster),FUN=mean)
```

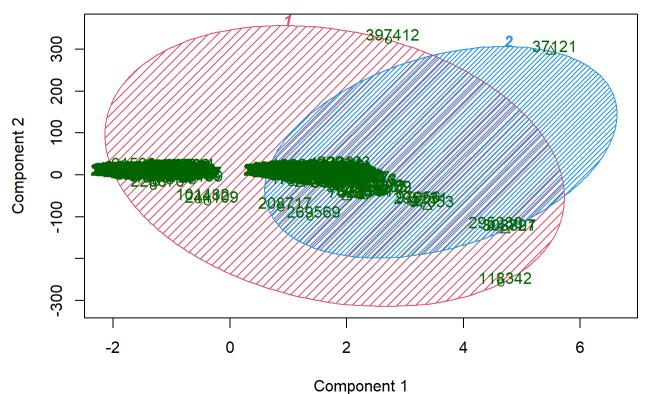
```
##
    Group.1
                 Quantity
                              UnitPrice CustomerID
## 1
           1 0.0001321754 0.0003526209
                                          17562.37
           2 0.0001416246 0.0003649616
## 2
                                          16091.85
## 3
           3 0.0001432305 0.0003924587
                                          14572.58
## 4
           4 0.0001771505 0.0004194005
                                          13017.31
```

```
data_norm2K <- data.frame(data_norm, retail2KCluster$cluster)
data_norm3K <- data.frame(data_norm, retail3KCluster$cluster)
data_norm4K <- data.frame(data_norm, retail4KCluster$cluster)</pre>
```

Now lets check the results of the clustering.

```
clusplot(data_norm2K, retail2KCluster$cluster, color=TRUE, shade=TRUE,
    labels=2, lines=0)
```

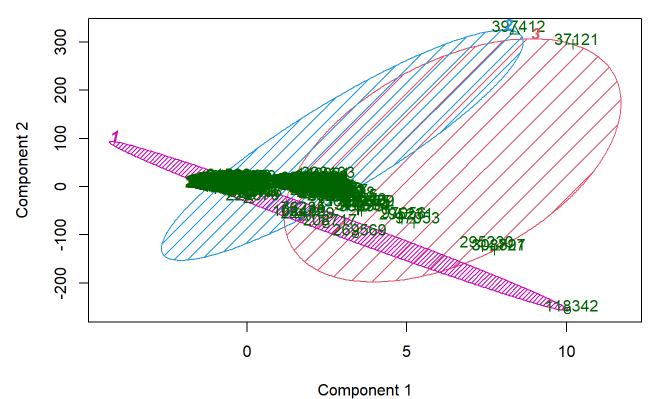
# CLUSPLOT( data\_norm2K )



These two components explain 71.72 % of the point variability.

```
clusplot(data_norm3K, retail3KCluster$cluster, color=TRUE, shade=TRUE,
    labels=2, lines=0)
```

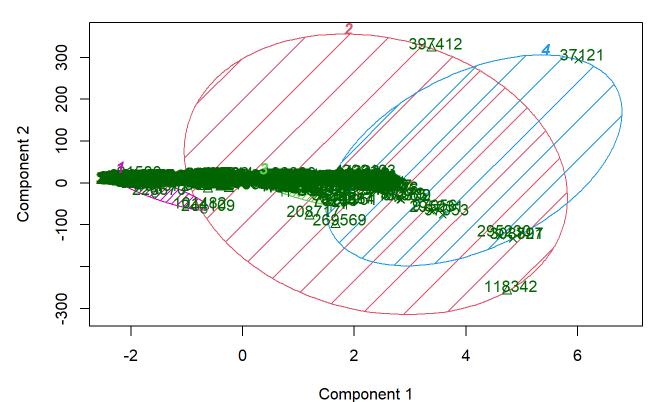
# CLUSPLOT( data\_norm3K )



These two components explain 60.46 % of the point variability.

clusplot(data\_norm4K, retail4KCluster\$cluster, color=TRUE, shade=TRUE,
 labels=2, lines=0)

## CLUSPLOT( data\_norm4K )



These two components explain 74.36 % of the point variability.

```
summary(retail2KCluster)
```

```
Length Class Mode
                397884 -none- numeric
## cluster
## centers
                     6 -none- numeric
## totss
                     1 -none- numeric
## withinss
                     2 -none- numeric
## tot.withinss
                     1 -none- numeric
## betweenss
                     1 -none- numeric
## size
                     2 -none- numeric
## iter
                     1 -none- numeric
## ifault
                     1 -none- numeric
```

summary(retail3KCluster)

```
##
                Length Class Mode
                397884 -none- numeric
## cluster
## centers
                     9 -none- numeric
## totss
                    1 -none- numeric
## withinss
                    3 -none- numeric
## tot.withinss 1 -none- numeric
## betweenss 1 -none- numeric
## size
                    3 -none- numeric
## iter
                    1 -none- numeric
## ifault
                     1 -none- numeric
```

### summary(retail4KCluster)

```
Length Class Mode
##
                397884 -none- numeric
## cluster
## centers
                   12 -none- numeric
## totss
                    1 -none- numeric
## withinss
                    4 -none- numeric
## tot.withinss 1 -none- numeric
## betweenss 1 -none- numeric
## size
                    4 -none- numeric
## iter
                    1 -none- numeric
## ifault
                      1 -none- numeric
```

Heirachical Agglomerative The data's far too big for this clustering method, so I'm cutting it down drastically.

```
temp <- sample(1:nrow(data_norm), 0.1*nrow(data_norm), replace=FALSE)
reduced_normalized <- data[temp,]

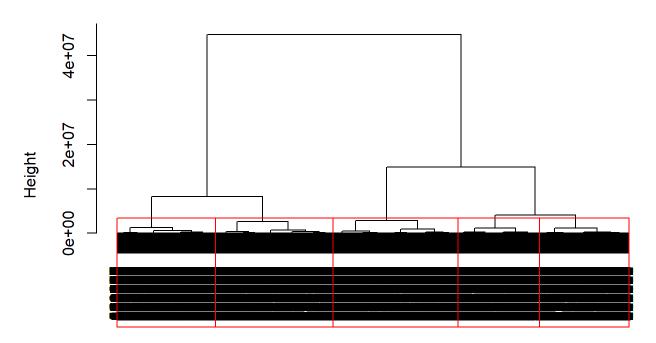
# Ward Hierarchical Clustering
euclidean_distance <- dist(reduced_normalized, method = "euclidean") # distance matrix

## Warning in dist(reduced_normalized, method = "euclidean"): NAs introduced by
## coercion</pre>
retailHeirachical <- hclust(euclidean_distance, method="ward")
```

```
## The "ward" method has been renamed to "ward.D"; note new "ward.D2"
```

```
plot(retailHeirachical) # display dendogram
groups <- cutree(retailHeirachical, k=5) # cut tree into 5 clusters
# draw dendogram with red borders around the 5 clusters
rect.hclust(retailHeirachical, k=5, border="red")</pre>
```

### **Cluster Dendrogram**



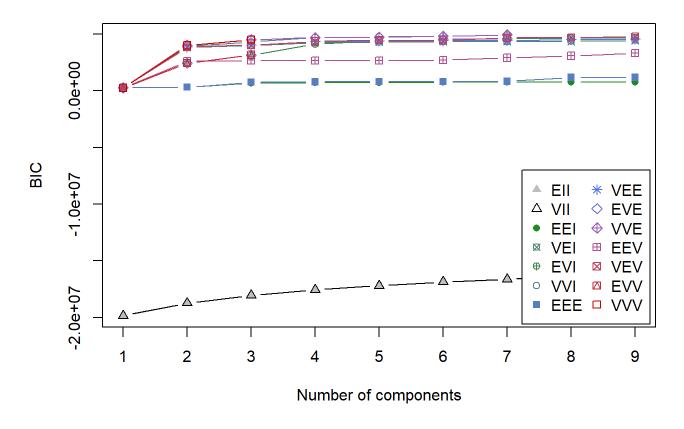
euclidean\_distance
hclust (\*, "ward.D")

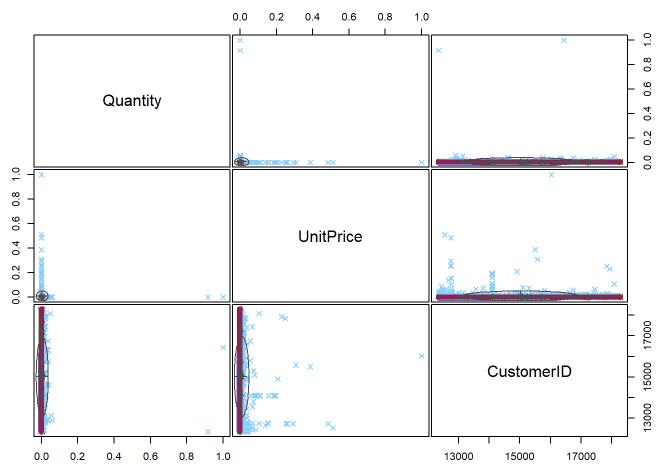
#### Model Based

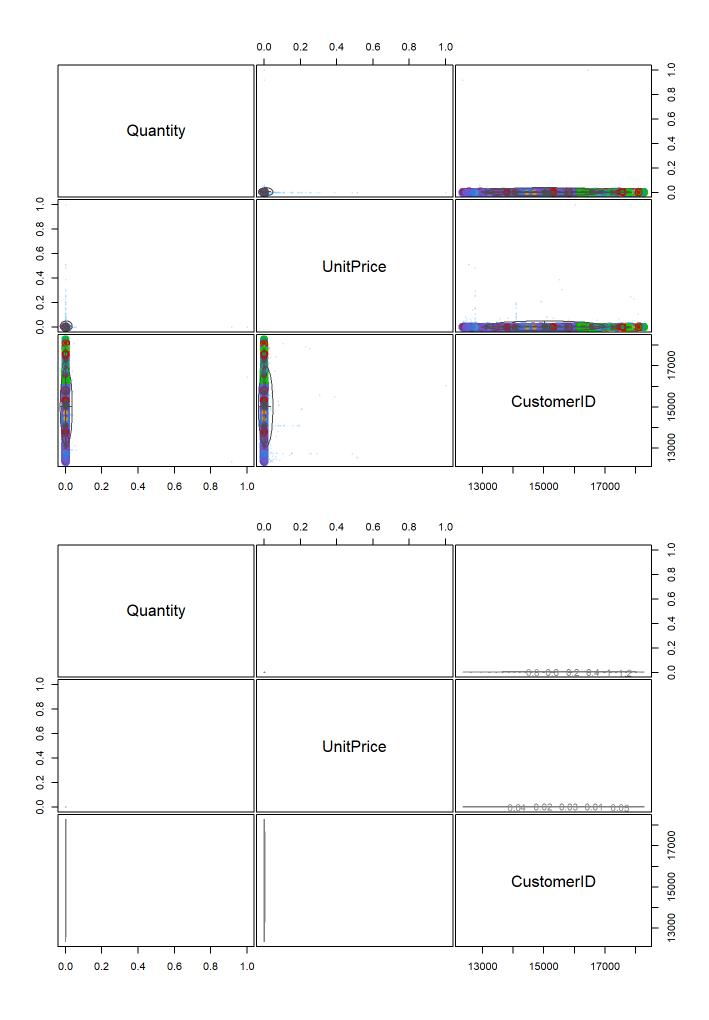
```
# Model Based Clustering
retailModel <- Mclust(data_norm)
summary(retailModel) # display the best model</pre>
```

```
## Gaussian finite mixture model fitted by EM algorithm
##
##
## Mclust VVE (ellipsoidal, equal orientation) model with 7 components:
##
                                 BIC
##
   log-likelihood
                        n df
           2467357 397884 51 4934057 4788163
##
##
## Clustering table:
##
             2
                   3
                               5
                                      6
## 91586 20463 52536 98097 85396 1722 48084
```

```
plot(retailModel)
```







The results for each of the models varied in its usefulness. The most useful I would say for this data set was the K means clustering algorithm. Thought I had to specify the number of clusters to use, the sum of squares algorithm identified three clusters that might be useful in 2, 3, and 4 clusters. 3 and 4 did not provide any useful information. But 2 classified information into 2 usable clusters that could be used with predictions. The heirarchical clustering is not useful in a dataset of this size. I attempted to cut the data set down to a 1/10th of its original size and still could not create a legible clustering graph. Going down to smaller sizes provided a more legible graph but due to the random nature of picking such a small number of values from the dataset, it did not provide a consistent method of identifying groups. Finally, the model based clustering was better than the heirachical but due to its more general nature, did not compare with the output produced by the K means cluster.