

Practicum-3.2

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```
#Importing Libraries  
library(fpc)
```

```
## Warning: package 'fpc' was built under R version 3.6.3
```

```
library(arules)
```

```
## Warning: package 'arules' was built under R version 3.6.3
```

```
## Loading required package: Matrix
```

```
##
```

```
## Attaching package: 'arules'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      abbreviate, write
```

```
#unloadNamespace("arules")  
#update.packages("arules")  
#library(arules)
```

---Problem 2---

1. Download the data set Plant Disease Data Set. Note that the data file does not contain header names; you may wish to add those. The description of each column can be found in the data set explanation. This assignment must be completed within a separate R Markdown Notebook. Use `read.transaction()` from the `arules` package to read the data.
 2. Explore the data set as you see fit and that allows you to get a sense of the data and get comfortable with it. Is there distributional skew in any of the features? Is there a need to apply a transform?
- I used `read.transactions` function to import the plants transaction data
 - Since this the data is transaction i.e sparse matrix we cannot observe the skewness or distribution of the data
 - Using summary and inspect I did explanatory analysis

```

#Importing Dataset
plant_data <- read.transactions("C:\\Users\\harsh\\Desktop\\Introduction to Machine learning and Data M

#Observing transactions
summary(plant_data)

```

```

## transactions as itemMatrix in sparse format with
## 34781 rows (elements/itemsets/transactions) and
## 34851 columns (items) and a density of 0.0002779312
##
## most frequent items:
##      ca      tx      or      az      fl (Other)
## 11676   8483   7028   6778   6621 296309
##
## element (itemset/transaction) length distribution:
## sizes
##      2      3      4      5      6      7      8      9     10     11     12     13     14
## 11566  4874  2954  2107  1366  1094   859   744   655   562   503   421   421
##      15     16     17     18     19     20     21     22     23     24     25     26     27
##      333    322    284    252    241    249    207    200    212    198    195    155    152
##      28     29     30     31     32     33     34     35     36     37     38     39     40
##      190    179    159    146    140    146    148    147    119    123    110    124    118
##      41     42     43     44     45     46     47     48     49     50     51     52     53
##      114     83     85    102     90     90     75     64     89     63     60     64     68
##      54     55     56     57     58     59     60     61     62     63     64     65     66
##      56     54     51     59     47     45     39     59     52     43     39     43     47
##      67     68     69     70
##      60     35     26     4
##
##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.
##      2.000   2.000   4.000   9.686  10.000  70.000
##
## includes extended item information - examples:
##           labels
## 1      *achnella
## 2 *achnella caduca
## 3      *agroelymus

```

```
inspect(plant_data[1:5])
```

```

##      items
## [1] {abelia,
##      fl,
##      nc}
## [2] {abelia x grandiflora,
##      fl,
##      nc}
## [3] {abelmoschus,
##      ct,
##      dc,
##      fl,
##      hi,

```

```

##      il,
##      ky,
##      la,
##      md,
##      mi,
##      ms,
##      nc,
##      pr,
##      sc,
##      va,
##      vi}
## [4] {abelmoschus esculentus,
##      ct,
##      dc,
##      fl,
##      il,
##      ky,
##      la,
##      md,
##      mi,
##      ms,
##      nc,
##      pr,
##      sc,
##      va,
##      vi}
## [5] {abelmoschus moschatus,
##      hi,
##      pr}

```

```

#Checking top 20 items using itemfrequency plot
itemFrequency(plant_data[,1:5])

```

```

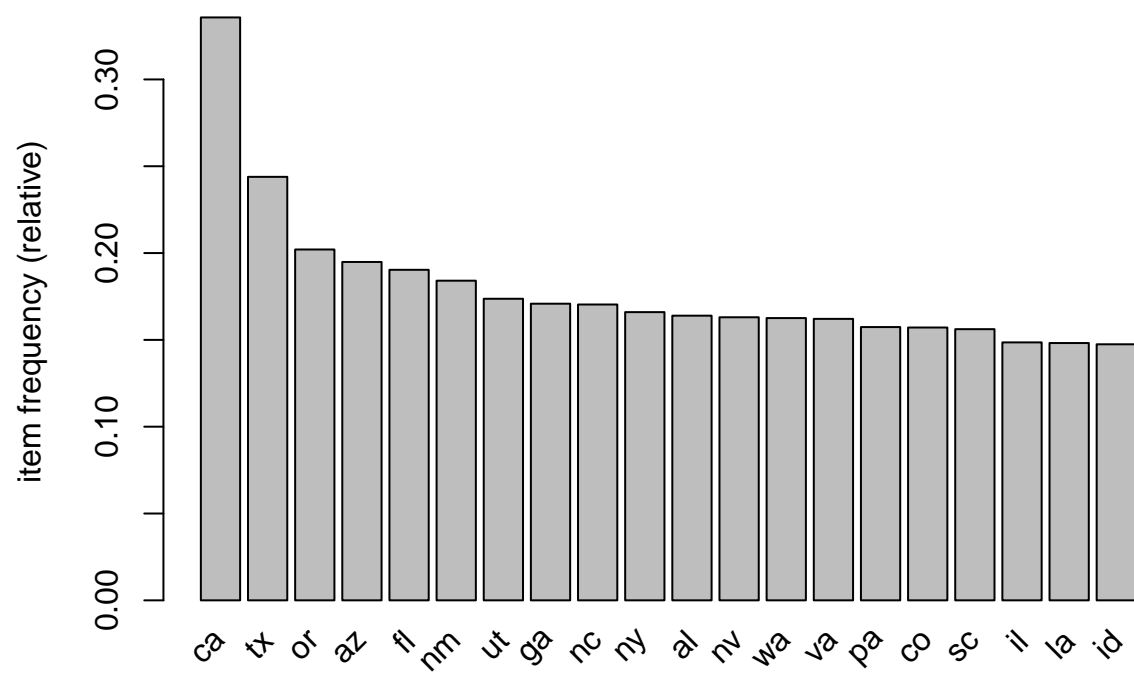
##          *achnella      *achnella caduca          *agroelymus
##          2.875133e-05      2.875133e-05      2.875133e-05
## *agroelymus adamsii *agroelymus bowdenii
##          2.875133e-05      2.875133e-05

```

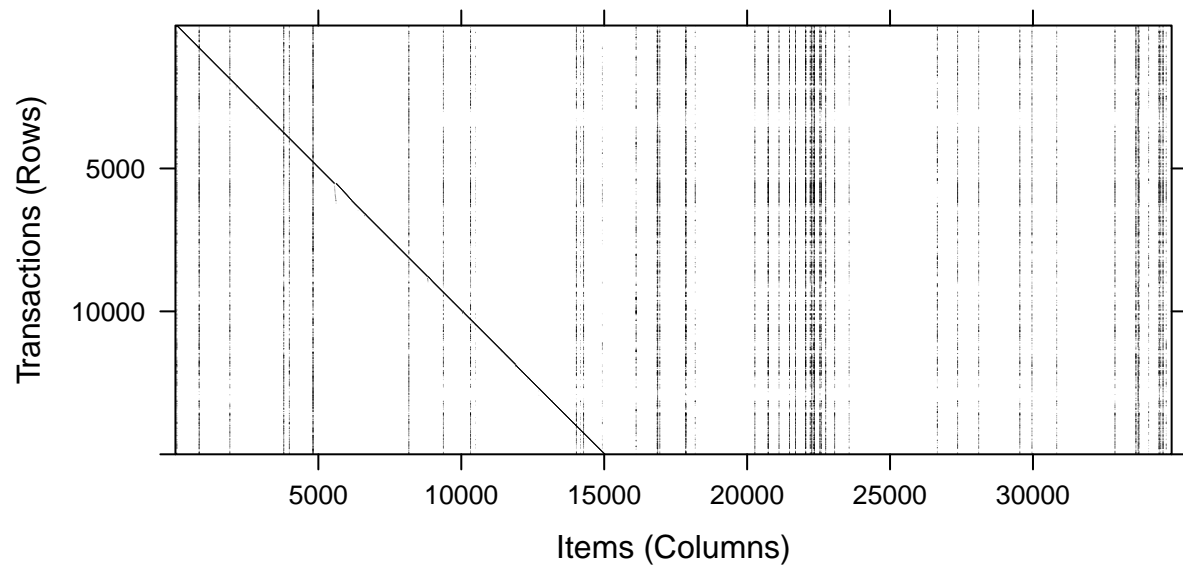
```

itemFrequencyPlot(plant_data,topN=20)

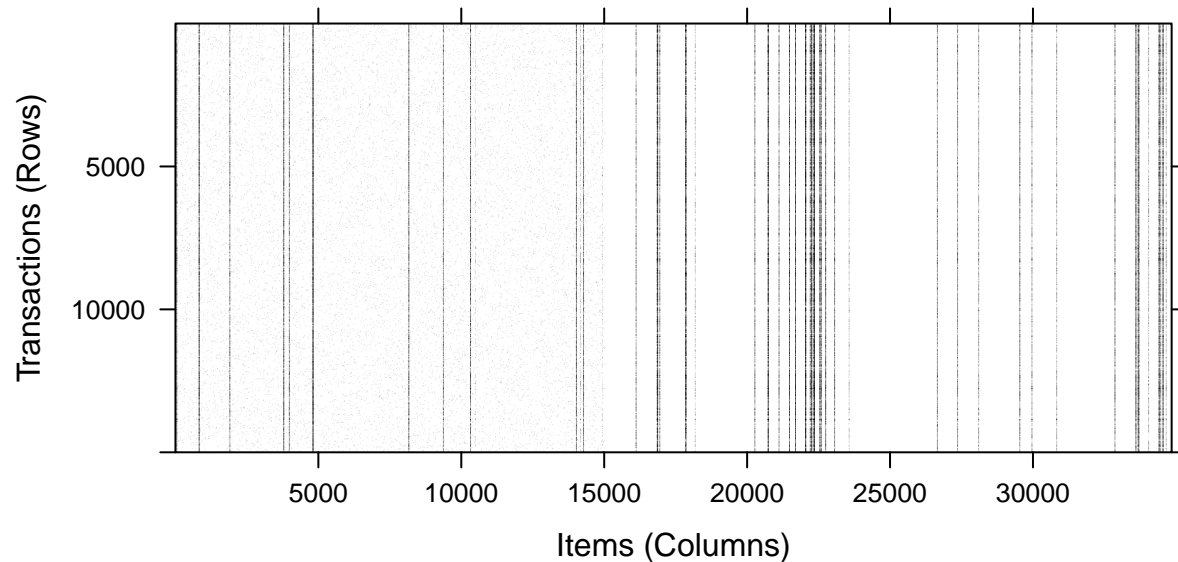
```



```
#Observing the image of the transaction data with and without sampling  
image(plant_data[1:15000])
```



```
image(sample(plant_data[1:15000]))
```



3. Use association rules to segment the data similar to what was done in Hämmäläinen, W., & Nykänen, M. (2008, December). Efficient discovery of statistically significant association rules. In Data Mining, 2008. ICDM'08. Eighth IEEE International Conference on (pp. 203-212). IEEE.

- Using apriori function to build association rules for plants dataset
- Improving the model by adjusting the support and confidence
- Using Inspect and summary, we can observe the output

```
#using apriori function to build rules
default_rules <- apriori(plant_data)
```

```
## Apriori
##
## Parameter specification:
## confidence minval smax arem aval originalSupport maxtime support minlen
##          0.8    0.1    1 none FALSE              TRUE     5     0.1     1
## maxlen target  ext
##          10  rules TRUE
##
## Algorithmic control:
## filter tree heap memopt load sort verbose
##    0.1 TRUE TRUE  FALSE TRUE    2    TRUE
##
## Absolute minimum support count: 3478
##
```

```
## set item appearances ...[0 item(s)] done [0.00s].
## set transactions ...[34851 item(s), 34781 transaction(s)] done [0.14s].
## sorting and recoding items ... [49 item(s)] done [0.01s].
## creating transaction tree ... done [0.01s].
## checking subsets of size 1 2 3 4 5 done [0.03s].
## writing ... [506 rule(s)] done [0.00s].
## creating S4 object ... done [0.00s].
```

```
summary(default_rules)
```

```
## set of 506 rules
##
## rule length distribution (lhs + rhs):sizes
##   2   3   4   5
## 89 307 100  10
##
##   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  2.000   3.000   3.000   3.061   3.000   5.000
##
## summary of quality measures:
##      support      confidence      coverage      lift
## Min.   :0.1001   Min.   :0.8001   Min.   :0.1015   Min.   :2.470
## 1st Qu.:0.1015   1st Qu.:0.8582   1st Qu.:0.1110   1st Qu.:5.449
## Median :0.1043   Median :0.8980   Median :0.1175   Median :5.793
## Mean   :0.1071   Mean   :0.8969   Mean   :0.1200   Mean   :5.767
## 3rd Qu.:0.1100   3rd Qu.:0.9414   3rd Qu.:0.1247   3rd Qu.:6.092
## Max.   :0.1426   Max.   :0.9893   Max.   :0.1708   Max.   :7.424
##      count
## Min.   :3480
## 1st Qu.:3532
## Median :3626
## Mean   :3726
## 3rd Qu.:3825
## Max.   :4959
##
## mining info:
##      data ntransactions support confidence
## plant_data      34781      0.1      0.8
```

```
inspect(default_rules[1:5])
```

```
##      lhs      rhs support confidence coverage lift      count
## [1] {nv} => {ca} 0.1351600 0.8291005 0.1630200 2.469762 4701
## [2] {wy} => {mt} 0.1098013 0.8108280 0.1354188 5.875294 3819
## [3] {wy} => {co} 0.1104339 0.8154989 0.1354188 5.190095 3841
## [4] {mt} => {id} 0.1123027 0.8137500 0.1380064 5.518237 3906
## [5] {id} => {or} 0.1207268 0.8186781 0.1474656 4.051571 4199
```

```
#Improving rules by changing default parameters
```

```
new_rules <- apriori(plant_data, parameter = list (support = 0.001, confidence = 0.75))
```

```
## Apriori
```

```
##
## Parameter specification:
## confidence minval smax arem aval originalSupport maxtime support minlen
##      0.75    0.1    1 none FALSE          TRUE      5  0.001    1
## maxlen target  ext
##      10   rules TRUE
##
## Algorithmic control:
## filter tree heap memopt load sort verbose
##      0.1 TRUE TRUE  FALSE TRUE    2    TRUE
##
## Absolute minimum support count: 34
##
## set item appearances ...[0 item(s)] done [0.00s].
## set transactions ...[34851 item(s), 34781 transaction(s)] done [0.12s].
## sorting and recoding items ... [70 item(s)] done [0.01s].
## creating transaction tree ... done [0.01s].
## checking subsets of size 1 2 3 4 5

## Warning in apriori(plant_data, parameter = list(support = 0.001, confidence =
## 0.75)): Mining stopped (time limit reached). Only patterns up to a length of 5
## returned!

## done [17.61s].
## writing ... [43995074 rule(s)] done [5.66s].
## creating S4 object ... done [13.08s].
```

```
summary(new_rules)
```

```
## set of 43995074 rules
##
## rule length distribution (lhs + rhs):sizes
##      2      3      4      5
##    469    68831  2201628 41724146
##
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##    2.000  5.000   5.000   4.947   5.000   5.000
##
## summary of quality measures:
##      support      confidence      coverage      lift
##    Min.   :0.001006   Min.   :0.7500   Min.   :0.001006   Min.   : 2.234
##    1st Qu.:0.007360   1st Qu.:0.8729   1st Qu.:0.007935   1st Qu.: 6.013
##    Median :0.015411   Median :0.9399   Median :0.016791   Median : 7.018
##    Mean   :0.017252   Mean   :0.9202   Mean   :0.018989   Mean   : 7.515
##    3rd Qu.:0.024381   3rd Qu.:0.9788   3rd Qu.:0.027055   3rd Qu.: 8.652
##    Max.   :0.158765   Max.   :1.0000   Max.   :0.202064   Max.   :67.350
##      count
##    Min.   : 35
##    1st Qu.: 256
##    Median : 536
##    Mean   : 600
##    3rd Qu.: 848
##    Max.   :5522
```



```
##
## mining info:
##      data ntransactions support confidence
## plant_data      34781    0.001      0.75
```

```
inspect(new_rules[1:5])
```

```
##      lhs      rhs      support      confidence coverage      lift      count
## [1] {gl}    => {qc}    0.01236307 0.7664884 0.01612950 6.240457 430
## [2] {dengl} => {lb}    0.01204681 0.8747390 0.01377189 21.231192 419
## [3] {dengl} => {fraspm} 0.01377189 1.0000000 0.01377189 28.744628 479
## [4] {dengl} => {yt}    0.01049424 0.7620042 0.01377189 12.620603 365
## [5] {dengl} => {nt}    0.01081050 0.7849687 0.01377189 13.489128 376
```

4. Are there clusters in the data? Can plants be segmented into groups? Build a k-means clustering model to investigate.

- First I converted the transactions to separate items and transaction ID by importing it again
- Later I stored this data in a dataframe in matrix format
- Using kmeans function, I built clusters of the data by using $k = 10$ for creating 10 clusters
- Using size and centers we can observe the center and size of the cluster
- Using plotCluster function I visualize the clusters
- For this problem, I referred a few links I have added them in this chunk

```
#Importing the data as separate columns
```

```
plants <- read.transactions("C:\\Users\\harsh\\Desktop\\Introduction to Machine learning and Data Mining\\")
```

```
#Observing the new data
```

```
summary(plants)
```

```
## transactions as itemMatrix in sparse format with
## 34781 rows (elements/itemsets/transactions) and
## 70 columns (items) and a density of 0.1240883
##
```

```
## most frequent items:
```

```
##      ca      tx      or      az      fl (Other)
## 11676    8483    7028    6778    6621 261528
```

```
##
```

```
## element (itemset/transaction) length distribution:
```

```
## sizes
```

```
##      1      2      3      4      5      6      7      8      9     10     11     12     13
## 11566 4874 2954 2107 1366 1094 859 744 655 562 503 421 421
## 14     15     16     17     18     19     20     21     22     23     24     25     26
## 333   322   284   252   241   249   207   200   212   198   195   155   152
## 27     28     29     30     31     32     33     34     35     36     37     38     39
## 190   179   159   146   140   146   148   147   119   123   110   124   118
## 40     41     42     43     44     45     46     47     48     49     50     51     52
## 114    83    85    102    90    90    75    64    89    63    60    64    68
## 53     54     55     56     57     58     59     60     61     62     63     64     65
## 56     54     51     59     47     45     39     59     52     43     39     43     47
## 66     67     68     69
## 60     35     26     4
```

```
##
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##    1.000   1.000   3.000   8.686   9.000  69.000
##
## includes extended item information - examples:
##   labels
## 1      ab
## 2      ak
## 3      al
##
## includes extended transaction information - examples:
##           transactionID
## 1                abelia
## 2 abelia x grandiflora
## 3                abelmoschus
```

```
#Converting the data to matrix format and as integer datatype
plants_matrix <- as.data.frame(as(plants,"matrix"))
plants_matrix[,1:70] <- lapply(plants_matrix[,1:70],as.integer)
```

```
#Verifying the data
str(plants_matrix)
```

```
## 'data.frame':   34781 obs. of  70 variables:
## $ ab      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ ak      : int  0 0 0 0 0 1 0 1 0 0 ...
## $ al      : int  0 0 0 0 0 0 0 0 0 0 ...
## $ ar      : int  0 0 0 0 0 0 0 0 0 0 ...
## $ az      : int  0 0 0 0 0 1 0 0 0 0 ...
## $ bc      : int  0 0 0 0 0 1 0 1 0 0 ...
## $ ca      : int  0 0 0 0 0 1 0 1 0 0 ...
## $ co      : int  0 0 0 0 0 1 0 0 0 0 ...
## $ ct      : int  0 0 1 1 0 1 0 0 1 1 ...
## $ dc      : int  0 0 1 1 0 0 0 0 0 0 ...
## $ de      : int  0 0 0 0 0 0 0 0 0 0 ...
## $ deng1   : int  0 0 0 0 0 0 0 0 0 0 ...
## $ fl      : int  1 1 1 1 0 0 0 0 0 0 ...
## $ frasp   : int  0 0 0 0 0 1 0 0 1 1 ...
## $ ga      : int  0 0 0 0 0 1 0 0 0 0 ...
## $ gl      : int  0 0 0 0 0 0 0 0 0 0 ...
## $ hi      : int  0 0 1 0 1 0 0 0 0 0 ...
## $ ia      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ id      : int  0 0 0 0 0 1 0 0 0 0 ...
## $ il      : int  0 0 1 1 0 0 0 0 0 0 ...
## $ in      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ ks      : int  0 0 0 0 0 0 0 0 0 0 ...
## $ ky      : int  0 0 1 1 0 0 0 0 0 0 ...
## $ la      : int  0 0 1 1 0 0 0 0 0 0 ...
## $ lb      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ ma      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ mb      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ md      : int  0 0 1 1 0 1 0 0 1 1 ...
## $ me      : int  0 0 0 0 0 1 0 0 1 1 ...
## $ mi      : int  0 0 1 1 0 1 0 0 1 1 ...
```

```
## $ mn      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ mo      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ ms      : int 0 0 1 1 0 0 0 0 0 0 ...
## $ mt      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ nb      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ nc      : int 1 1 1 1 0 1 1 0 0 0 ...
## $ nd      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ ne      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ nf      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ nh      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ nj      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ nm      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ ns      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ nt      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ nu      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ nv      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ ny      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ oh      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ ok      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ on      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ or      : int 0 0 0 0 0 1 0 1 0 0 ...
## $ pa      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ pe      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ pr      : int 0 0 1 1 1 0 0 0 0 0 ...
## $ qc      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ ri      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ sc      : int 0 0 1 1 0 0 0 0 0 0 ...
## $ sd      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ sk      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ tn      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ tx      : int 0 0 0 0 0 0 0 0 0 0 ...
## $ ut      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ va      : int 0 0 1 1 0 1 0 0 1 1 ...
## $ vi      : int 0 0 1 1 0 0 0 0 0 0 ...
## $ vt      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ wa      : int 0 0 0 0 0 1 0 1 0 0 ...
## $ wi      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ wv      : int 0 0 0 0 0 1 0 0 1 1 ...
## $ wy      : int 0 0 0 0 0 1 0 0 0 0 ...
## $ yt      : int 0 0 0 0 0 1 0 0 0 0 ...
```

```
#Creating the clusters using kmeans function
clusters <- kmeans(plants_matrix,10)
clusters$size
```

```
## [1] 2097 2454 12769 6474 1346 2009 1376 2582 1757 1917
```

```
clusters$centers
```

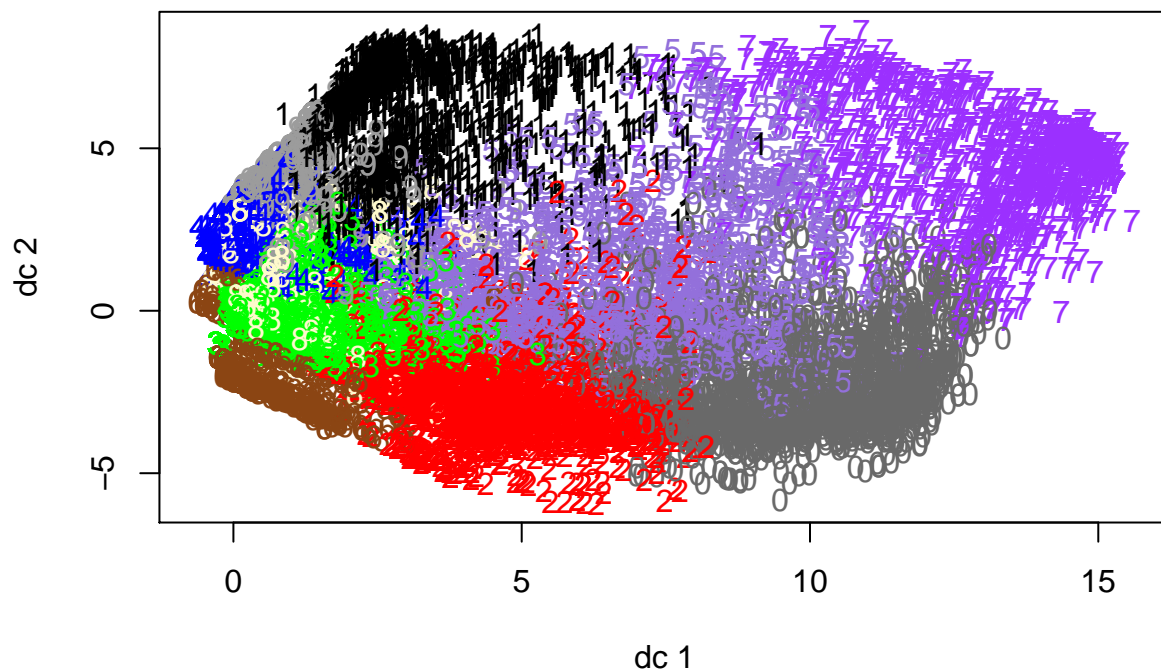
```
##          ab          ak          al          ar          az          bc
## 1  0.695278970 0.3681449690 0.009060563 0.042918455 0.44110634 0.7758702909
## 2  0.003259984 0.0020374898 0.901792991 0.473512632 0.10024450 0.0199674002
## 3  0.024590806 0.0689952228 0.033440363 0.022867883 0.05059128 0.0517659958
```

## 4	0.005560704	0.0194624652	0.004633920	0.002316960	0.06889095	0.0755329008
## 5	0.292719168	0.2555720654	0.101040119	0.053491828	0.03417533	0.4108469539
## 6	0.000000000	0.0009955202	0.057242409	0.004977601	0.04330513	0.0009955202
## 7	0.806686047	0.5813953488	0.702761628	0.743459302	0.79360465	0.9091569767
## 8	0.006196747	0.0007745933	0.013942680	0.077846631	0.66227730	0.0023237800
## 9	0.002276608	0.0056915196	0.005691520	0.016505407	0.73875925	0.0313033580
## 10	0.035472092	0.0140845070	0.912363067	0.895148670	0.14710485	0.0949400104
##	ca	co	ct	dc	de	dengl
## 1	0.65951359	0.7663328565	0.025274201	0.008583691	0.008106819	0.013352408
## 2	0.14384678	0.0346373268	0.077424613	0.109209454	0.188264059	0.000000000
## 3	0.000000000	0.0305427207	0.017699115	0.006265173	0.007674837	0.001801237
## 4	1.000000000	0.0038616002	0.005715168	0.001390176	0.001699104	0.000463392
## 5	0.20059435	0.1560178306	0.797919762	0.195393759	0.320208024	0.101783061
## 6	0.04629169	0.0004977601	0.002986560	0.006470881	0.000000000	0.000000000
## 7	0.89098837	0.9353197674	0.950581395	0.725290698	0.795784884	0.209302326
## 8	0.08520527	0.3508907823	0.001936483	0.001161890	0.001936483	0.000000000
## 9	0.72737621	0.3932840068	0.004553216	0.002276608	0.001138304	0.000000000
## 10	0.19770475	0.1371935316	0.774126239	0.742827334	0.787167449	0.000000000
##	fl	fraspm	ga	gl	hi	ia
## 1	0.006199332	0.021459227	0.011444921	0.077730091	0.01430615	0.140677158
## 2	0.788508557	0.000407498	0.902200489	0.000407498	0.11776691	0.042787286
## 3	0.093351085	0.004777195	0.043308012	0.020675072	0.16477406	0.011355627
## 4	0.012511585	0.001390176	0.004324992	0.001235712	0.02301514	0.002008032
## 5	0.038632987	0.312778603	0.155274889	0.037890045	0.02748886	0.329866270
## 6	0.608262817	0.000000000	0.062220010	0.000000000	0.24240916	0.000000000
## 7	0.577034884	0.421511628	0.726017442	0.049418605	0.34229651	0.943313953
## 8	0.024012393	0.000000000	0.006196747	0.000000000	0.00929512	0.033307514
## 9	0.013659647	0.000000000	0.007398976	0.000000000	0.02105862	0.001707456
## 10	0.649973918	0.048513302	0.918622848	0.003129890	0.09181012	0.658841941
##	id	il	in	ks	ky	la
## 1	0.8950882213	0.116833572	0.0557939914	0.172627563	0.021459227	0.023843586
## 2	0.0114099430	0.238386308	0.1483292584	0.142216789	0.346780766	0.740016300
## 3	0.0422116062	0.021379904	0.0144882136	0.009476075	0.016367766	0.030307777
## 4	0.0586963238	0.004324992	0.0009267841	0.001235712	0.002162496	0.008958913
## 5	0.2169390788	0.560921248	0.4680534918	0.077265973	0.284546805	0.040861813
## 6	0.0004977601	0.001991040	0.0000000000	0.001493280	0.002488800	0.100049776
## 7	0.9367732558	0.984011628	0.9527616279	0.821220930	0.852470930	0.670058140
## 8	0.0081332301	0.039116964	0.0151045701	0.181642138	0.008520527	0.061967467
## 9	0.3386454183	0.008537279	0.0028457598	0.042117245	0.001707456	0.017643711
## 10	0.0558163798	0.942618675	0.9300991132	0.651538863	0.965049557	0.768909755
##	lb	ma	mb	md	me	mi
## 1	0.1092036242	0.061516452	0.323319027	0.027181688	0.082021936	0.1468764902
## 2	0.0008149959	0.107986960	0.003259984	0.345558272	0.027302363	0.0383048085
## 3	0.0184039471	0.031404182	0.018560576	0.022789568	0.020910016	0.0240426032
## 4	0.0015446401	0.018690145	0.003398208	0.007568736	0.004170528	0.0046339203
## 5	0.3112927192	0.869985141	0.445765230	0.531203566	0.852897474	0.8068350669
## 6	0.0000000000	0.006968641	0.000000000	0.017421603	0.000000000	0.0009955202
## 7	0.3626453488	0.962209302	0.848110465	0.918604651	0.926598837	0.9811046512
## 8	0.0000000000	0.009682417	0.006196747	0.014717273	0.005809450	0.0123934934
## 9	0.0000000000	0.015367103	0.000569152	0.013659647	0.005122368	0.0062606716
## 10	0.0203442879	0.775169536	0.153364632	0.932185707	0.515910276	0.7897756912
##	mn	mo	ms	mt	nb	nc
## 1	0.227467811	0.092513114	0.01144492	0.9713876967	0.0629470672	0.013352408
## 2	0.015077425	0.298288509	0.77791361	0.0126324368	0.0052974735	0.781581092

## 3	0.014174955	0.018638891	0.01417495	0.0335969927	0.0116688856	0.034145195
## 4	0.001390176	0.004016064	0.00308928	0.0154464010	0.0018535681	0.006641952
## 5	0.602526003	0.208766716	0.03789004	0.2637444279	0.6901931649	0.377414562
## 6	0.000000000	0.002488800	0.03235441	0.0009955202	0.0000000000	0.014932802
## 7	0.963662791	0.880087209	0.64970930	0.9636627907	0.8241279070	0.842296512
## 8	0.020914020	0.072037180	0.02052672	0.0395042603	0.0003872967	0.005422153
## 9	0.001707456	0.018212863	0.00569152	0.1548093341	0.0005691520	0.003414912
## 10	0.537819510	0.904016693	0.83881064	0.0761606677	0.2529994784	0.930620762
##	nd	ne	nf	nh	nj	nm
## 1	0.3290414878	0.290414878	0.105388650	0.049594659	0.025274201	0.522174535
## 2	0.0073349633	0.035452323	0.002852486	0.025264874	0.216381418	0.100651997
## 3	0.0056386561	0.007126635	0.020283499	0.011590571	0.019735296	0.011042368
## 4	0.0009267841	0.000617856	0.002316960	0.001390176	0.009113377	0.001699104
## 5	0.2043090639	0.115156018	0.514115899	0.779346211	0.684249629	0.061664190
## 6	0.0009955202	0.000000000	0.000000000	0.000000000	0.007964161	0.019910403
## 7	0.8604651163	0.871366279	0.615552326	0.902616279	0.947674419	0.872093023
## 8	0.0251742835	0.099535244	0.000000000	0.000000000	0.004647560	0.929899303
## 9	0.0039840637	0.025611838	0.000000000	0.000000000	0.004553216	0.459305635
## 10	0.1893583725	0.435054773	0.076682316	0.532603026	0.869066249	0.197183099
##	ns	nt	nu	nv	ny	oh
## 1	0.056270863	0.2665712923	0.113018598	0.67858846	0.113972341	0.051502146
## 2	0.015077425	0.0008149959	0.000407498	0.02770986	0.165036675	0.134881826
## 3	0.012138774	0.0364946354	0.023337771	0.01777743	0.046283969	0.020440128
## 4	0.003398208	0.0023169601	0.001390176	0.16203275	0.014519617	0.003707136
## 5	0.685735513	0.2243684993	0.138930163	0.03863299	0.955423477	0.621842496
## 6	0.000000000	0.000000000	0.000000000	0.00248880	0.006470881	0.006470881
## 7	0.809593023	0.4854651163	0.177325581	0.74345930	0.992005814	0.974563953
## 8	0.000000000	0.0003872967	0.000000000	0.03253292	0.006971340	0.009295120
## 9	0.001138304	0.0011383039	0.000000000	0.93454752	0.023904382	0.005122368
## 10	0.246739697	0.0046948357	0.001564945	0.05059990	0.897235263	0.951486698
##	ok	on	or	pa	pe	pr
## 1	0.135431569	0.253695756	0.798760134	0.048640916	0.019551741	0.001430615
## 2	0.326405868	0.032599837	0.060717196	0.219233904	0.001629992	0.182966585
## 3	0.021928107	0.041663404	0.059284204	0.033440363	0.003837419	0.135484376
## 4	0.002316960	0.006796416	0.295180723	0.014674081	0.001390176	0.004633920
## 5	0.028231798	0.868499257	0.299405646	0.795690936	0.451708767	0.014858841
## 6	0.006968641	0.001493280	0.003484321	0.011448482	0.000000000	0.988053758
## 7	0.736191860	0.980377907	0.932412791	0.983284884	0.655523256	0.199854651
## 8	0.244771495	0.002323780	0.007358637	0.006196747	0.000000000	0.013555383
## 9	0.075128059	0.006829824	0.317017644	0.013659647	0.001138304	0.011383039
## 10	0.752217006	0.699530516	0.139280125	0.952008346	0.117892540	0.122065728
##	qc	ri	sc	sd	sk	tn
## 1	0.2060085837	0.009537434	0.020028612	0.407725322	0.433953267	0.017644254
## 2	0.0097799511	0.048899756	0.834148329	0.017522412	0.002852486	0.491850041
## 3	0.0448743050	0.006030229	0.026548673	0.006030229	0.012686976	0.018638891
## 4	0.0052517763	0.000308928	0.006641952	0.000772320	0.002008032	0.001699104
## 5	0.8142644874	0.549777117	0.152303120	0.169390788	0.343982169	0.277117385
## 6	0.000000000	0.000000000	0.026879044	0.000000000	0.000000000	0.003484321
## 7	0.9389534884	0.859011628	0.710755814	0.899709302	0.825581395	0.834302326
## 8	0.0003872967	0.001161890	0.012780790	0.068164214	0.008520527	0.010457010
## 9	0.0011383039	0.002276608	0.009106431	0.020489471	0.002845760	0.001138304
## 10	0.4267083985	0.598330725	0.873761085	0.274908712	0.066770996	0.965049557
##	tx	ut	va	vi	vt	wa
## 1	0.18454936	0.751549833	0.023843586	0.0009537434	0.0548402480	0.820219361

```
## 2 0.62184189 0.046454768 0.578239609 0.1022819886 0.0126324368 0.032599837
## 3 0.12444201 0.046049025 0.023885974 0.0048555094 0.0132351789 0.054037121
## 4 0.01513747 0.005715168 0.004324992 0.0006178560 0.0033982082 0.136546185
## 5 0.03789004 0.121099554 0.514858841 0.0052005944 0.7971768202 0.355126300
## 6 0.15181682 0.001991040 0.007964161 0.7849676456 0.0000000000 0.001991040
## 7 0.74491279 0.861918605 0.916424419 0.1010174419 0.9156976744 0.951308140
## 8 0.68048025 0.187838885 0.004260263 0.0061967467 0.0003872967 0.002711077
## 9 0.22936824 0.963005122 0.003984064 0.0073989755 0.0022766079 0.137734775
## 10 0.70005216 0.102243088 0.964006260 0.0594679186 0.5414710485 0.125195618
##      wi      wv      wy      yt
## 1 0.141154030 0.0157367668 0.880305198 0.3171196948
## 2 0.027302363 0.1572942135 0.010187449 0.0004074980
## 3 0.020283499 0.0118255149 0.030777665 0.0402537395
## 4 0.002625888 0.0013901761 0.005560704 0.0035526722
## 5 0.720653789 0.4316493314 0.176077266 0.1827637444
## 6 0.001493280 0.0024888004 0.000000000 0.0000000000
## 7 0.983284884 0.8706395349 0.906250000 0.4622093023
## 8 0.017041053 0.0003872967 0.127807901 0.0003872967
## 9 0.002845760 0.0022766079 0.253272624 0.0011383039
## 10 0.681794471 0.8836724048 0.078768910 0.0062597809
```

```
#Visualizing the clusters
plotcluster(plants_matrix,clusters$cluster)
```



Reference: <https://stats.stackexchange.com/questions/31083/how-to-produce-a-pretty-plot-of-the-results-of-k-means-cluster-analysis> <https://stackoverflow.com/questions/41972270/how-to-convert-object-of-transaction-to-dataframe-in-r>