Mtcars analysis

Wojciech Kolasa

2025-07-23

cat('\n')

apply(mtcars[,] , 2 ,skewness)

The data comes from R's built-in dataset called mtcars. Each observation consists of 11 features describing various parameters of selected car models. The aim of this project is to perform both supervised and unsupervised classification and to describe each variable.

Analysis Plan:

- 1. Presentation of summary statistics including histograms
- 2. Univariate analysis of outliers and multimodality
- 3. Correlation analysis and dimensionality reduction

```
4. Clustering analysis
 5. Supervised classification using random forests
cat('Basic statistical measures','\n')
## Basic statistical measures
cat('\n')
apply(mtcars[,], 2 ,summary)
##
                               disp
                                                 drat
                mpg
                       cyl
                                          hp
                                                            wt
                                                                   qsec
## Min.
           10.40000 4.0000 71.1000
                                     52.0000 2.760000 1.51300 14.50000 0.0000
## 1st Qu. 15.42500 4.0000 120.8250
                                     96.5000 3.080000 2.58125 16.89250 0.0000
## Median 19.20000 6.0000 196.3000 123.0000 3.695000 3.32500 17.71000 0.0000
           20.09062 6.1875 230.7219 146.6875 3.596563 3.21725 17.84875 0.4375
## 3rd Qu. 22.80000 8.0000 326.0000 180.0000 3.920000 3.61000 18.90000 1.0000
## Max.
           33.90000 8.0000 472.0000 335.0000 4.930000 5.42400 22.90000 1.0000
##
                am
                    gear
        0.00000 3.0000 1.0000
## Min.
## 1st Qu. 0.00000 3.0000 2.0000
## Median 0.00000 4.0000 2.0000
## Mean 0.40625 3.6875 2.8125
## 3rd Qu. 1.00000 4.0000 4.0000
## Max.
         1.00000 5.0000 8.0000
cat('\n')
cat('Standard deviation','\n')
## Standard deviation
cat('\n')
apply(mtcars[,] , 2 , sd)
##
           mpg
                       cyl
                                  disp
                                                                  0.9784574
##
     6.0269481
                 1.7859216 123.9386938 68.5628685
                                                     0.5346787
##
          asec
                                    am
                                                           carb
                        ٧S
                                              gear
     1.7869432
                 0.5040161
                             0.4989909
                                         0.7378041
                                                      1.6152000
cat('\n')
cat('Skewness','\n')
## Skewness
```

```
##
          mpg
                     cyl
                               disp
                                            hp
                                                      drat
##
   0.6404399 -0.1831287
                          0.4002724
                                     0.7614356  0.2788734  0.4437855  0.3870456
##
                               gear
                                          carb
##
   0.2519763 0.3817709 0.5546495 1.1021304
cat('\n')
cat('Kurtosis','\n')
## Kurtosis
cat('\n')
apply(mtcars[,] , 2 ,kurtosis)
                 cyl
                         disp
                                    hp
       mpg
                                                              qsec
## 2.799467 1.319032 1.910317 3.052233 2.435116 3.172471 3.553753 1.063492
##
                gear
                         carb
        am
## 1.145749 2.056790 4.536121
```

Summary and Conclusions:

The variables disp and hp have the highest minimum values and also the highest mean and median. They also have high standard deviation values, indicating low concentration around the mean. The rest of the variables show low standard deviation, suggesting high concentration around the mean. All variables have kurtosis greater than 0, indicating leptokurtic distributions. The variable carb has the highest kurtosis and the strongest presence of extreme values.

```
for(i in 1:11) print(grubbs.test(mtcars[,i]))
```

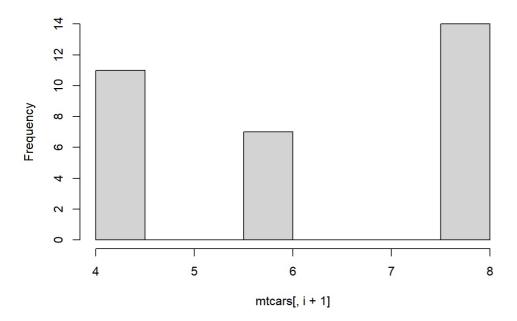
```
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 2.29127, U = 0.82518, p-value = 0.276
## alternative hypothesis: highest value 33.9 is an outlier
##
##
##
    Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 1.22486, U = 0.95004, p-value = 1
## alternative hypothesis: lowest value 4 is an outlier
##
##
##
    Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 1.9468, U = 0.8738, p-value = 0.7363
## alternative hypothesis: highest value 472 is an outlier
##
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 2.74657, U = 0.74881, p-value = 0.05564
## alternative hypothesis: highest value 335 is an outlier
##
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 2.4939, U = 0.7929, p-value = 0.1419
## alternative hypothesis: highest value 4.93 is an outlier
##
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 2.25534, U = 0.83063, p-value = 0.3083
## alternative hypothesis: highest value 5.424 is an outlier
##
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 2.82675, U = 0.73393, p-value = 0.04021
## alternative hypothesis: highest value 22.9 is an outlier
##
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 1.11604, U = 0.95853, p-value = 1
## alternative hypothesis: highest value 1 is an outlier
##
##
##
   Grubbs test for one outlier
## data: mtcars[, i]
## G = 1.18990, U = 0.95285, p-value = 1
## alternative hypothesis: highest value 1 is an outlier
##
##
    Grubbs test for one outlier
##
##
## data: mtcars[, i]
## G = 1.77893, U = 0.89462, p-value = 1
## alternative hypothesis: highest value 5 is an outlier
##
##
##
   Grubbs test for one outlier
##
## data: mtcars[, i]
## G = 3.21168, U = 0.65653, p-value = 0.006787
## alternative hypothesis: highest value 8 is an outlier
```

Conclusions:

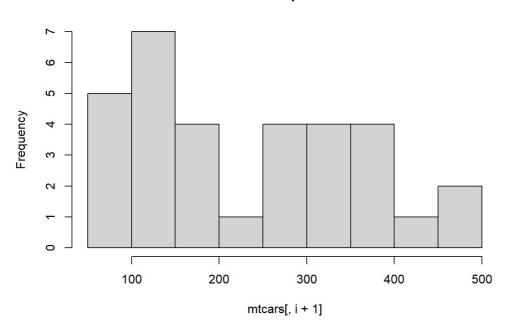
For the variables <code>qsec</code> and <code>carb</code>, the Grubbs test p-value is less than 0.05, indicating the presence of outliers that must be considered during clustering analysis.

for(i in 1:10) hist(mtcars[,i+1], main=colnames(mtcars)[i+1])

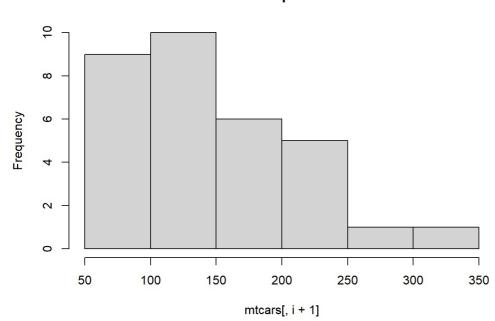




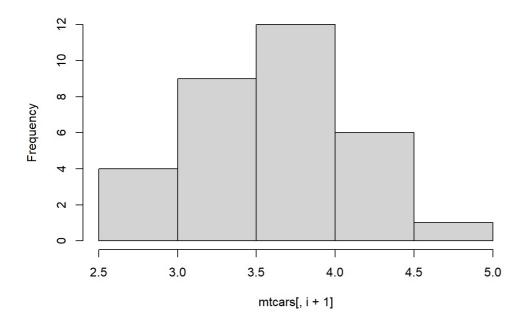
disp

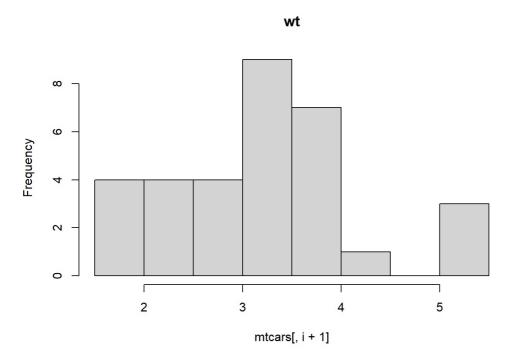


hp

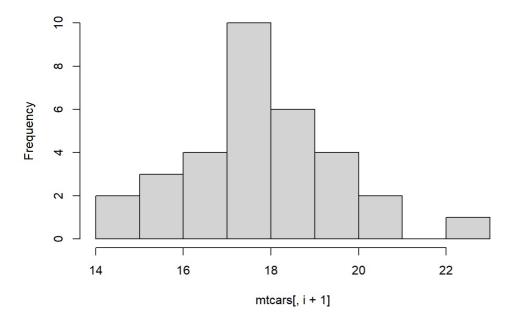




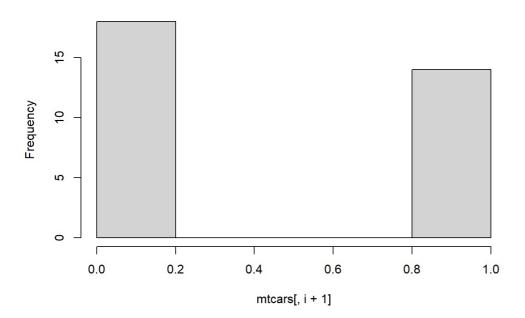


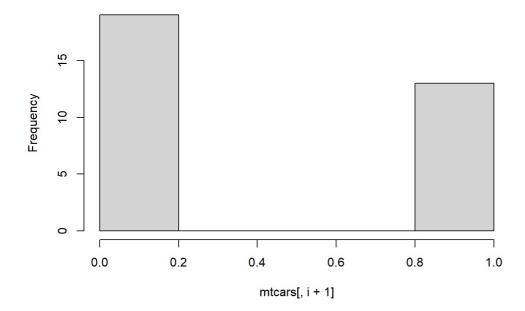


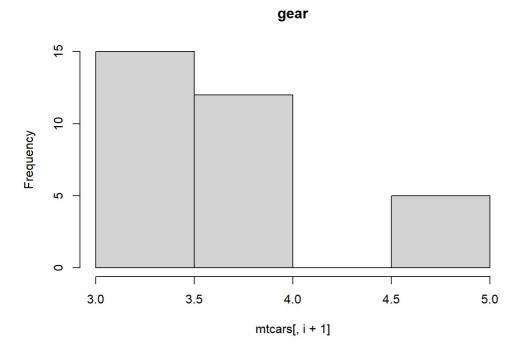




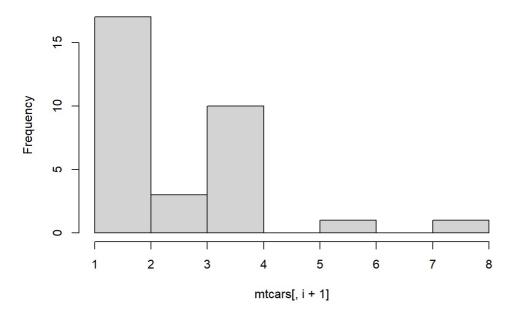
٧s







carb



Conclusions:

Histogram analysis suggests that normality should be tested for hp and drat , and multimodality for disp and wt .

Normality Tests:

```
shapiro.test(mtcars$hp)

##
## Shapiro-Wilk normality test
##
## data: mtcars$hp
## W = 0.93342, p-value = 0.04881

shapiro.test(mtcars$drat)

##
## Shapiro-Wilk normality test
##
## data: mtcars$drat
##
## at a: mtcars$drat
##
## W = 0.94588, p-value = 0.1101
```

Conclusions:

The hypothesis of normality for variables hp and drat is rejected.

Multimodality Analysis:

```
silverman.test(mtcars$disp,k=1)

## Silvermantest: Testing the null hypothesis that the number of modes is <= 1
## The resulting p-value is 0.1781782

silverman.test(mtcars$wt,k=1)

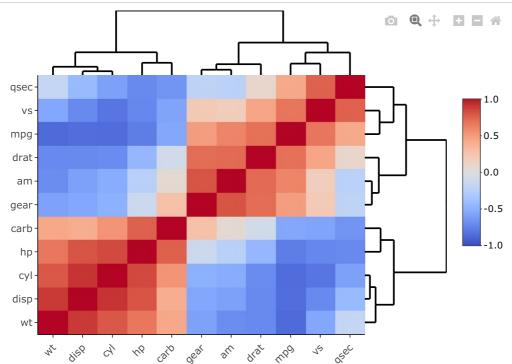
## Silvermantest: Testing the null hypothesis that the number of modes is <= 1
## The resulting p-value is 0.08608609</pre>
```

Conclusions:

The distributions of disp and wt are not multimodal, as the significance level exceeds 5%.

Correlation Analysis using Pearson's coefficient

heatmaply_cor(cor(mtcars[,],method='pearson'))

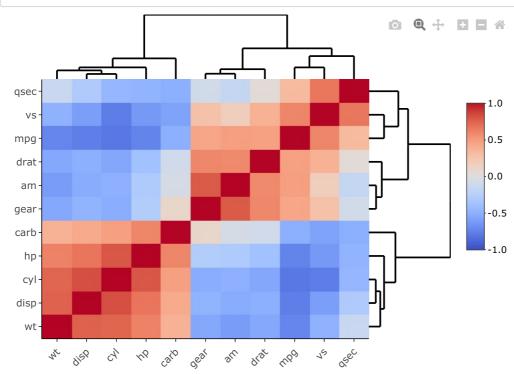


Conclusions:

Positive correlations are observed within the groups: (wt, disp, cyl, hp), (gear, am, drat), and (vs, qsec). A weaker positive correlation is present between vs and carb.

Correlation Analysis using Kendall's coefficient

heatmaply_cor(cor(mtcars[,],method='kendall'))



Conclusions:

The variables are less correlated in the same areas as with Pearson's method.

```
prcomp(mtcars[,-1])->pca.mtcars
summary(pca.mtcars)
## Importance of components:
                               PC1
## Standard deviation 136.4339 38.14648 1.31542 0.96553 0.76789 0.32620 0.2892
## Proportion of Variance 0.9273 0.07249 0.00009 0.00005 0.00003 0.00001 0.0000
   Cumulative Proportion 0.9273 0.99982 0.99991 0.99995 0.99998 0.99999 1.0000
##
                            PC8 PC9
                                        PC10
## Standard deviation
                          0.2508 0.222 0.1999
## Proportion of Variance 0.0000 0.000 0.0000
## Cumulative Proportion 1.0000 1.000 1.0000
pca.mtcars$rotation[,1]
            cyl
##
    0.012042615 \quad 0.900235270 \quad 0.435074057 \quad -0.002661394 \quad 0.006242550 \quad -0.006676533
##
            VS
                          am
                                     gear
                                                   carb
   -0.002731293 -0.001963245 -0.002606103 0.005767541
```

```
df.pca=data.frame(pcl=pca.mtcars$x[,1],pc2=pca.mtcars$x[,2],pc3=pca.mtcars$x[,3],kl=as.factor(mtcars$cyl))
plot_ly(df.pca,x=~pc1,y=~pc2,z=~pc3,color = ~kl,type='scatter3d')
```

```
## No scatter3d mode specifed:
## Setting the mode to markers
## Read more about this attribute -> https://plotly.com/r/reference/#scatter-mode
```



WebGL is not supported by your browser - visit https://get.webgl.org for more info

Conclusions:

The first principal component explains 92.73% of the total variance. The variable am has the strongest influence on PC1. Clear cluster separation is visible along the PC1 axis

Unsupervised classification using k-means clustering

```
kmeans(mtcars[,-1], centers=3)->km.mtcars.3
table(km.mtcars.3$cluster,mtcars$cyl)
```

```
##
## 4 6 8
## 1 11 5 0
## 2 0 2 5
## 3 0 0 9
```

 $df.km<-data.frame(x=pca.mtcars$x[,1],y=pca.mtcars$x[,2],z=pca.mtcars$x[,3],type=as.factor(km.mtcars.3$cluster)) \\ plot_ly(df.km,x=~x,y=~y, z=~z,color=~type,type ='scatter3d')$

```
## No scatter3d mode specifed:
## Setting the mode to markers
## Read more about this attribute -> https://plotly.com/r/reference/#scatter-mode
```

```
1
2
3
```

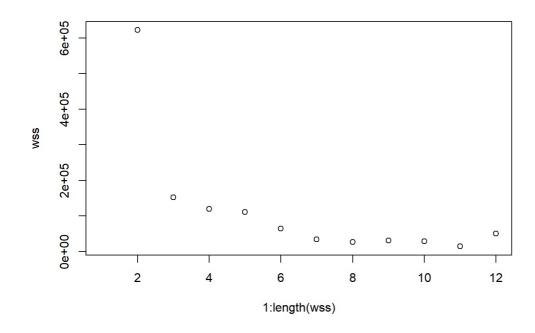
WebGL is not supported by your browser - visit https://get.webgl.org for more info

Conclusions:

A clear division into 3 clusters is visible, with one of the clusters being the most dispersed.

Determining the optimal number of clusters

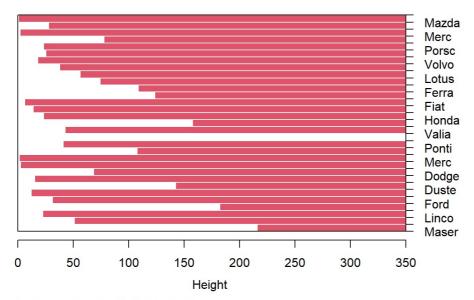
```
wss<-NA
for(i in 1:11) wss<-c(wss,kmeans(mtcars[,-1], centers=i)$tot.withinss)
plot(1:length(wss),wss)</pre>
```



Conclusion:

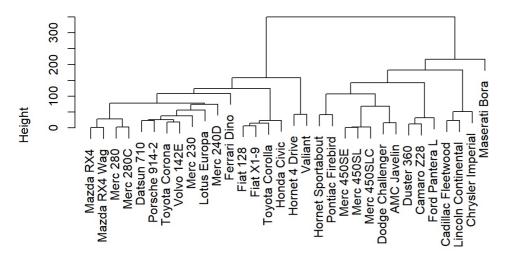
Using the elbow method, it can be concluded that 3 or 4 clusters are most optimal.

Banner of agnes(x = dist(mtcars[,], method = "minkowski", p = 1 diss = T)



Agglomerative Coefficient = 0.91

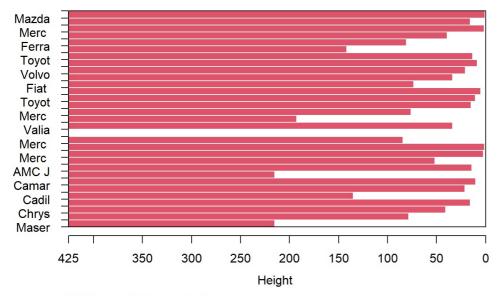
Dendrogram of agnes(x = dist(mtcars[,], method = "minkowski", p = 1] diss = T)



dist(mtcars[,], method = "minkowski", p = 1) Agglomerative Coefficient = 0.91

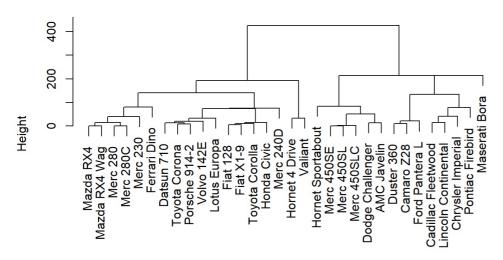
plot(diana(mtcars[,]))

Banner of diana(x = mtcars[,])



Divisive Coefficient = 0.93

Dendrogram of diana(x = mtcars[,])



mtcars[,]
Divisive Coefficient = 0.93

Conclusions:

Both methods yield similar results. The coefficients differ by only 0.02.

Supervised Classification

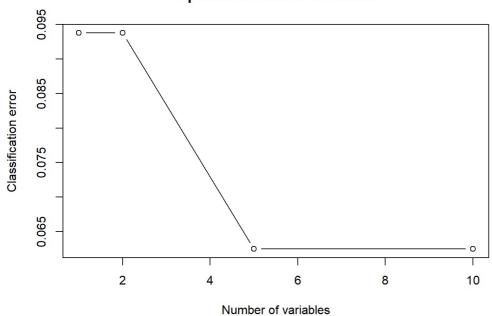
We use the random forest method.

```
rfcv(mtcars[,-2],as.factor(mtcars$cyl))->rf.ic
rf.ic$error.cv
```

```
## 10 5 2 1
## 0.06250 0.06250 0.09375 0.09375
```

```
plot(rf.ic$n.var, rf.ic$error.cv, type="b",
    xlab="Number of variables", ylab="Classification error",
    main="Optimal number of variables")
```

Optimal number of variables

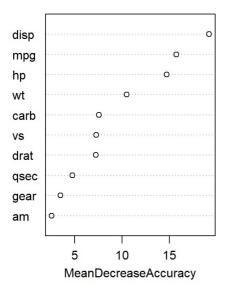


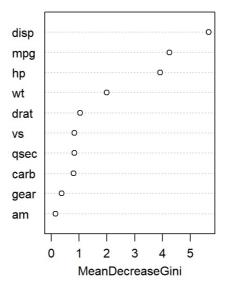
Conclusion:

The smallest cross-validation error occurs when the number of variables is between 2 and 5.

```
randomForest(mtcars[,-2],as.factor(mtcars$cyl),importance = T)->rf.mtcars.imp
varImpPlot(rf.mtcars.imp)
```

rf.mtcars.imp





Conclusions:

According to the accuracy drop criterion, the most important variables are disp, mpg, hp, and wt. This number of variables is acceptable based on previous results.

```
randomForest(mtcars[,c(3,1,4,6)],as.factor(mtcars$cyl))->rf.mtcars.sel
rf.mtcars.sel$confusion
```

Conclusion:

Using these variables, the maximum classification error is 28%.

Final Conclusions

The variables disp and hp have high mean and standard deviation values. The Grubbs test indicated outliers in qsec and carb, which should be considered in cluster analysis. Histogram analysis helped identify variables for normality and multimodality testing. Normality tests showed that hp and drat do not follow a normal distribution, while Silverman's test did not confirm multimodality for disp and wt. PCA analysis showed that the first principal component (PC1) explains 92.73% of the variance, mainly influenced by am. K-means clustering and the elbow method indicated 3 or 4 optimal clusters. Random forests are not an optimal classification model in this case, as the maximum classification error is 28%.