

## Correlation Analysis

The correlation coefficient  $r$  measures the strength and direction of a linear relationship,

- 1 indicates a perfect positive correlation.
- -1 indicates a perfect negative correlation.
- 0 indicates that there is no relationship between the different variables.

Values between -1 and 1 denote the strength of the correlation, as shown in the example below.

In this tutorial, we will explain the different ways of executing correlation plots in R

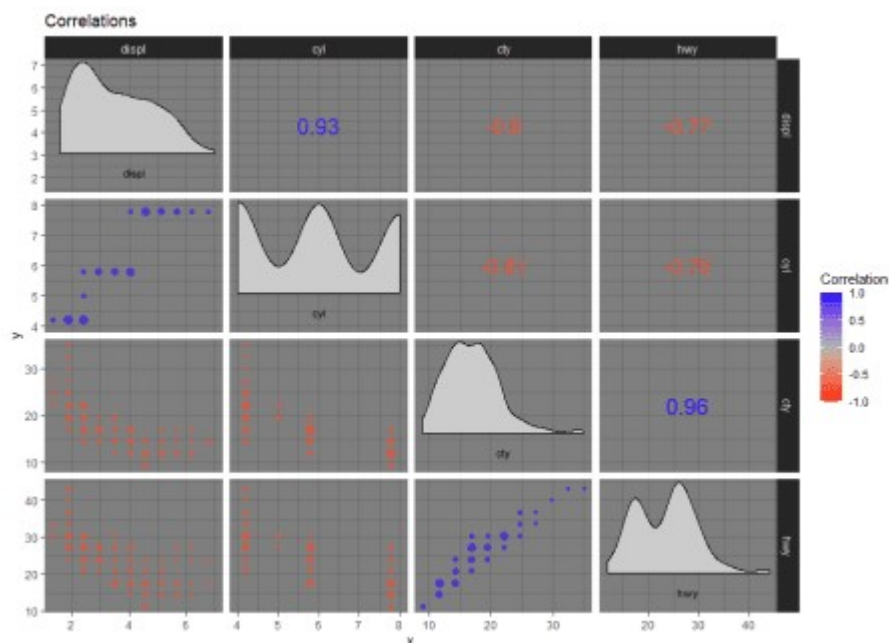
## Cormorant Package

```
# remotes::install_github("r-link/cormorant")
```

```
library(cormorant)
library(tidyverse)
library(dplyr)
```

We are selecting only quantitative variables for further analysis

```
mpg <- select(mpg, displ, cyl, cty, hwy)
cormorant(mpg, style = "binned") +
  theme_dark() +
  labs(title = "Correlations")
```



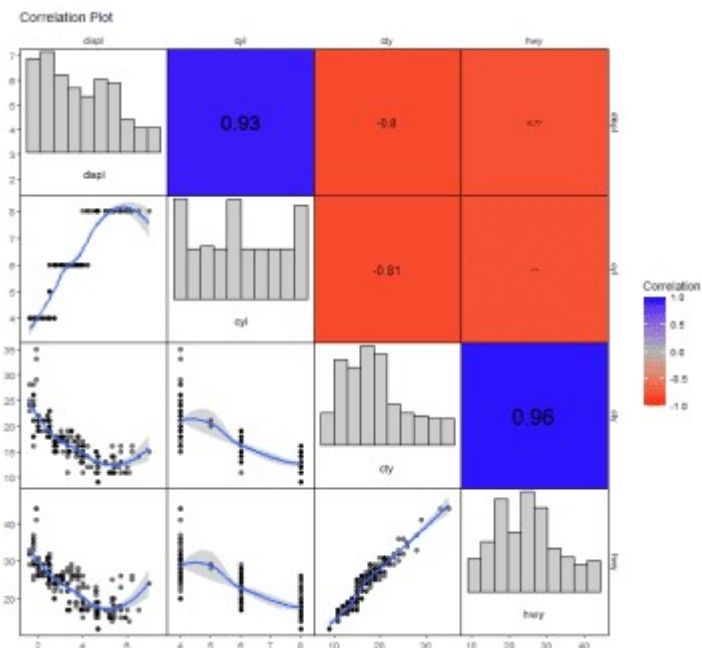
Customized Plot from ggcorrmm

```
ggcorrmm(data = mpg) +
  lotri(geom_point(alpha = 0.5)) +
  lotri(geom_smooth()) +
  utri_heatmap() +
  utri_corrtext() +
  dia_names(y_pos = 0.15, size = 3) +
```

```

dia_histogram(lower = 0.3, fill = "grey80", color = 1) +
scale_fill_corr() +
labs(title = "Correlation Plot")

```



### Visualize correlation matrix using corrplot

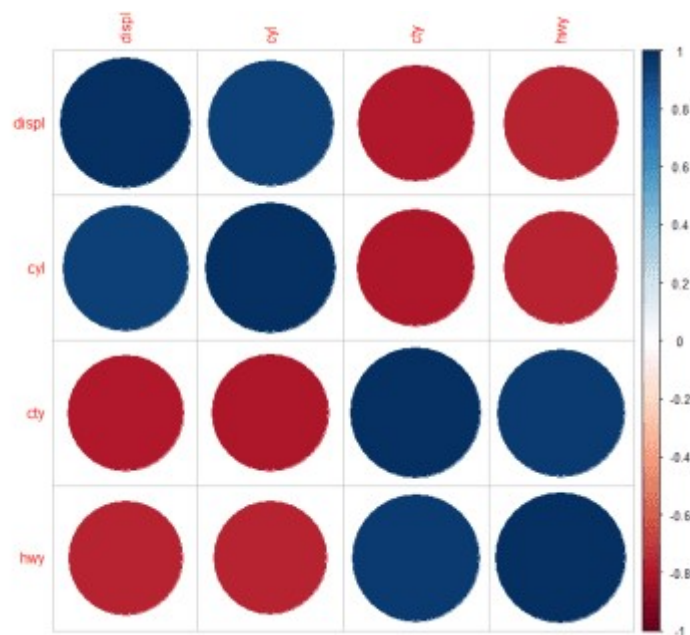
Following plots, correlation coefficients are colored according to the value. The correlation matrix can be also reordered according to the degree of association between variables.

#### How to learn statistics?

```

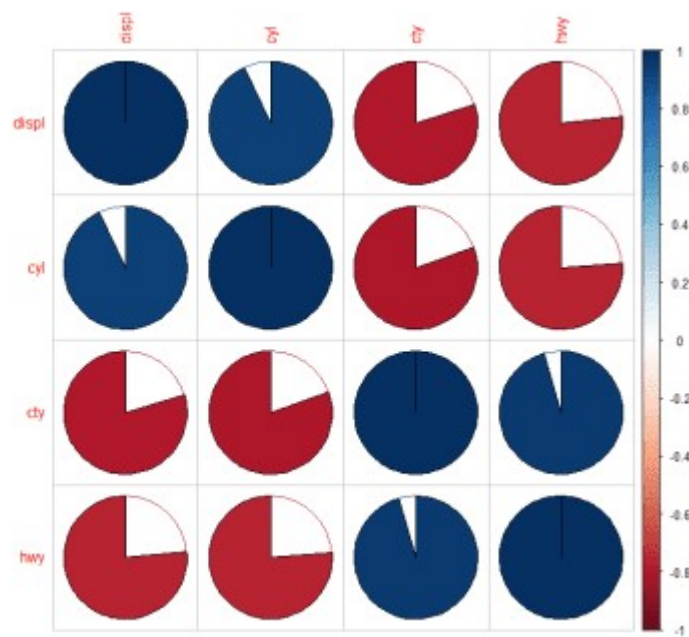
library(corrplot)
library(RColorBrewer)
M <- cor(mpg)
corrplot(M, method="circle")

```

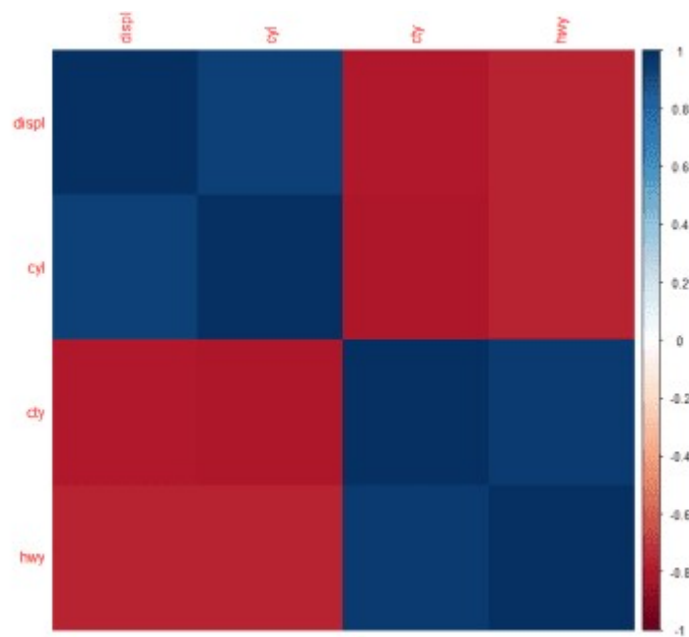


```
corrplot(M, method="pie")
```

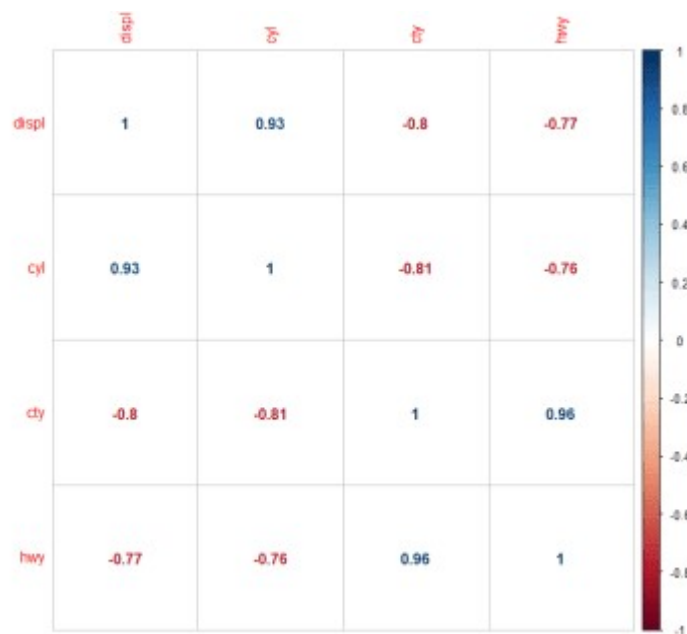
## Difference between association and correlation



```
corrplot(M, method="color")
```



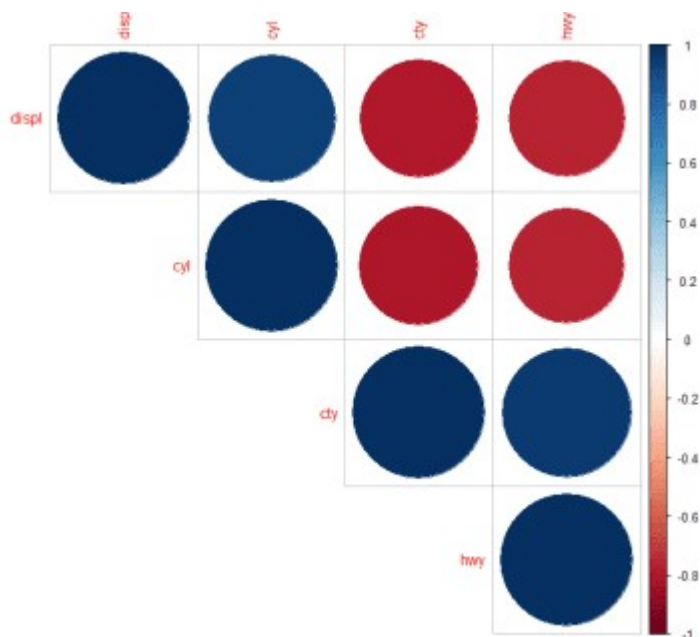
```
corrplot(M, method="number")
```



There are three types of layout :

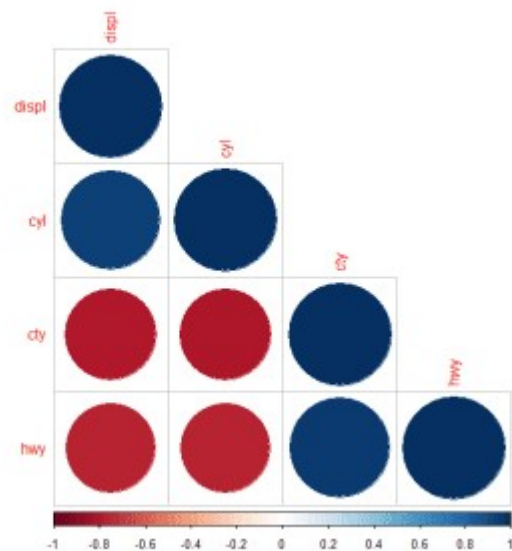
- “full” (default) : display full correlation matrix
- “upper”: display upper triangular of the correlation matrix
- “lower”: display lower triangular of the correlation matrix

```
corrplot(M, type="upper")
```

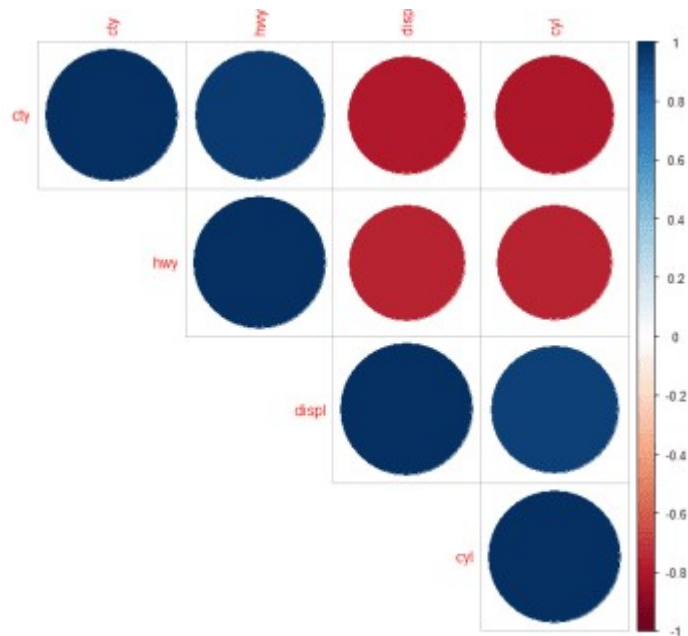


```
corrplot(M, type="lower")
```

Types of data visualization charts



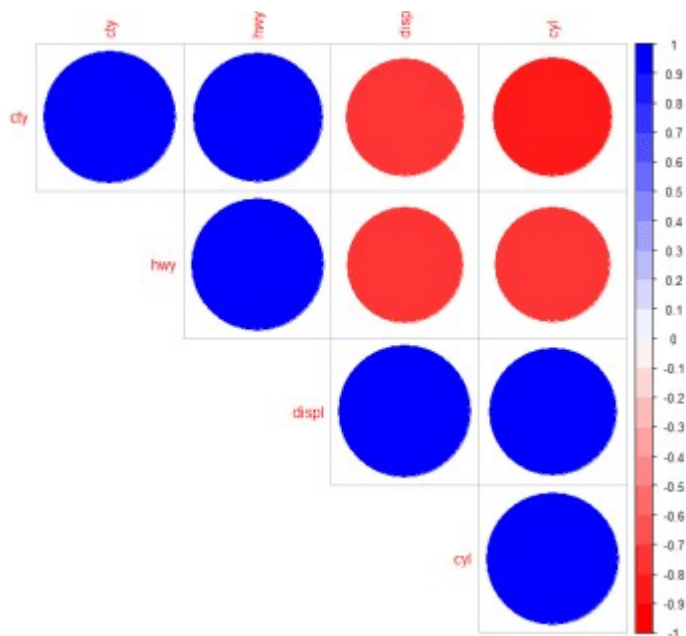
```
corrplot(M, type="upper", order="hclust")
```



## Using different color spectrum

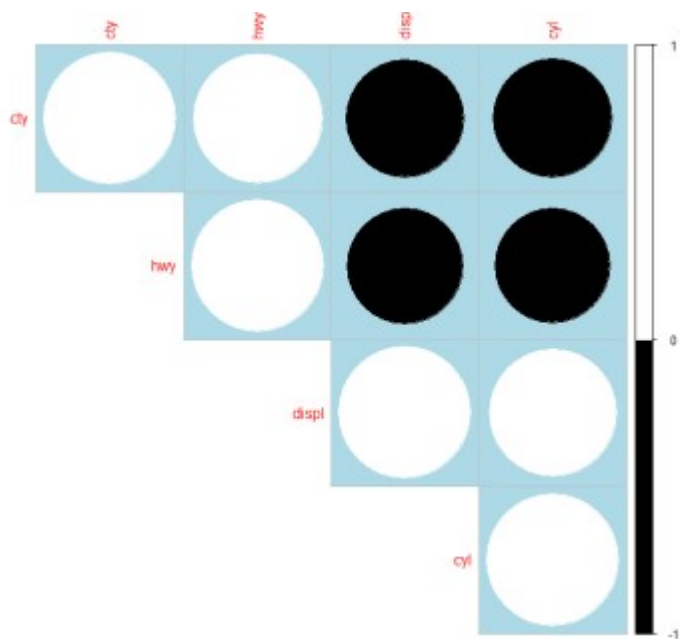
### [Sample size calculation in R](#)

```
col<- colorRampPalette(c("red", "white", "blue"))(20)
corrplot(M, type="upper", order="hclust", col=col)
```



Change background color to lightblue

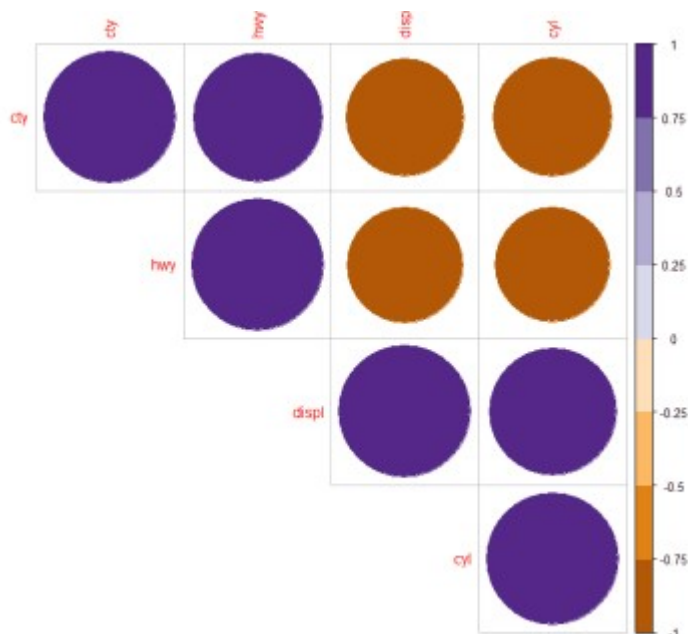
```
corrplot(M, type="upper", order="hclust", col=c("black", "white"),
         bg="lightblue")
```



Changing the color of the plot

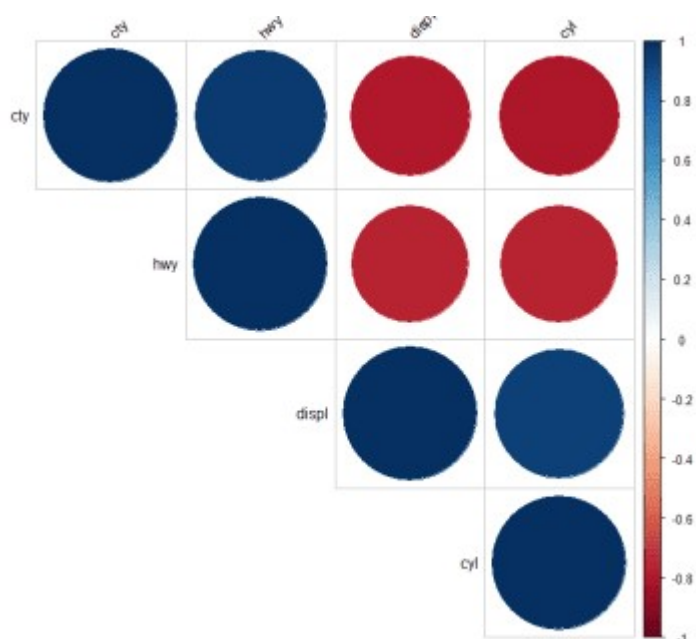
[Random Forest Feature selection in R](#)

```
corrplot(M, type="upper", order="hclust",
         col=brewer.pal(n=8, name="PuOr"))
```



Changing the color and the rotation of text labels

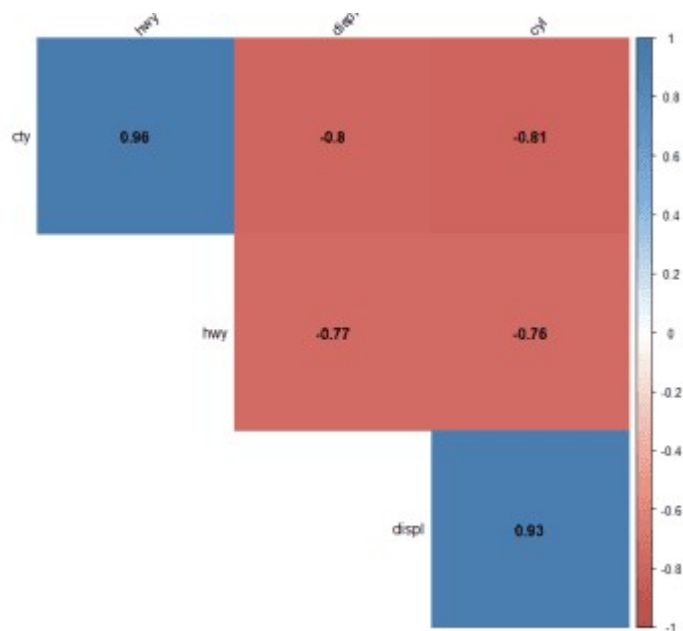
```
corrplot(M, type="upper", order="hclust", tl.col="black", tl.srt=45)
```



Customize the corrplot

### Handling Imbalanced data in R

```
col <- colorRampPalette(c("#BB4444", "#EE9988", "#FFFFFF", "#77AADD",
"#4477AA"))
Specialized the insignificant value according to the significant level
corrplot(M, type="upper", order="hclust",
          p.mat = p.mat, sig.level = 0.01)
corrplot(M, method="color", col=col(200),
          type="upper", order="hclust",
          addCoef.col = "black", # Add coefficient of correlation
          tl.col="black", tl.srt=45, #Text label color and rotation
```



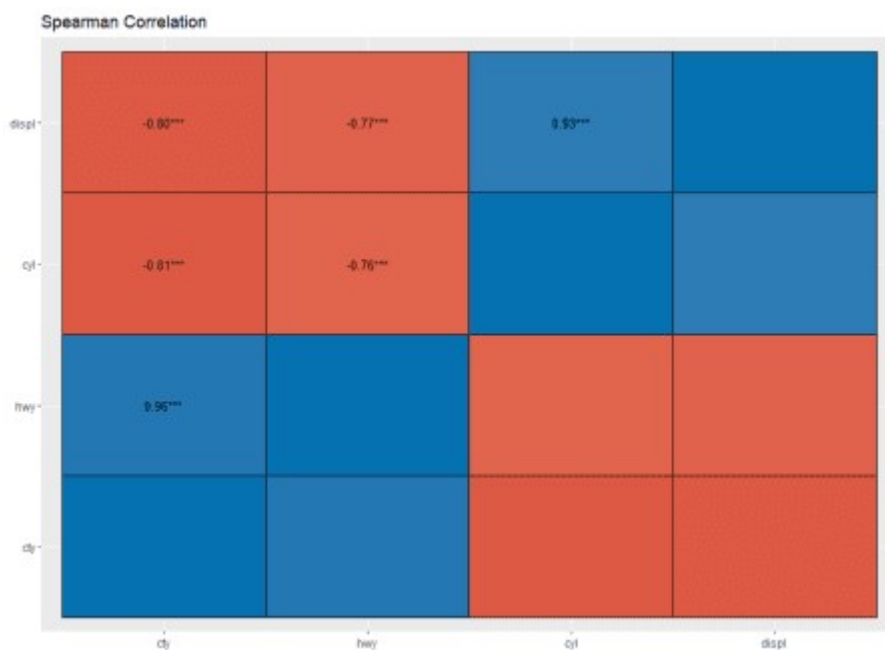
## sjPlot Package

```
sjp.corr( data, title = NULL, axis.labels = NULL, sort.corr = TRUE,
  decimals = 3, na.deletion = c("listwise", "pairwise"), corr.method =
  c("pearson", "spearman", "kendall"), geom.colors = "RdBu", wrap.title =
  50, wrap.labels = 20, sjp.corr 65 show.legend = FALSE, legend.title =
  NULL, show.values = TRUE, show.p = TRUE, p.numeric = FALSE )
```

sjplot is very useful for small number of variables.

[ggside in R](#)

```
library(sjPlot)
sjp.corr(mpg, title = "Spearman Correlation", decimals = 2)
```

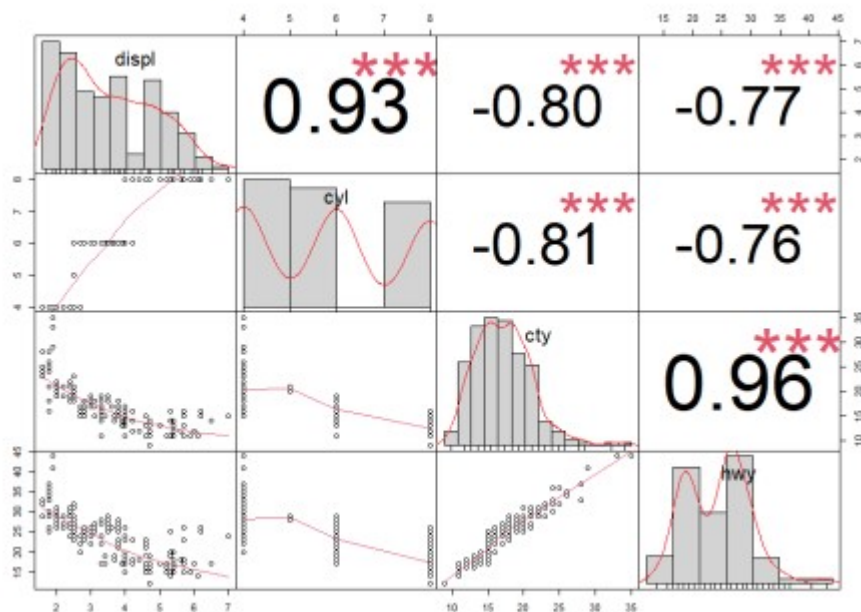


## PerformanceAnalytics Package

```
library(PerformanceAnalytics)
```



```
chart.Correlation(mpg, histogram=TRUE, pch="+")
```



Correlation plots are the best way to show the pattern and relationship.