## R Markdown Presentation & Plotly

## Instructions

These are the aspect that we will be covering with the presentation: This introduction in which we are dealing with rmarkdown 1. A second one in which run R code inside the slide. 2. And finally we will draw a chart with plotly.

```
library(plotly)
## Warning: package 'plotly' was built under R version 3.4.4
## Loading required package: ggplot2
## Warning: package 'ggplot2' was built under R version 3.4.4
## Attaching package: 'plotly'
## The following object is masked from 'package:ggplot2':
##
##
       last_plot
## The following object is masked from 'package:stats':
##
##
       filter
## The following object is masked from 'package:graphics':
##
##
       layout
```

## Loading required package: ggplot2

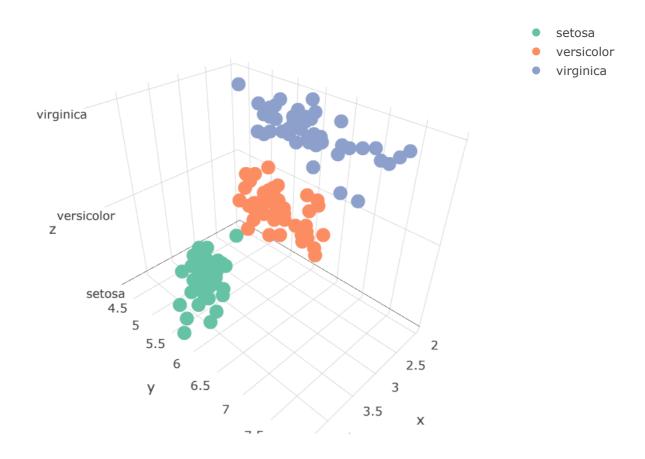
Attaching package: 'plotly' The following object is masked from 'package:ggplot2': last\_plot The following object is masked from 'package:graphics': layout

```
data(iris)
head(iris)
```

```
##
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1
              5.1
                          3.5
                                       1.4
                                                    0.2 setosa
## 2
              4.9
                          3.0
                                       1.4
                                                    0.2 setosa
              4.7
## 3
                          3.2
                                        1.3
                                                    0.2 setosa
              4.6
## 4
                          3.1
                                        1.5
                                                    0.2 setosa
              5.0
                          3.6
## 5
                                        1.4
                                                    0.2 setosa
## 6
              5.4
                          3.9
                                       1.7
                                                    0.4 setosa
```

```
plot_ly(x= iris$Sepal.Width, y= iris$Sepal.Length, z = iris$Species,
type = "scatter3d", mode = "markers", color = iris$Species)
```

```
## Warning: package 'bindrcpp' was built under R version 3.4.4
```



Warning in arrange\_impl(.data, dots): '.Random.seed' is not an integer vector but of type 'NULL', so ignored ==========