

Cluster Analysis

Cesar Conejo Villalobos

1/29/2021

R Markdown

```
getwd()
```

```
## [1] "C:/Users/CCONE/Documents/Cesar/Projects/RControlVersion/UC3M/MultiAnalysis"
```

```
#load('rda/clinical_trial_complete.rda')
```

```
#summary(clinical_trial_complete)
```

we back one

```
#dir.create(path = '../Report3-Cluster_Factor_Analysis')
```

```
dir()
```

```
## [1] "data"                "figure_output"
## [3] "rda"                 "README.md"
## [5] "Report1-Data_Presentation" "Report2-PCA"
## [7] "Report3-Cluster_Factor_Analysis" "scripts"
## [9] "UC3M_MultiAnalysis_Project_1.Rproj"
```

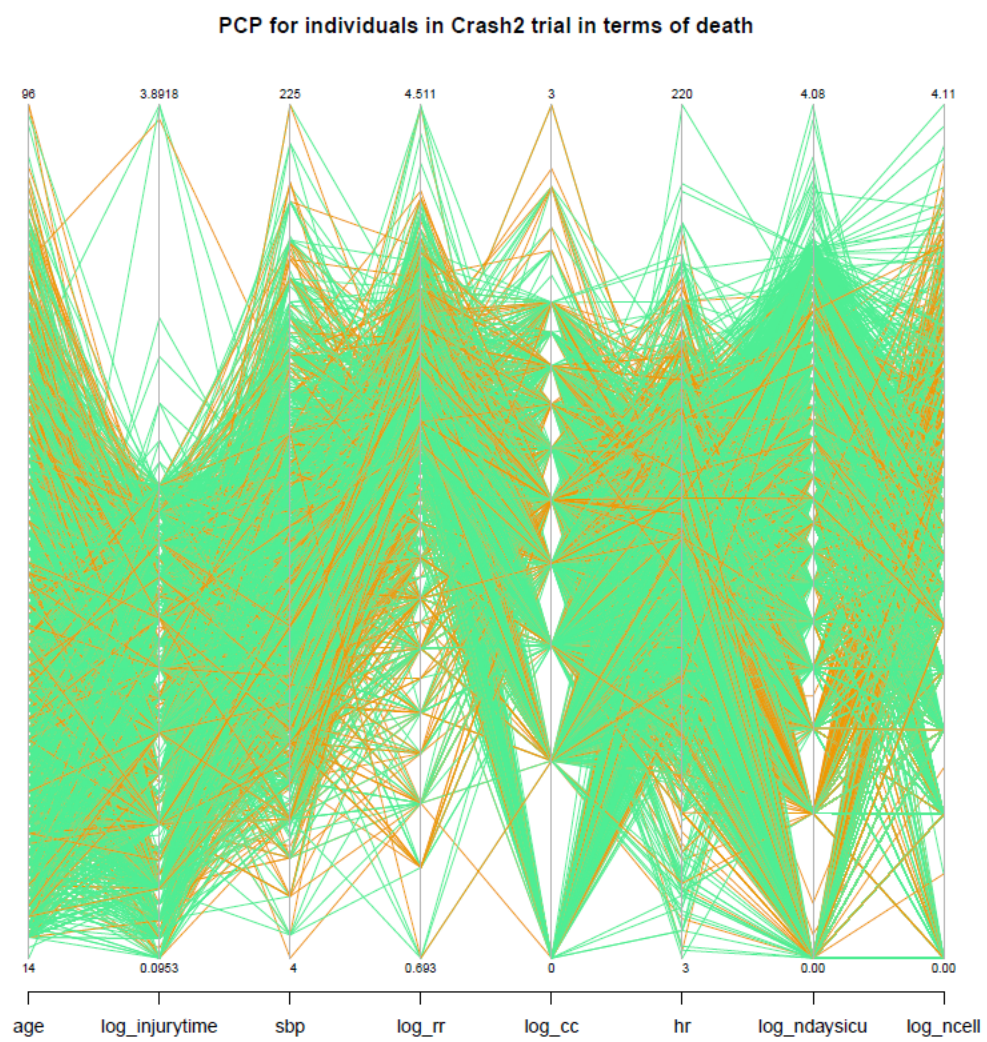


Figure 1: PCP plot of all quantitative variables