Step1 Team Project Multivariate Analysis

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Introduction data set

We have selected the CRASH-2 data set provided by Vanderbilt School of Biostatistics for our project. It describes the outcome of a randomized controlled trial and economic valuation of the effects of tranexamic acid on death, vascular occlusive events, and transfusion requirement in bleeding trauma patients. Tranexamic acid reduces bleeding in trauma patients undergoing surgery but is an expensive treatment option. The trial's objective was to assess the effects and cost-effectiveness of an early administration of this medication.

Participants of the study were adults with, or at risk of, significant bleeding within 8 hours of injury. Sample randomization was determined by the allocation of an eight-digit sequence randomly generated by a computer. Patients and staff were masked to the treatment allocation of the tranexamic acid.

We have adjusted the original data set to remove some variables that were not relevant to our investigation. We have removed variables regarding the exact surgical procedures administered to patients, various IDs, and details on the patient outcome. We removed the health outcome columns because of complications regarding missing data, where the boolean structure of the columns relating to specific outcomes, like stroke or pulmonary embolism, left a large number of cases with missing values. Instead, we added a boolean variable for a general outcome of survival to assess the efficacy of the procedure, rather than looking at particular health outcomes in post-surgery for living patients.

We will be using variables regarding the sex, age, and injury of the patient as well as certain biometrics, like blood pressure, respiratory and heart rates, details on surgical blood transfusion, and a boolean variable on the survival of the patient. Our selection provides us with a balance of continuous and categorical variables, many of which are boolean, with minimal complications due to missing data.

Summary variables in the data set

The variables in this dataset are the following:

1. sex: (Boolean) The sex of the patient (Male/Female)
2. age : (Numerical) Age of the patient(Years)
3. injurytime: (Numerical) Hours since injury (Hours)
4. injurytype: (Categorical) Type of injury {Blunt, Penetrating, Blunt and Penetrating}
5. sbp: (Numerical) Systolic Blood Pressure (mmHg)
6. rr: (Numerical) Respiratory Rate (rate per minute)
7. cc: (Numerical) Central Capillary Refille Time (seconds)
8. hr: (Numerical) Heart Rate (rate per minute)
9. ndaysicu: (Numerical) Number of days in ICU (days)
10. ncell: (Numerical) Number of Units of Red Call Products Transfused.

11. Death: (Boolean) Indicator if the patient survived after the procedure

A summary of the data type is the following:

variable	type_variable	sub_type_variable
sex	Qualitative	Nominal
age	Quantitative	Continuous
injurytime	Quantitative	Continuous
injurytype	Qualitative	Nominal
sbp	Quantitative	Continuous
rr	Quantitative	Continuous
cc	Quantitative	Continuous
hr	Quantitative	Continuous
ndaysicu	Quantitative	Discrete
ncell	Quantitative	Continuous
death	Qualitative	Nominal

A review of the structure of the dataset is the following:

```
## 'data.frame':    9497 obs. of  11 variables:
##  $ sex       : Factor w/ 2 levels "male","female": 1 1 1 1 1 1 1 1 1 2 ...
##  $ age       : int  50 30 40 19 27 16 29 41 56 37 ...
##  $ injurytime: num  1 1 2 3 0.5 1 1 0.5 0.5 8 ...
##  $ injurytype: Factor w/ 3 levels "blunt","penetrating",..: 1 1 2 2 2 2 1 2 1 2 ...
##  $ sbp       : int  75 70 60 90 90 90 116 120 60 104 ...
##  $ rr        : int  28 26 20 30 26 28 15 15 9 23 ...
##  $ cc        : int  5 6 5 5 5 2 3 3 3 5 ...
##  $ hr        : int  120 130 120 90 96 118 118 70 100 92 ...
##  $ ndaysicu  : num  0 6 2 9 7 0 7 7 23 2 ...
##  $ ncell     : num  1 2 4 2 1 1 16 8 4 4 ...
##  $ death     : Factor w/ 2 levels "0","1": 2 1 2 2 1 1 1 1 1 1 ...
```

A summary of the values in the data set are:

```
##      sex            age          injurytime                    injurytype
##  male  :7906   Min.   :14.00   Min.   : 0.100   blunt                :5211
##  female:1591   1st Qu.:24.00   1st Qu.: 1.000   penetrating          :2937
##                Median :31.00   Median : 3.000   blunt and penetrating:1349
##                Mean   :34.66   Mean   : 3.094
##                3rd Qu.:43.00   3rd Qu.: 4.500
##                Max.   :96.00   Max.   :48.000
##       sbp              rr              cc               hr
##  Min.   :  4.00   Min.   : 2.00   Min.   : 1.000   Min.   :  3.0
##  1st Qu.: 80.00   1st Qu.:20.00   1st Qu.: 2.000   1st Qu.: 96.0
##  Median : 90.00   Median :22.00   Median : 3.000   Median :110.0
##  Mean   : 93.13   Mean   :23.46   Mean   : 3.438   Mean   :108.1
##  3rd Qu.:104.00   3rd Qu.:28.00   3rd Qu.: 4.000   3rd Qu.:120.0
##  Max.   :225.00   Max.   :91.00   Max.   :20.000   Max.   :220.0
##     ndaysicu          ncell         death
##  Min.   : 0.000   Min.   : 0.000   0:7672
##  1st Qu.: 0.000   1st Qu.: 2.000   1:1825
##  Median : 1.000   Median : 3.000
##  Mean   : 4.137   Mean   : 3.912
##  3rd Qu.: 5.000   3rd Qu.: 5.000
##  Max.   :58.000   Max.   :60.000
```

Finally, the list of different values by column is the following:

Table 2: Count of distinct values of each variable

sex	age	injurytime	injurytype	sbp	rr	cc	hr	ndaysicu	ncell	death
2	81	78	3	153	58	16	154	47	47	2

Visual Analysis

First, we will review the distribution of the variables involve in the dataset

In the case of age, the figure 1 reflects how this variable appears to be largely weighted to the left, with lower ages featuring more frequently than those that are greater, possibly reflecting that younger people often take more risk and work higher at-risk occupations, raising their chance of experiencing trauma involving bleeding.

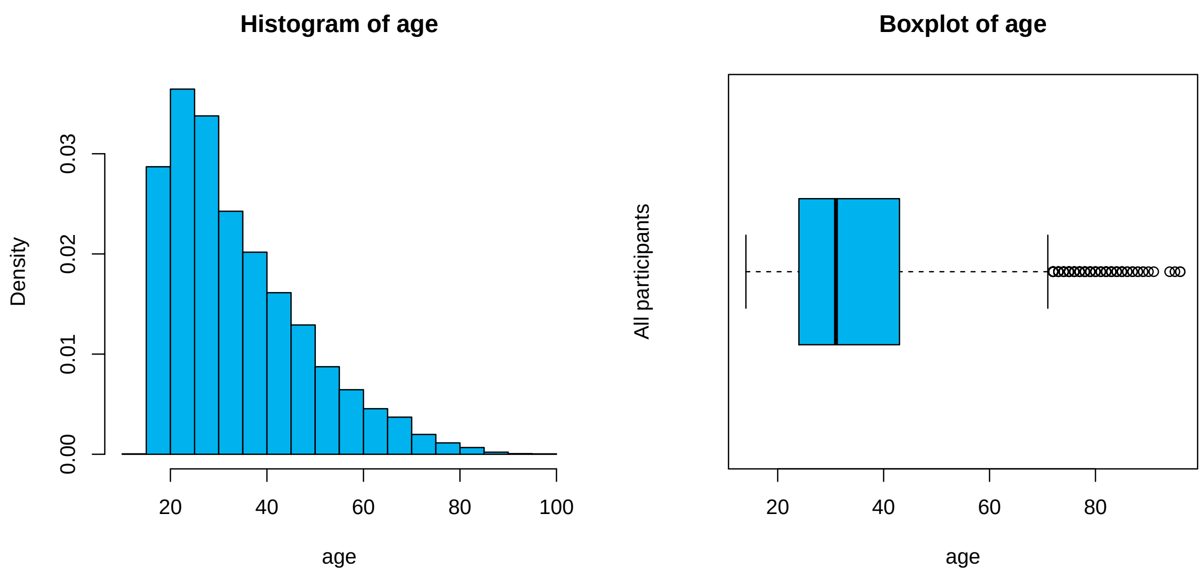

Figure 1: Distribution of age variable

The figure 2 show the distribution of the variable Injury time. We can see how this variable is highly positive skewed with almost all values falling below ten minutes since the injury was experienced. This is likely due to the fact that in cases of serious injury victims are brought to the hospital quite quickly. Then, we apply a *log* transformation `log_injurytime = log(injurytime + 1)` to this variables as we can see in figure 3.

For *sbp* (Systolic Blood Pressure), the distribution is a fairly centrally balanced distribution around 90 mmHg. This is logical as a sample of biological characteristics observed in a population is likely to have most people around the mean and then a reasonably tight distribution of those who differ, similar to that of other biological features like height. Furthermore, most people are fairly young in the sample and therefore would have rates that deviant less from the norm, at a healthy level. The distribution is given by figure 4.

In the case of *rr* (Respiratory Rate) appears, similar to sbp, resembling a moderately balanced distribution around 22 respirations per minute, although is weighted more to the right. The distribution of this variable is showed in figure 5. Taking a *log* transformation `log_rr = log(rr)`, we have the new distribution in figure 6.

In the case of *hr* (Heart rate), figure 7 show that this distribution seems fairly balanced at around 110, similar to the variables above, like *sbp* and *rr*.

For *cc* (Central capillary) refill has 75% of the observations below of 4 as we can appreciate in figure 8. However, the distribution is right-skewed. As a result, we apply a *log* transformation `log_cc = log(cc)` that is given in figure 9.

The figure 10 shows the distribution of the *ndaysicu:* variable. In this case, the distribution is heavily weighted to the left and right-skewed. Most patients it seems, with injuries at high risk of bleeding, do not often need to remain in the hospital for long. The transformed distribution `log_ndaysicu = log(ndaysicu + 1)` is given in the figure 11.

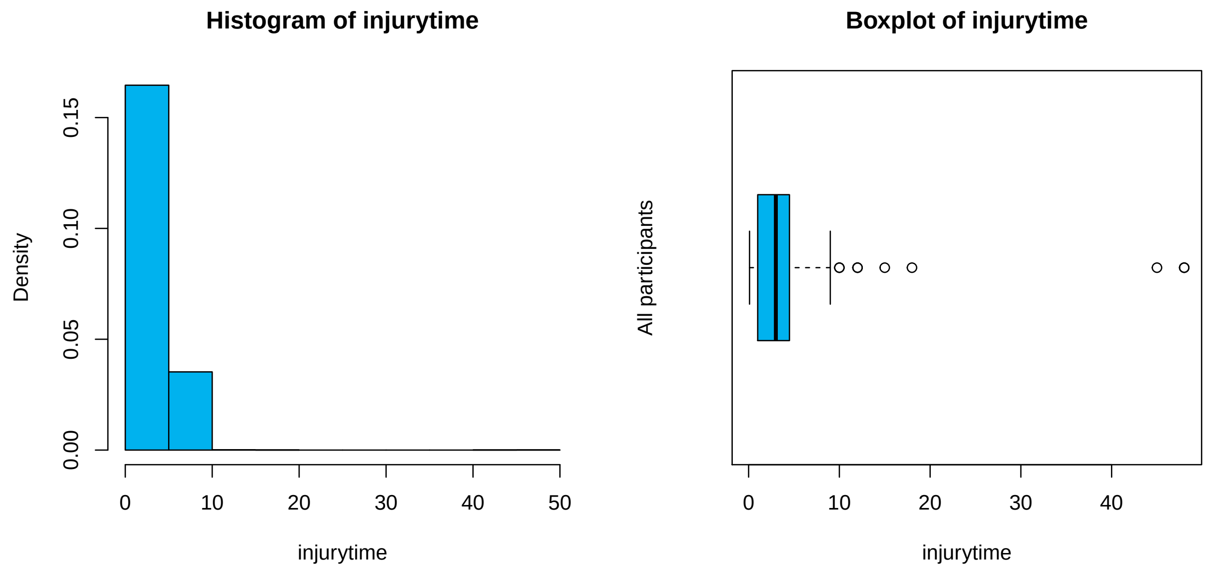

Figure 2: Distribution of injurytime variable

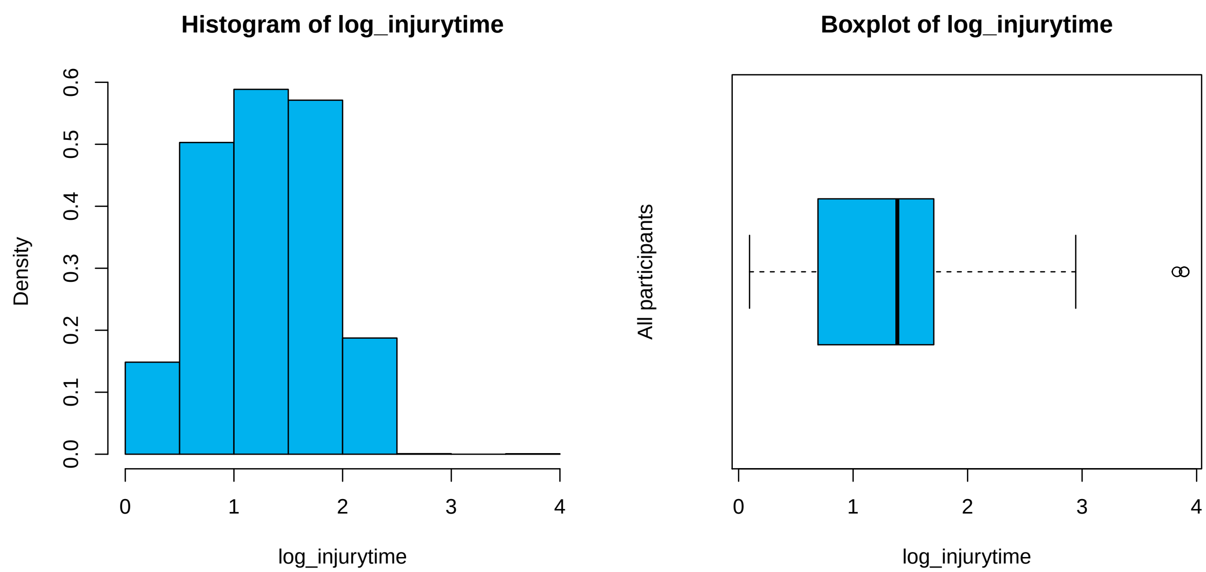

Figure 3: Distribution of the log injurytime variable

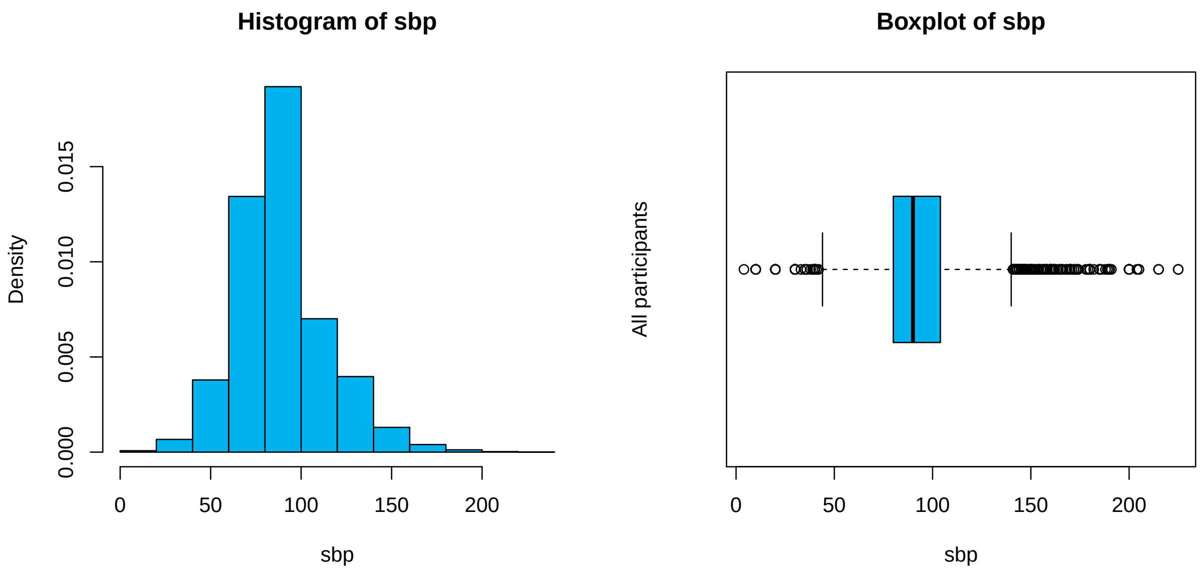

Figure 4: Distribution of sbp (Systolic Blood Pressure)

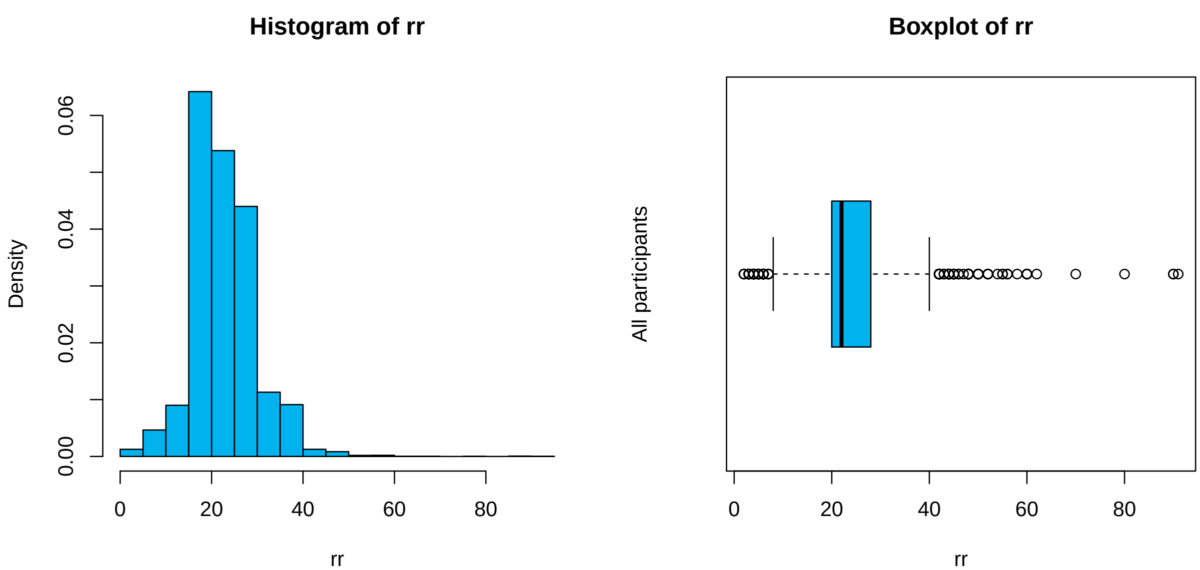

Figure 5: Distribution of rr (Respiratory Rate)

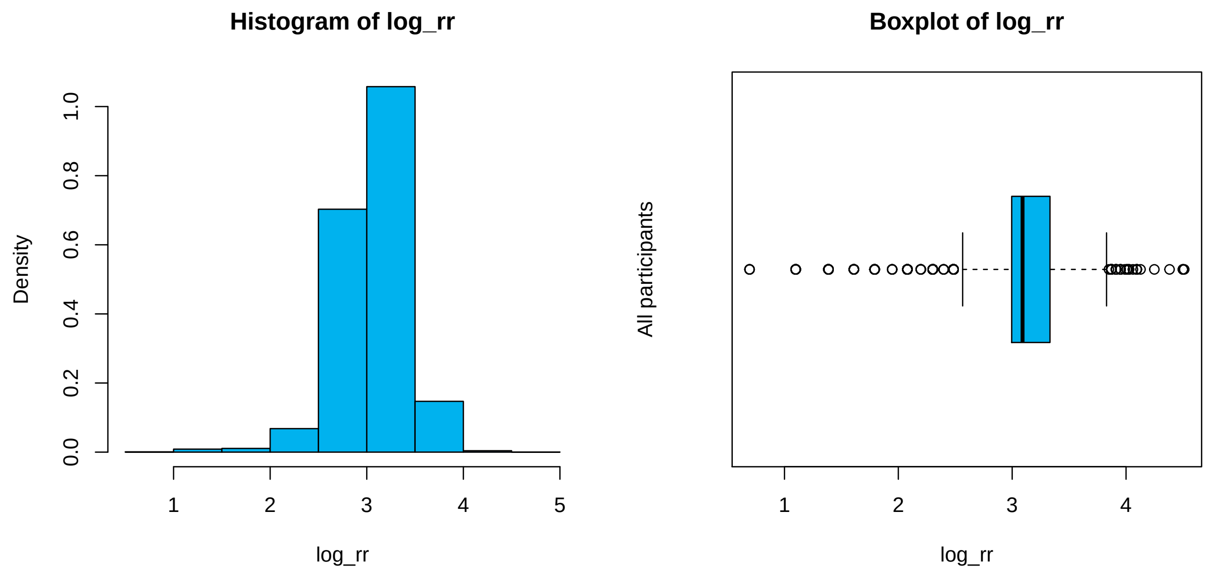

Figure 6: Distribution of log transformation of rr (Respiratory Rate)

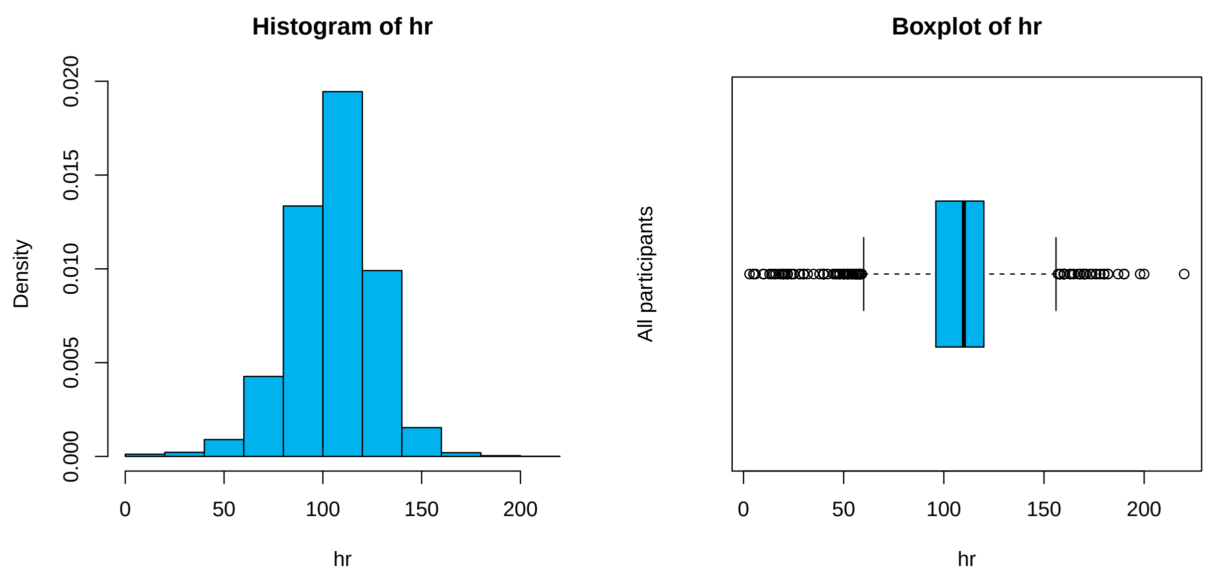

Figure 7: Distribution of hh (hearth Rate)

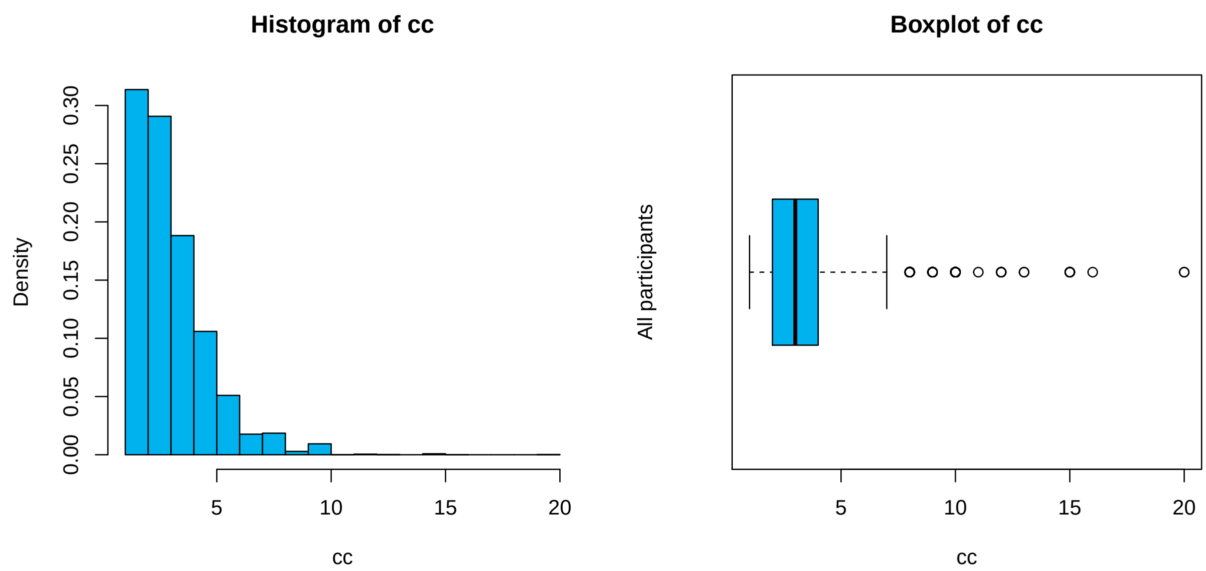

Figure 8: Distribution of cc (Central capillary)

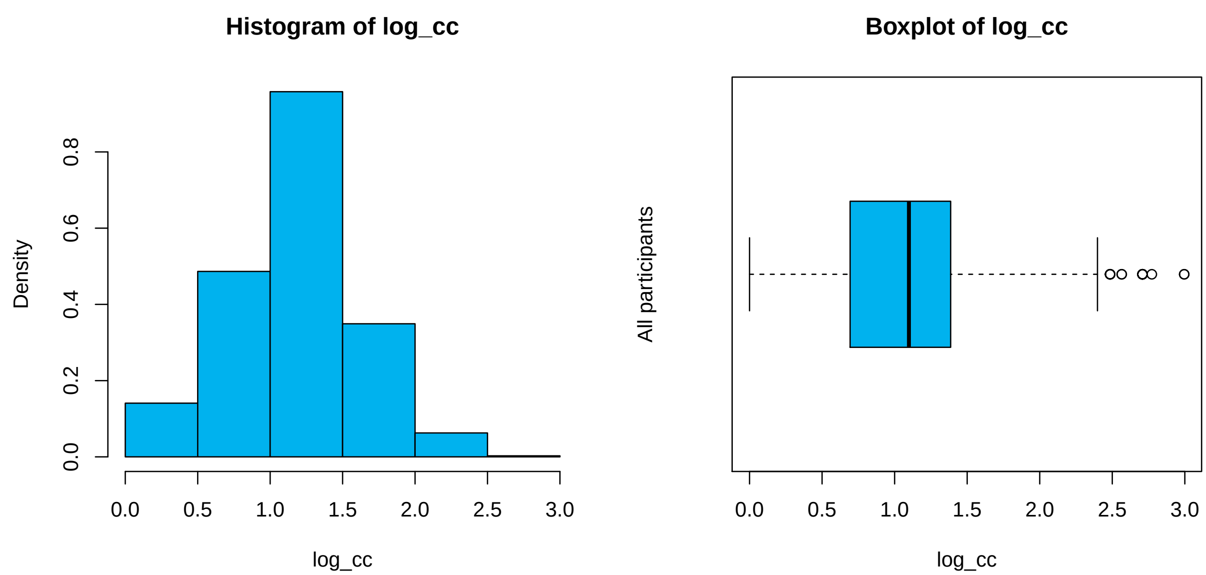

Figure 9: Distribution of log transformation of cc (Central capillary)

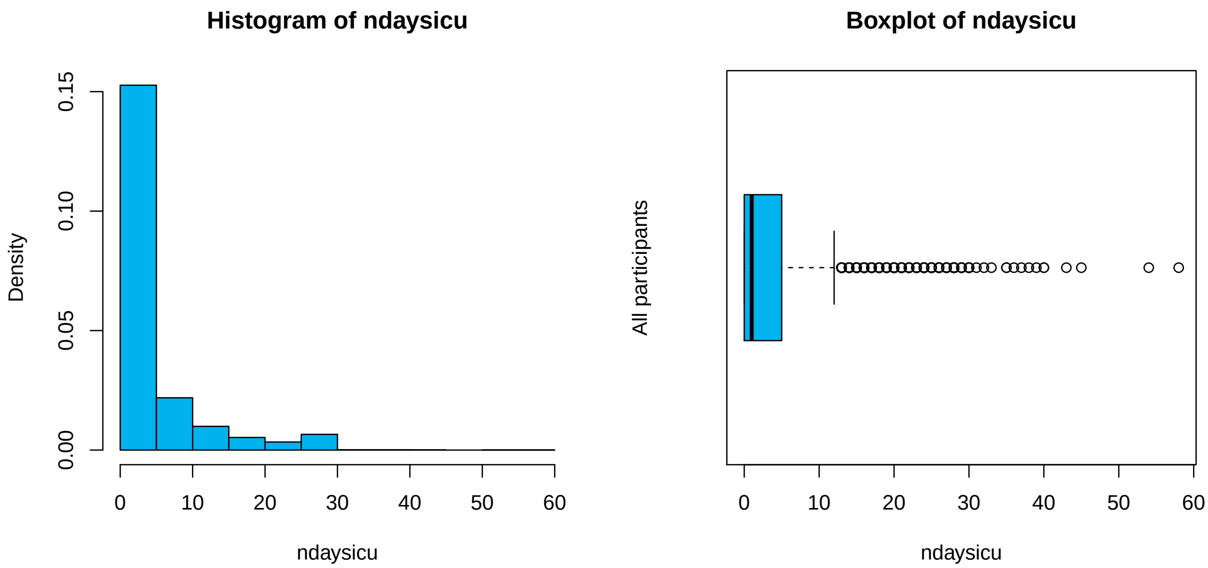

Figure 10: Distribution of ndaysicu

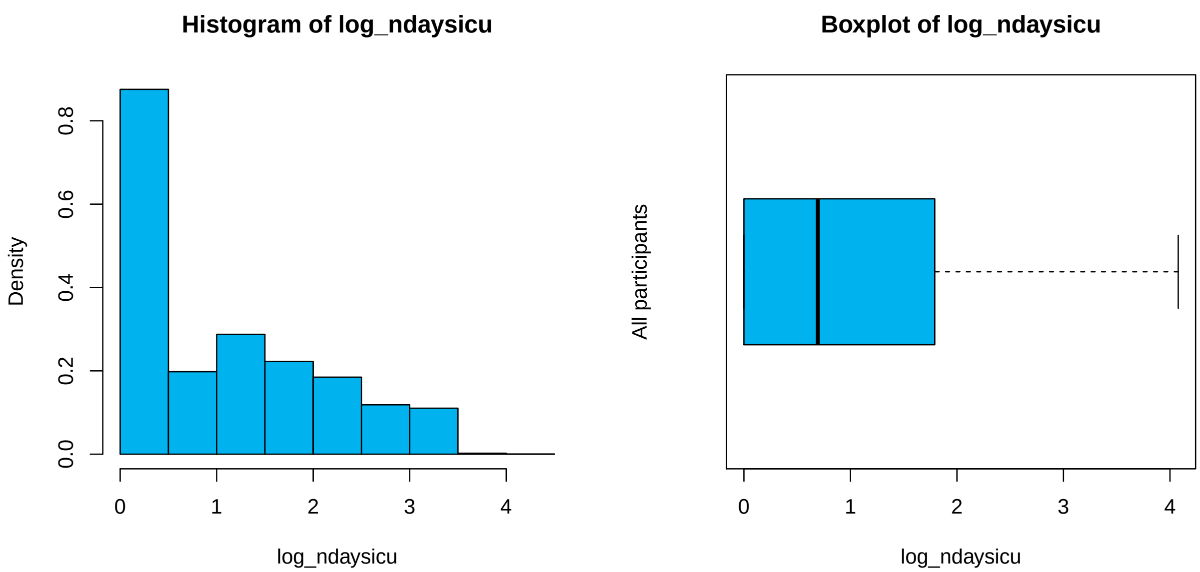

Figure 11: Distribution of log ndaysicu

Finally, for the *ncell*distribution is weighted to the left with a median of 3 as we can appreciate in the figure 12. The conclusion of this it that with many patients, only needing a small number of or zero units of red cell products transfused. Due to this variable is highly right skewed, we apply the *log* transformation `log_ncell = log(ncell + 1)` of the figure 13.

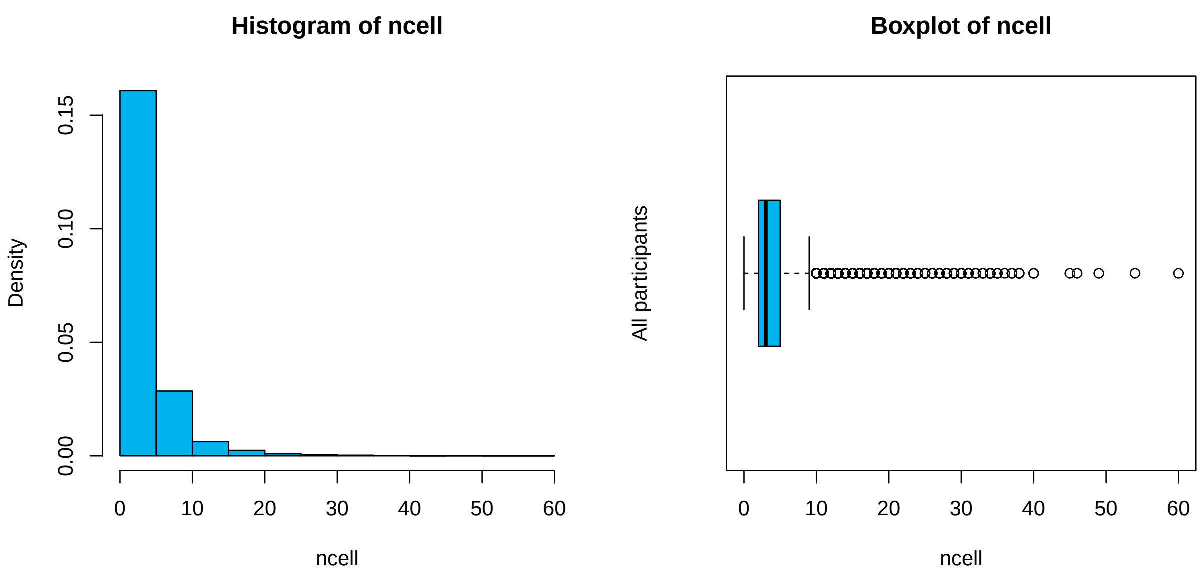

Figure 12: Distribution of ncell

For the categorical variables, we will focus on the distributions of deaths. The figure 14, shows that approximately for each death, 4 people survive. In the context of this problem, if we have an unbalanced proportion of people that survive can be considered as a sign that the drugs works.

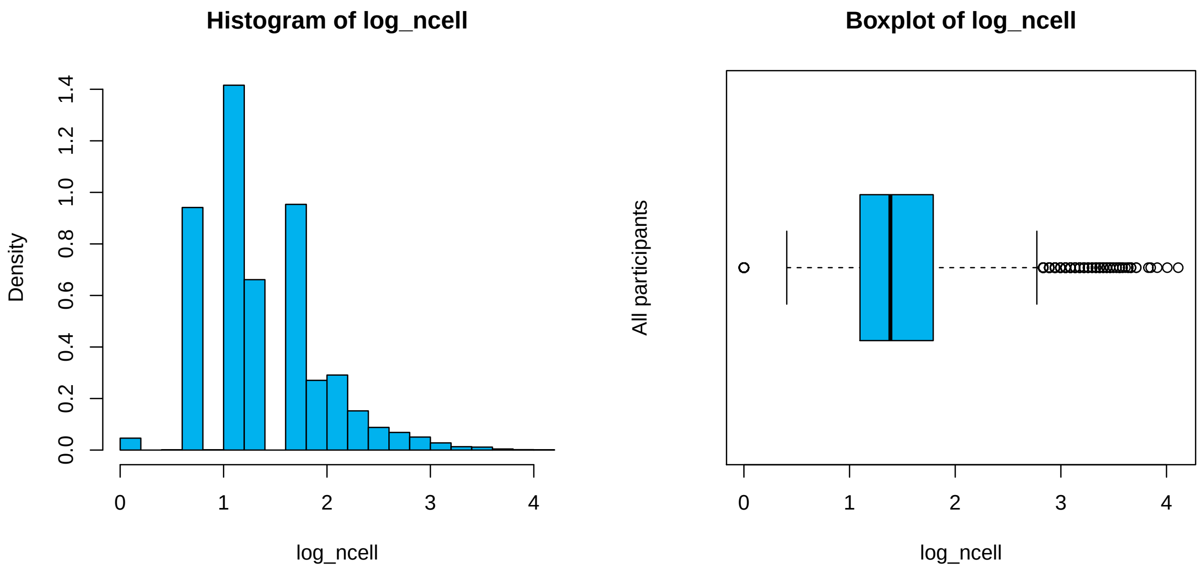

Figure 13: Distribution of log ncell

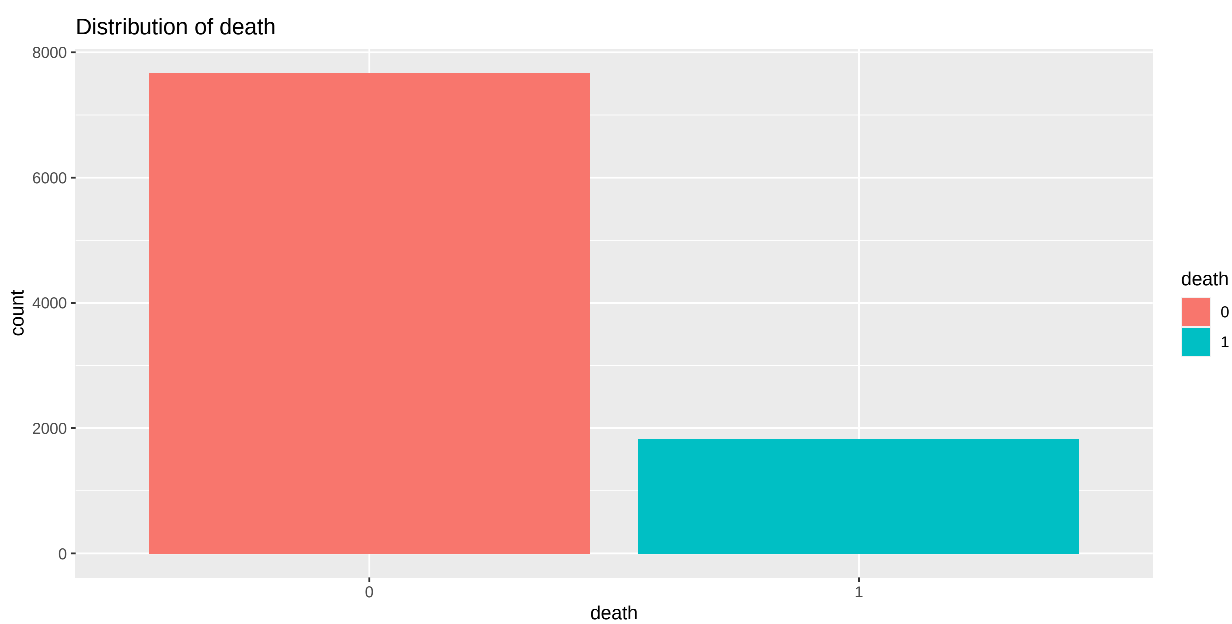

Figure 14: Distribution of deaths

On the other hand, we can study some relations of the quantitative variables in terms of the categorical variable death. Figure 15 . . .

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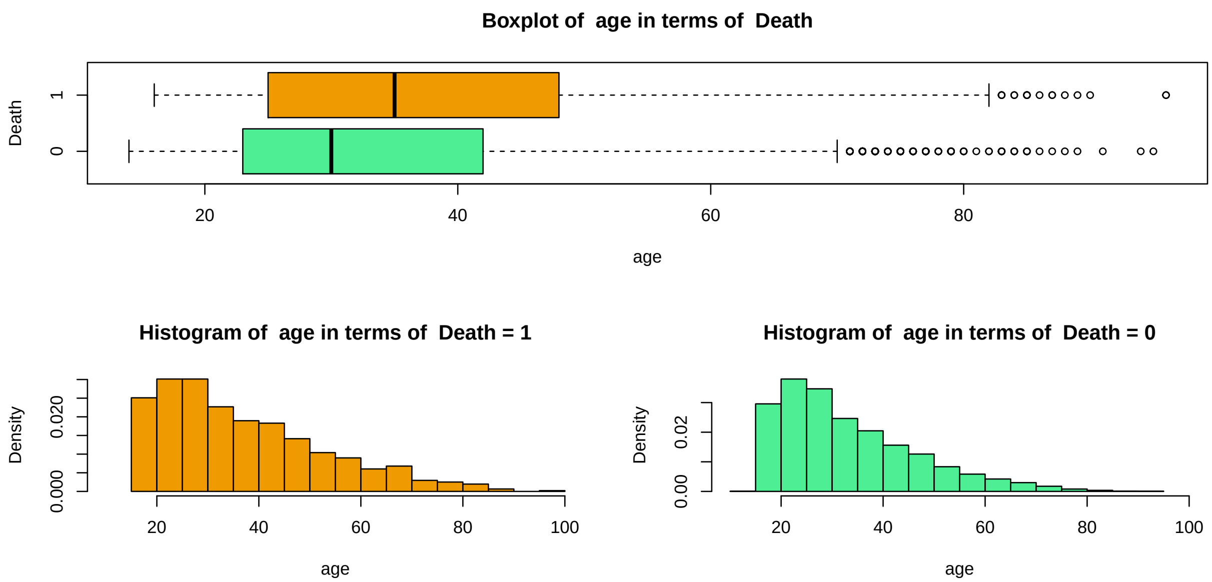

Figure 15: Distribution of age in terms of death

Figure 16 . . . .

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Figure 17 . . . .

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Figure 16 . . .

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Figure 19 . . .

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Figure 20 . . .

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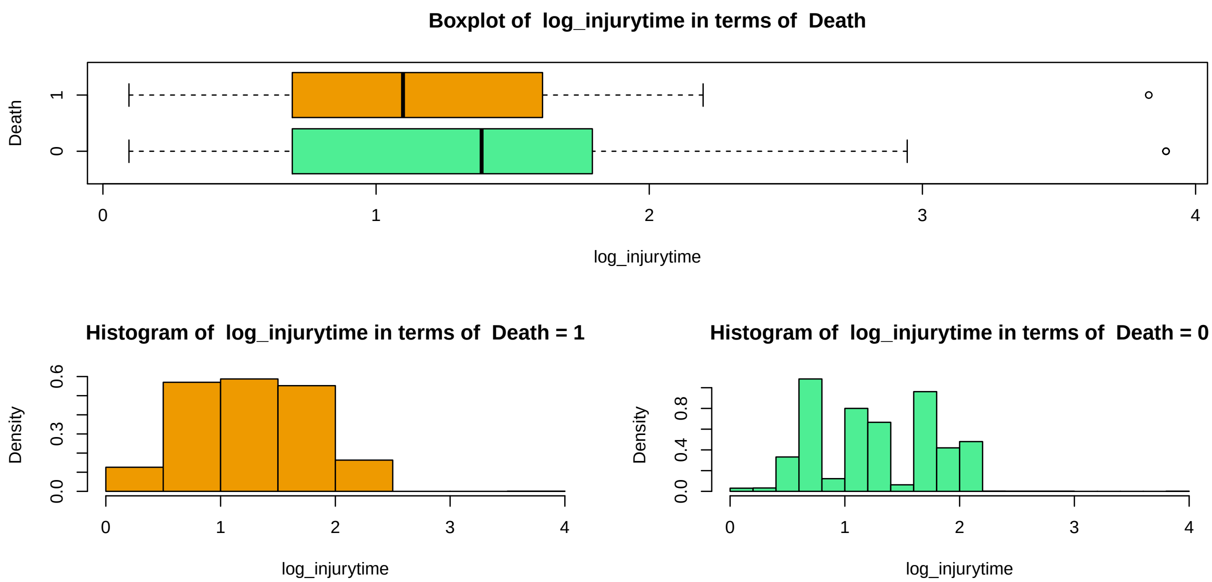

Figure 16: Distribution of the log injurytime in terms of death

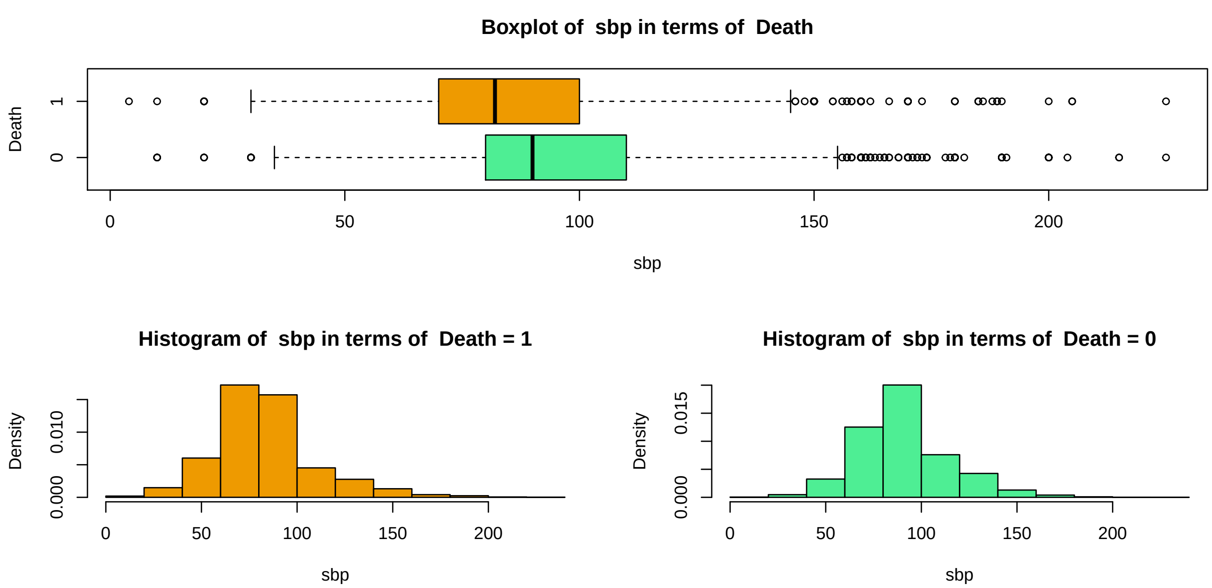

Figure 17: Distribution of sbp (Systolic Blood Pressure) in terms of death

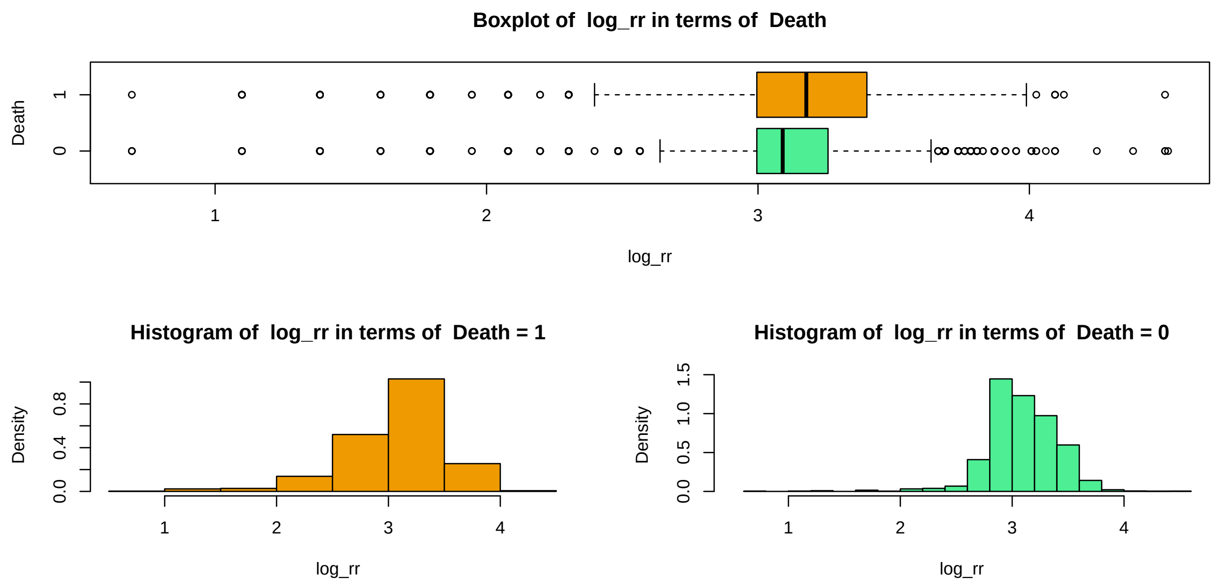

Figure 18: Distribution of log transformation of rr (Respiratory Rate) in terms of death

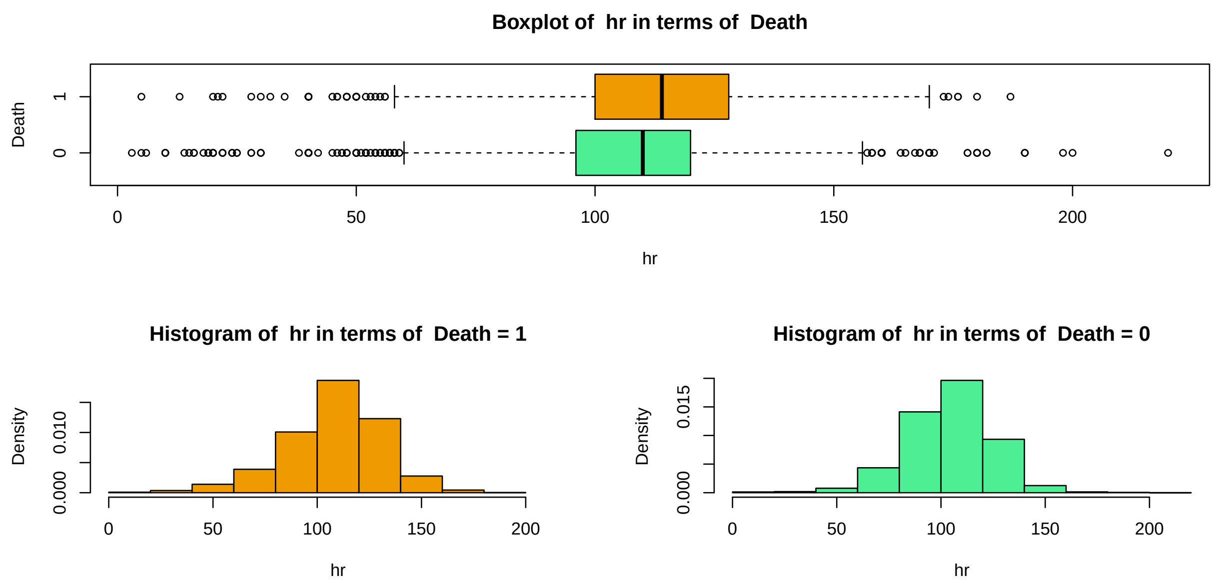

Figure 19: Distribution of hh (hearth Rate) in terms of death

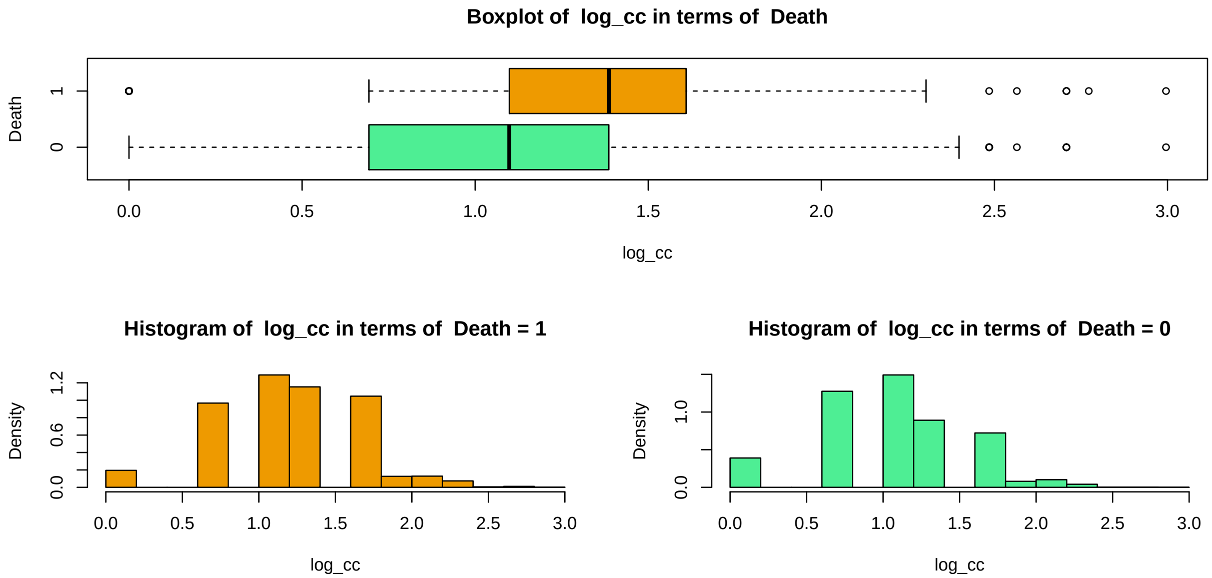

Figure 20: Distribution of log transformation of cc (Central capillary) in terms of death

Figure 21

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Figure 22 ...

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Finally, the scatter plot in figure 23 ...

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The PCP plot in figure 24...

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The Andrews' plot in figure 25...

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In conclusion ....

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In the next section we will make some inference using sample estimators of mean, covariance, correlation

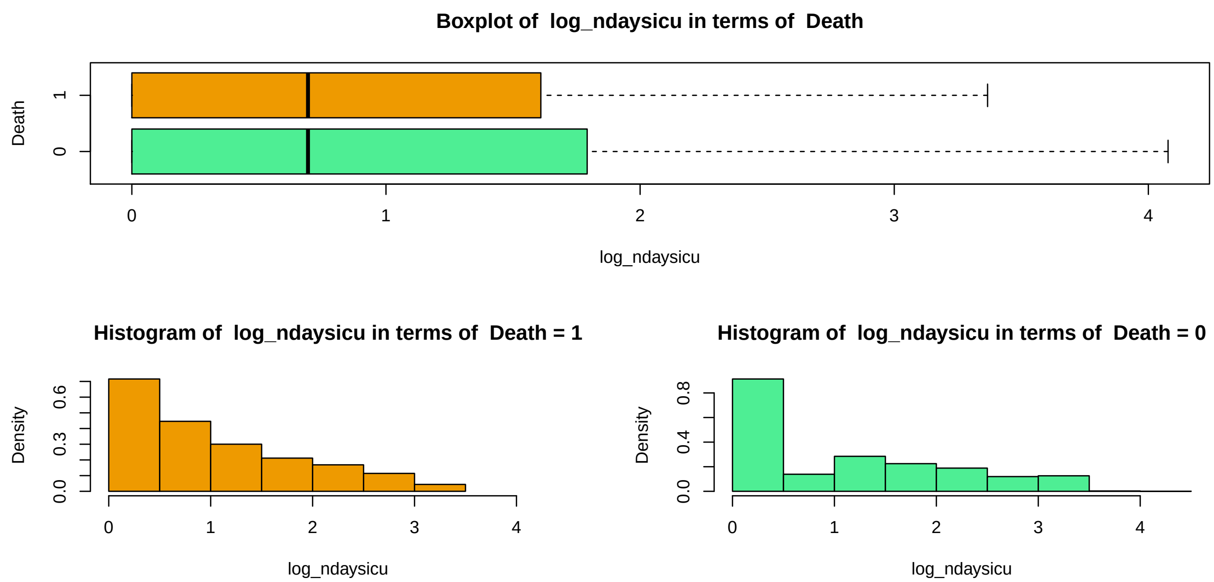

Figure 21: Distribution of log ndaysicu in terms of death

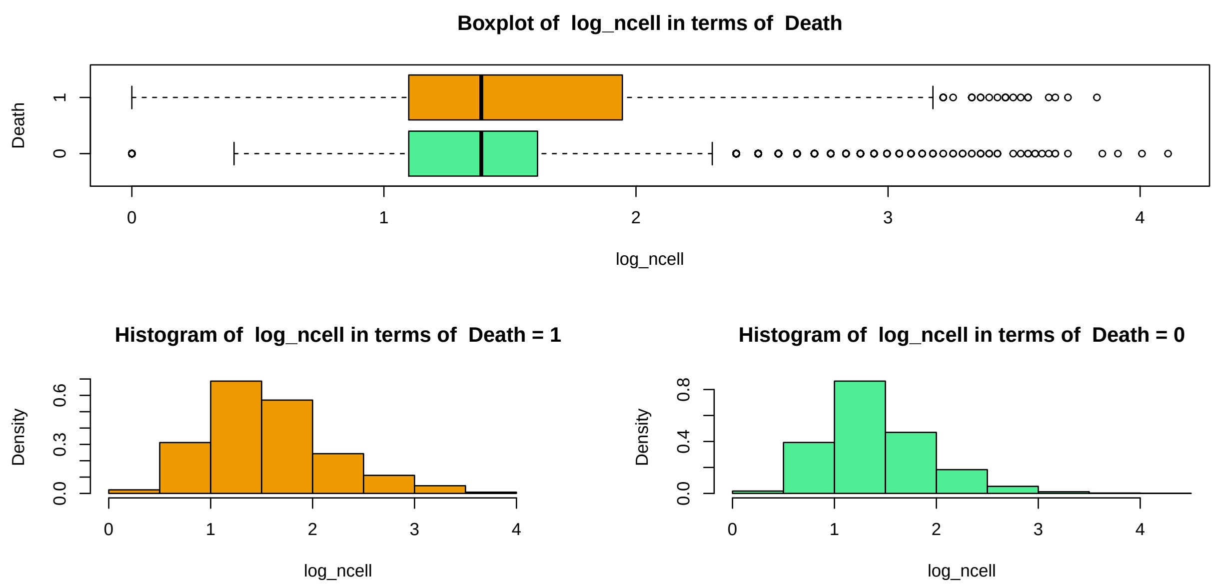

Figure 22: Distribution of log ncell in terms of death

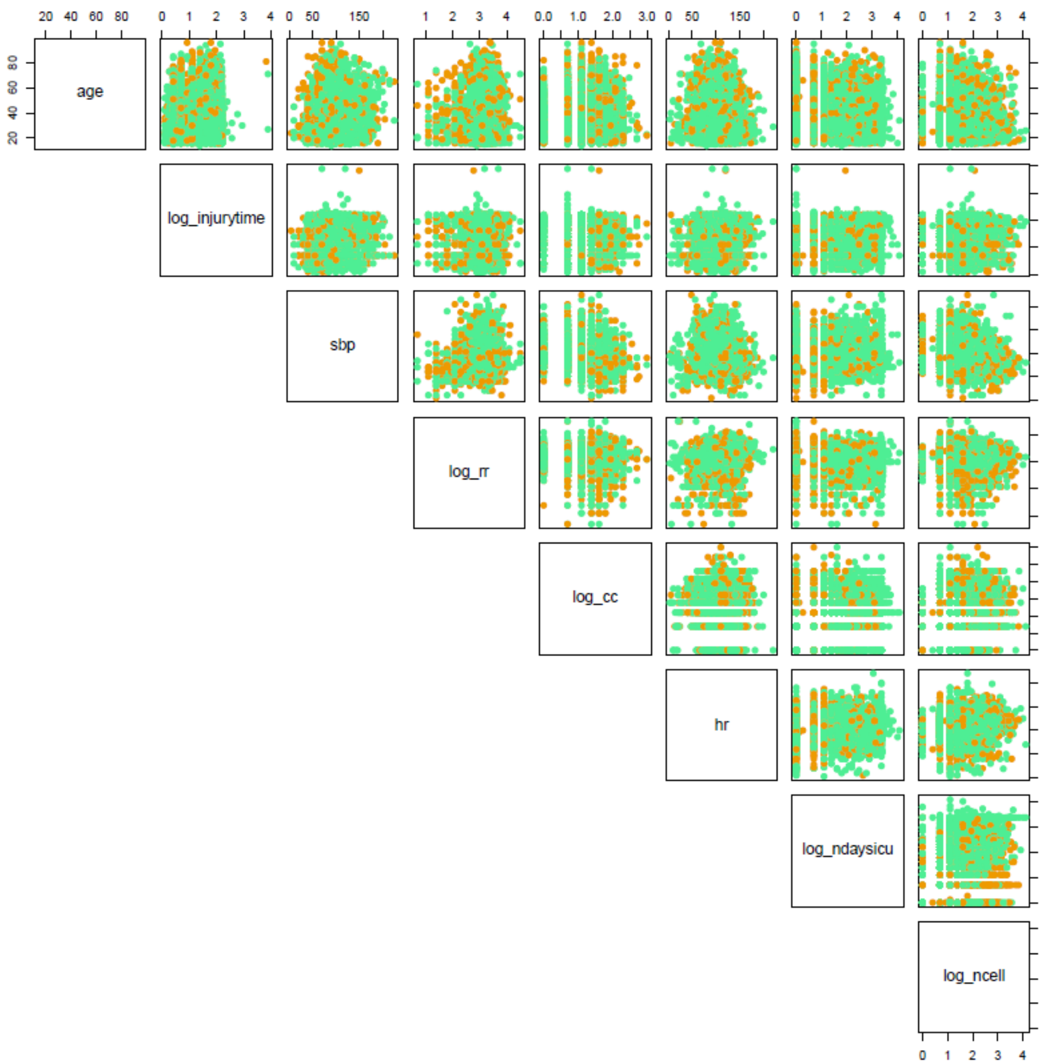

Figure 23: Scatter plot of all quantitiative variables

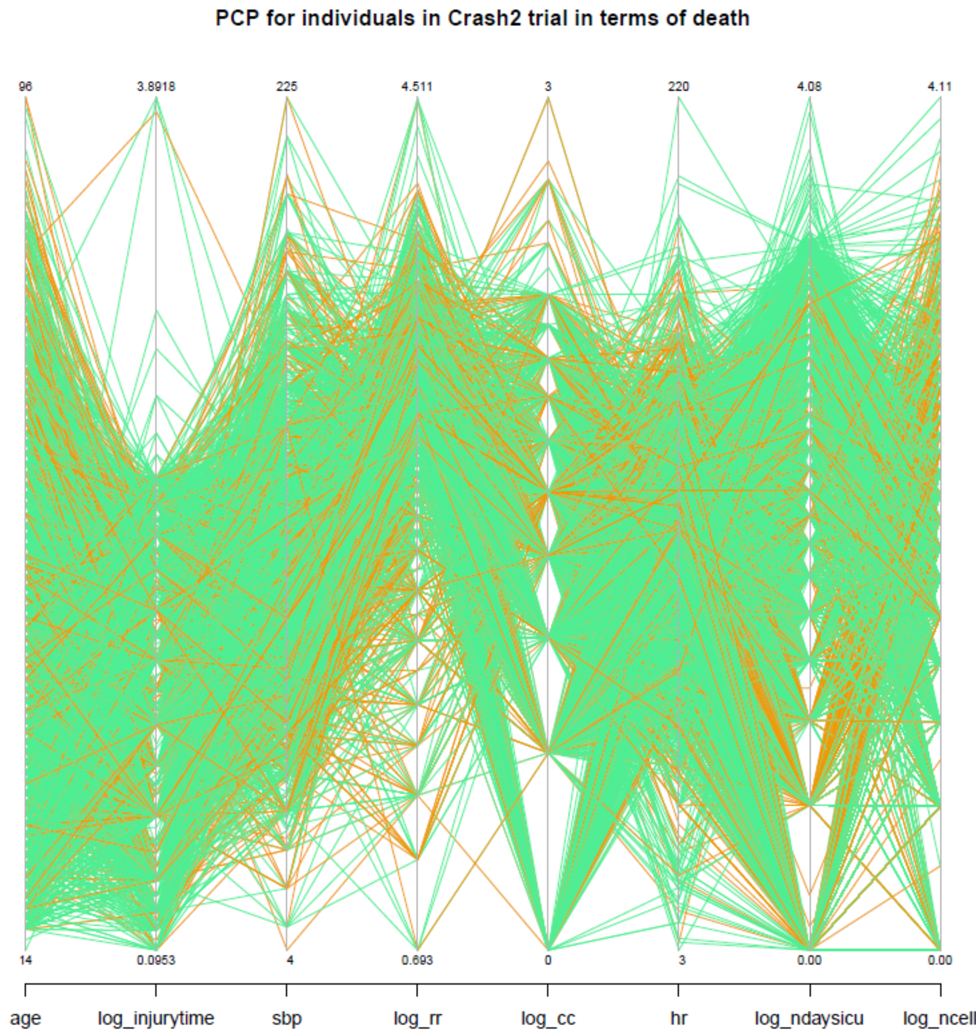

Figure 24: PCP plot of all quantitiative variables

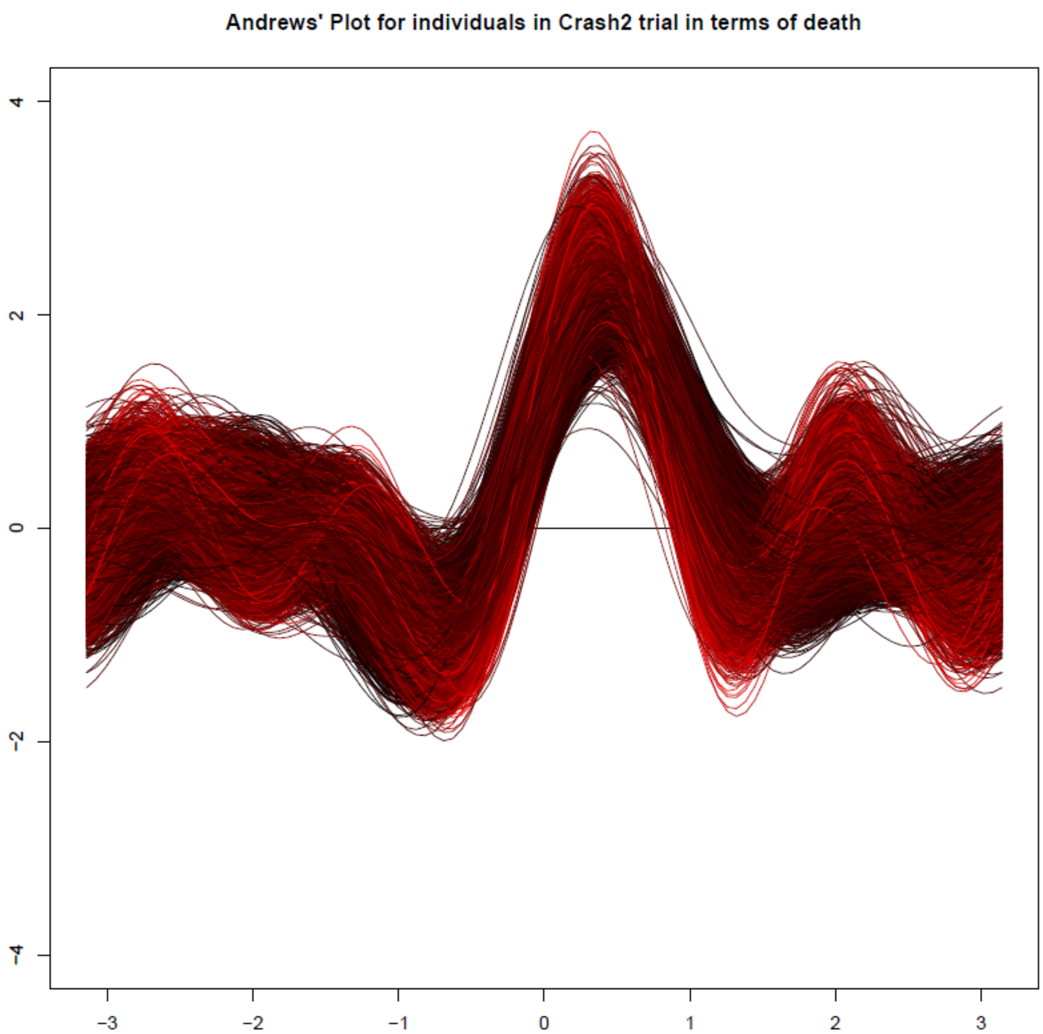

Figure 25: Andrews' plot of all quantitiative variables

Sample Estimators

Sample mean

All population...

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Table 3: Sample mean

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
34.66474	93.12909	108.0621	1.26759	3.106722	1.116876	0.9976935	1.394665

Population with death = 1 ...

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Table 4: Sample mean death

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
37.82849	87.33808	111.4559	1.240578	3.110272	1.232432	0.9390286	1.524647

Survival population

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Table 5: Sample mean survive

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
33.91215	94.50665	107.2548	1.274015	3.105878	1.089387	1.011649	1.363745

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Sample covariance

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Table 6: Sample covariance matrix

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
201.8020767	5.6687061	-26.9775900	0.5843356	0.0754442	0.3511496	1.1379316	0.1173186
5.6687061	604.1019094	-117.2846636	1.8258493	-0.3695028	-2.8828312	1.6117323	-1.9082923
-26.9775900	-117.2846636	459.8457610	-0.0753408	1.3004116	0.8685547	0.9063157	1.4739129
0.5843356	1.8258493	-0.0753408	0.2877629	-0.0037522	0.0126630	0.0863252	0.0034858
0.0754442	-0.3695028	1.3004116	-0.0037522	0.1092305	0.0119420	-0.0260511	-0.0025718
0.3511496	-2.8828312	0.8685547	0.0126630	0.0119420	0.2448313	0.0636306	0.0344845
1.1379316	1.6117323	0.9063157	0.0863252	-0.0260511	0.0636306	1.1508972	0.1949716
0.1173186	-1.9082923	1.4739129	0.0034858	-0.0025718	0.0344845	0.1949716	0.3404679

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Table 7: Sample covariance matrix death patients

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
261.7057666	17.272379	-51.3011548	0.8025150	0.0945272	0.3294187	1.1929405	-0.3849756
17.2723792	705.035309	-116.7715375	2.4101714	0.4360450	-3.2088292	4.1185430	-2.2891556
-51.3011548	-116.771537	564.8052079	0.0045026	1.7090123	1.1424702	0.7975679	1.9352979
0.8025150	2.410171	0.0045026	0.2799950	0.0020301	0.0093817	0.0999836	-0.0265201
0.0945272	0.436045	1.7090123	0.0020301	0.1899808	0.0014615	-0.0137661	-0.0143161
0.3294187	-3.208829	1.1424702	0.0093817	0.0014615	0.2228162	-0.0160445	0.0149818
1.1929405	4.118543	0.7975679	0.0999836	-0.0137661	-0.0160445	0.8604265	0.1140010
-0.3849756	-2.289156	1.9352979	-0.0265201	-0.0143161	0.0149818	0.1140010	0.4261349

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Table 8: Sample covariance matrix survival patients

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
184.6367863	8.3060201	-24.3595684	0.5577010	0.0676092	0.2486948	1.1796601	0.1156599
8.3060201	570.3044793	-111.6340107	1.6410806	-0.5550393	-2.6086143	0.9158253	-1.5962997
-24.3595684	-111.6340107	431.5566186	-0.0673383	1.1998768	0.6880413	0.9909257	1.2344841
0.5577010	1.6410806	-0.0673383	0.2894326	-0.0050994	0.0143642	0.0826221	0.0116551
0.0676092	-0.5550393	1.1998768	-0.0050994	0.0900403	0.0143148	-0.0289142	0.0000845
0.2486948	-2.6086143	0.6880413	0.0143642	0.0143148	0.2461653	0.0845804	0.0347028
1.1796601	0.9158253	0.9909257	0.0826221	-0.0289142	0.0845804	1.2191014	0.2164958
0.1156599	-1.5962997	1.2344841	0.0116551	0.0000845	0.0347028	0.2164958	0.3151667

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Sample correlation

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Table 9: Sample covariance matrix

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
1.0000000	0.0162355	-0.0885593	0.0766800	0.0160691	0.0499569	0.0746681	0.0141536
0.0162355	1.0000000	-0.2225256	0.1384817	-0.0454874	-0.2370449	0.0611251	-0.1330613
-0.0885593	-0.2225256	1.0000000	-0.0065495	0.1834860	0.0818573	0.0393963	0.1177952
0.0766800	0.1384817	-0.0065495	1.0000000	-0.0211642	0.0477075	0.1500037	0.0111366
0.0160691	-0.0454874	0.1834860	-0.0211642	1.0000000	0.0730252	-0.0734743	-0.0133361
0.0499569	-0.2370449	0.0818573	0.0477075	0.0730252	1.0000000	0.1198711	0.1194407
0.0746681	0.0611251	0.0393963	0.1500037	-0.0734743	0.1198711	1.0000000	0.3114691
0.0141536	-0.1330613	0.1177952	0.0111366	-0.0133361	0.1194407	0.3114691	1.0000000

Figure 26 …

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Table 10: Sample covariance matrix death patients

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
1.0000000	0.0402106	-0.1334354	0.0937500	0.0134059	0.0431388	0.0794978	-0.0364547
0.0402106	1.0000000	-0.1850471	0.1715408	0.0376765	-0.2560164	0.1672173	-0.1320676
-0.1334354	-0.1850471	1.0000000	0.0003580	0.1649837	0.1018409	0.0361794	0.1247455
0.0937500	0.1715408	0.0003580	1.0000000	0.0088021	0.0375608	0.2037027	-0.0767761
0.0134059	0.0376765	0.1649837	0.0088021	1.0000000	0.0071034	-0.0340485	-0.0503150
0.0431388	-0.2560164	0.1018409	0.0375608	0.0071034	1.0000000	-0.0366435	0.0486202
0.0794978	0.1672173	0.0361794	0.2037027	-0.0340485	-0.0366435	1.0000000	0.1882687

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
-0.0364547	-0.1320676	0.1247455	-0.0767761	-0.0503150	0.0486202	0.1882687	1.0000000

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Table 11: Sample covariance matrix survival patients

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
1.0000000	0.0255965	-0.0862962	0.0762902	0.0165817	0.0368888	0.0786281	0.0151619
0.0255965	1.0000000	-0.2250216	0.1277329	-0.0774554	-0.2201623	0.0347328	-0.1190669
-0.0862962	-0.2250216	1.0000000	-0.0060252	0.1924861	0.0667548	0.0432018	0.1058514
0.0762902	0.1277329	-0.0060252	1.0000000	-0.0315884	0.0538137	0.1390921	0.0385896
0.0165817	-0.0774554	0.1924861	-0.0315884	1.0000000	0.0961511	-0.0872717	0.0005018
0.0368888	-0.2201623	0.0667548	0.0538137	0.0961511	1.0000000	0.1543962	0.1245894
0.0786281	0.0347328	0.0432018	0.1390921	-0.0872717	0.1543962	1.0000000	0.3492684
0.0151619	-0.1190669	0.1058514	0.0385896	0.0005018	0.1245894	0.3492684	1.0000000

Figure 27…

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Outliers

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Table 12: Sample robust media all population

age	sbp	hr	log_injurytime	log_rr	log_cc	log_ndaysicu	log_ncell
33.94433	92.32304	108.848	1.271321	3.137716	1.106643	0.9422688	1.357334

Comparison of the eigenvalues of the covariance matrix for sample and robust is given by figure 28 …

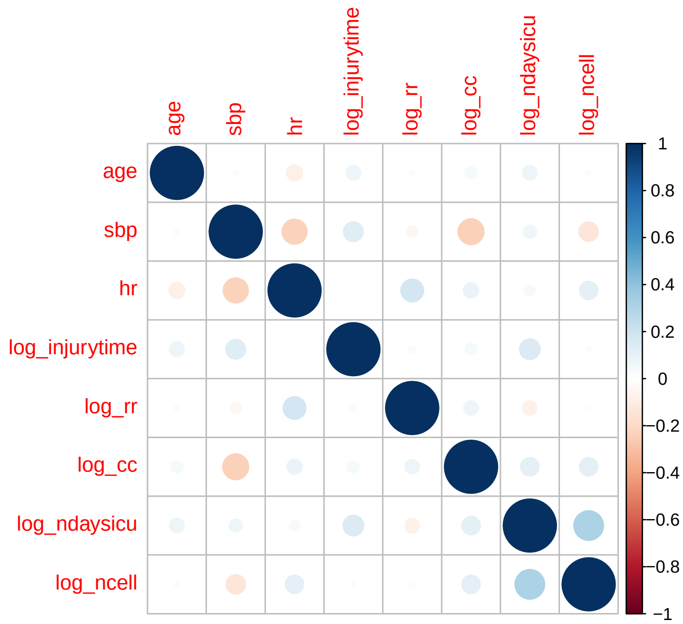
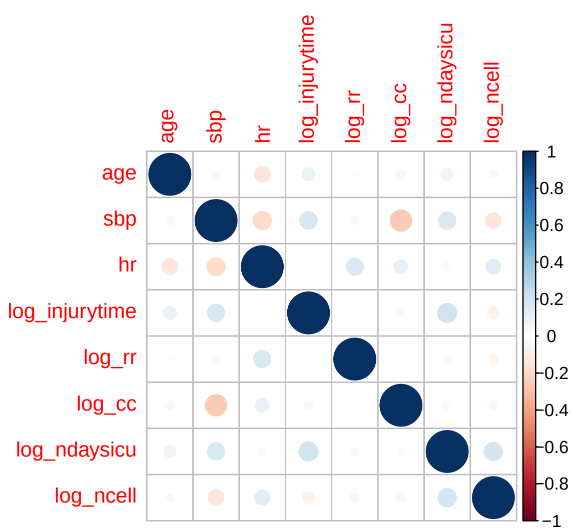


Figure 26: Sample Correlation Crash2 Dataset



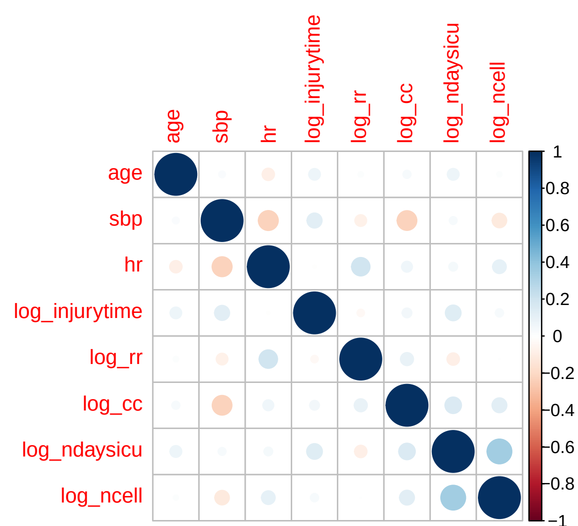


Figure 27: Sample Correlation for death and survival patients from Crash2 Dataset

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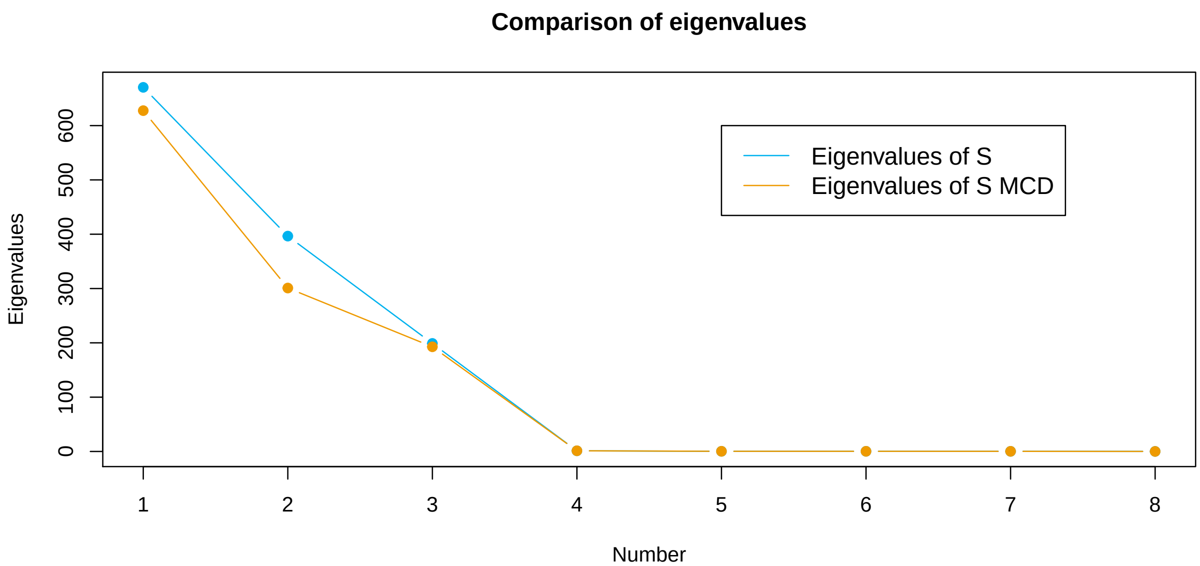

Figure 28: Comparison eigenvalues sample and robust covariance

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figure 29 Lorem ipsum dolor sit amet, consectetur adipiscing elit, sed do eiusmod tempor incididunt ut labore et dolore magna aliqua. Ut enim ad minim veniam, quis nostrud exercitation ullamco laboris nisi ut aliquip ex ea commodo consequat.

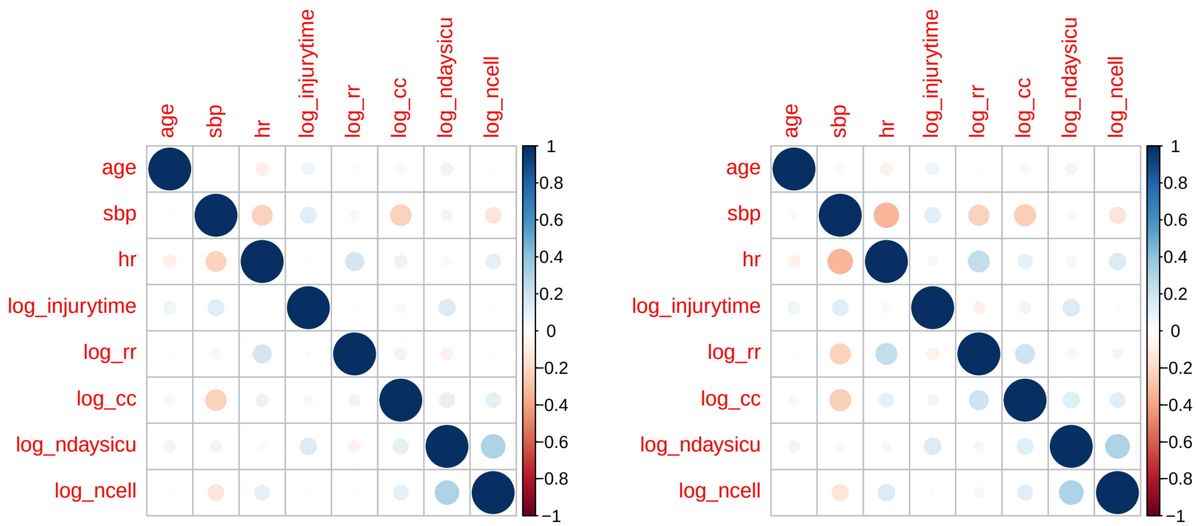

Figure 29: Correlation all population and MCD correlation

Principal Component Analysis

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PCA Complete dataset

figure 31...

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figure 31...

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figure 33...

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figure 35...

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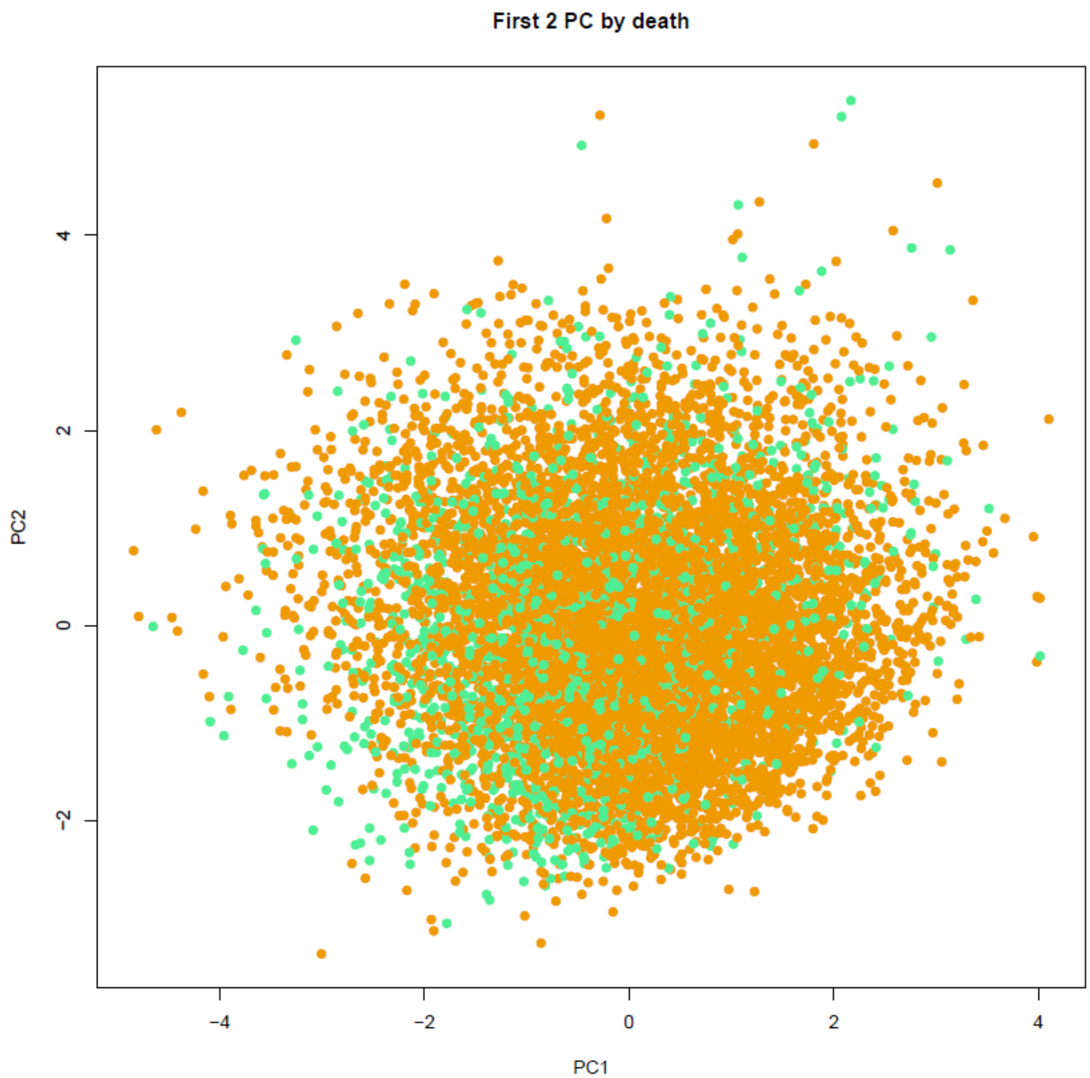

Figure 30: First two PCs based on the sample covariance all population

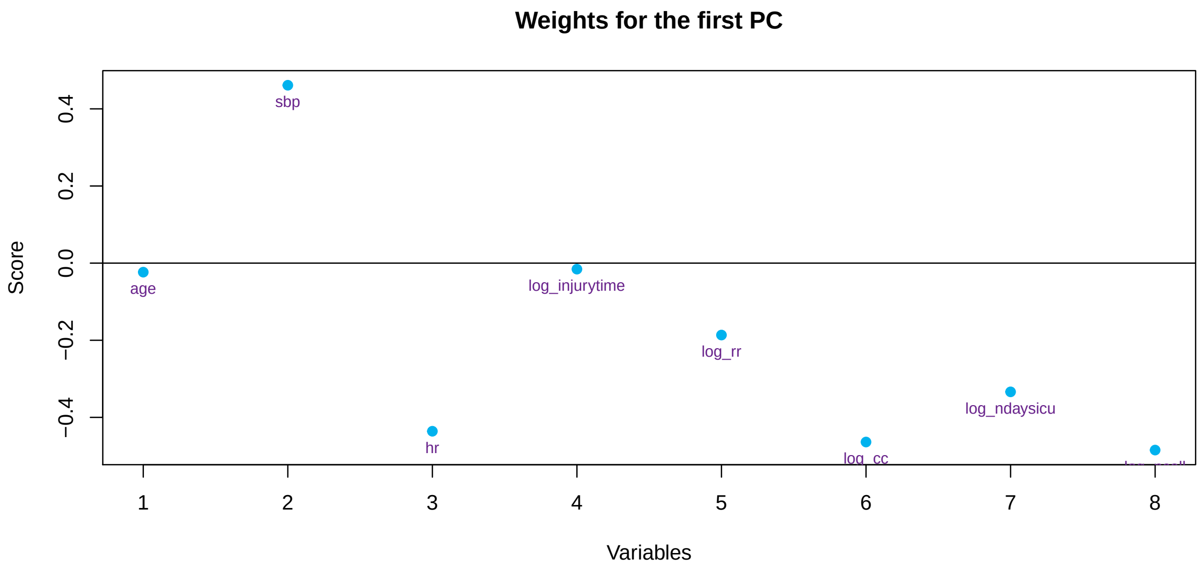

Figure 31: Loadings for the first PC

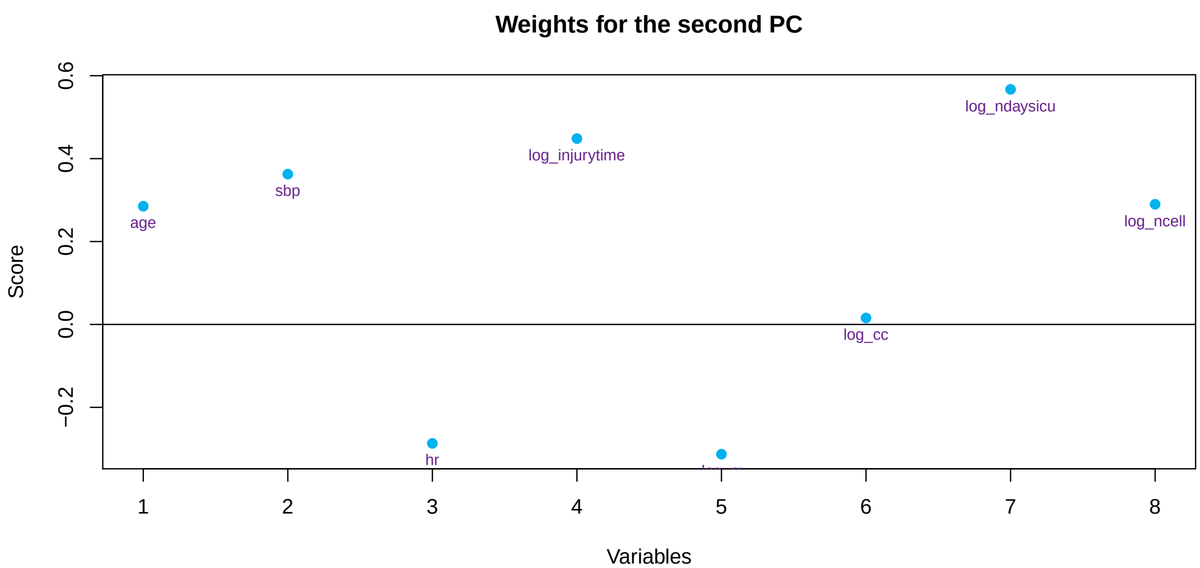

Figure 32: Loadings for the second PC

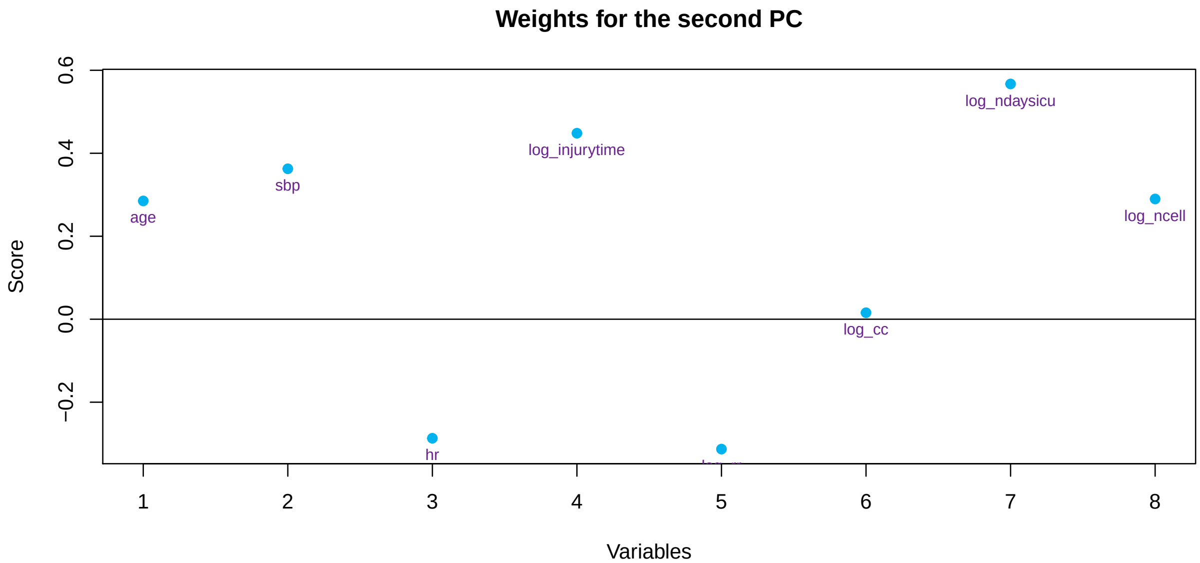

Figure 33: Loadings for the first two PC

commodo consequat.

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figure 37...

IS NECESSARY TO DO THE SAME BY CLASS?

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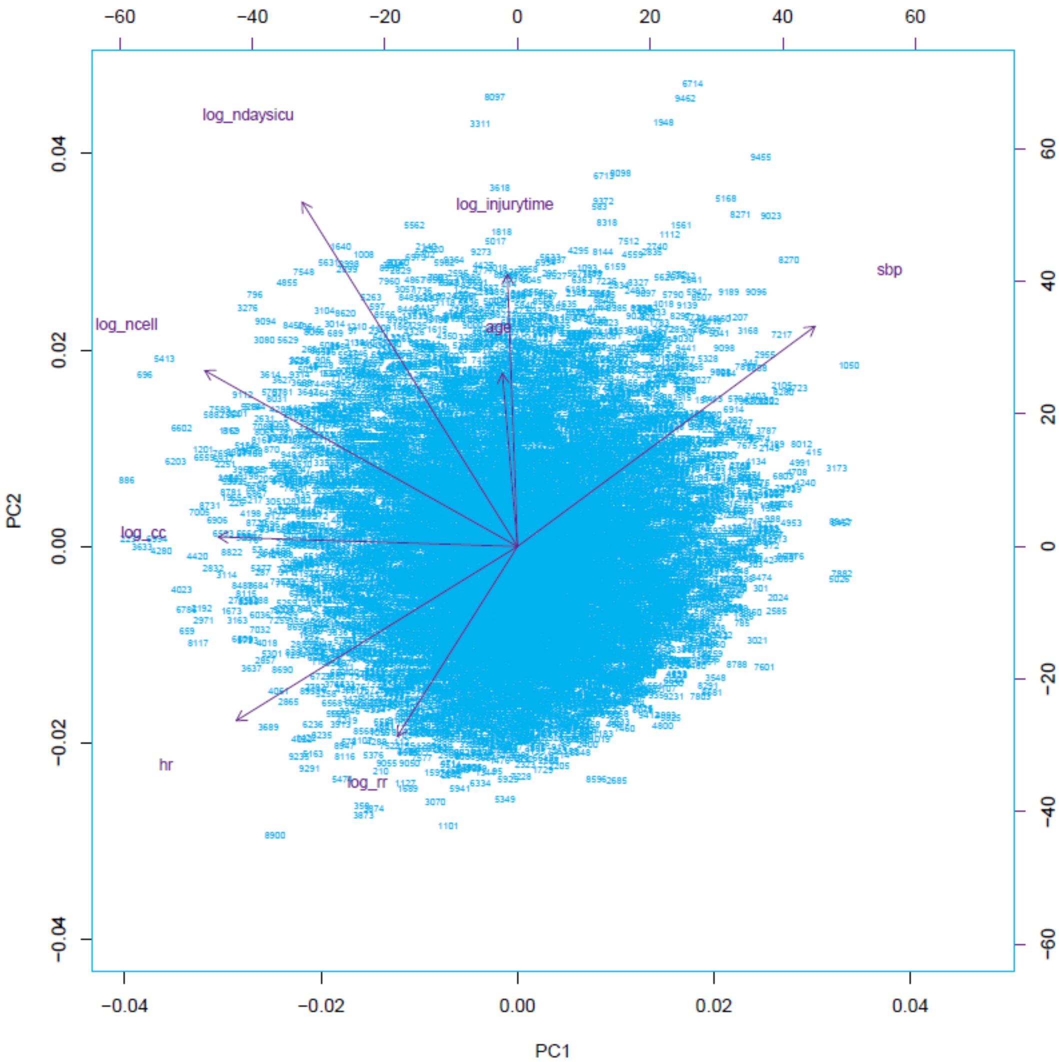

Figure 34: PC Scores and PC loading all population

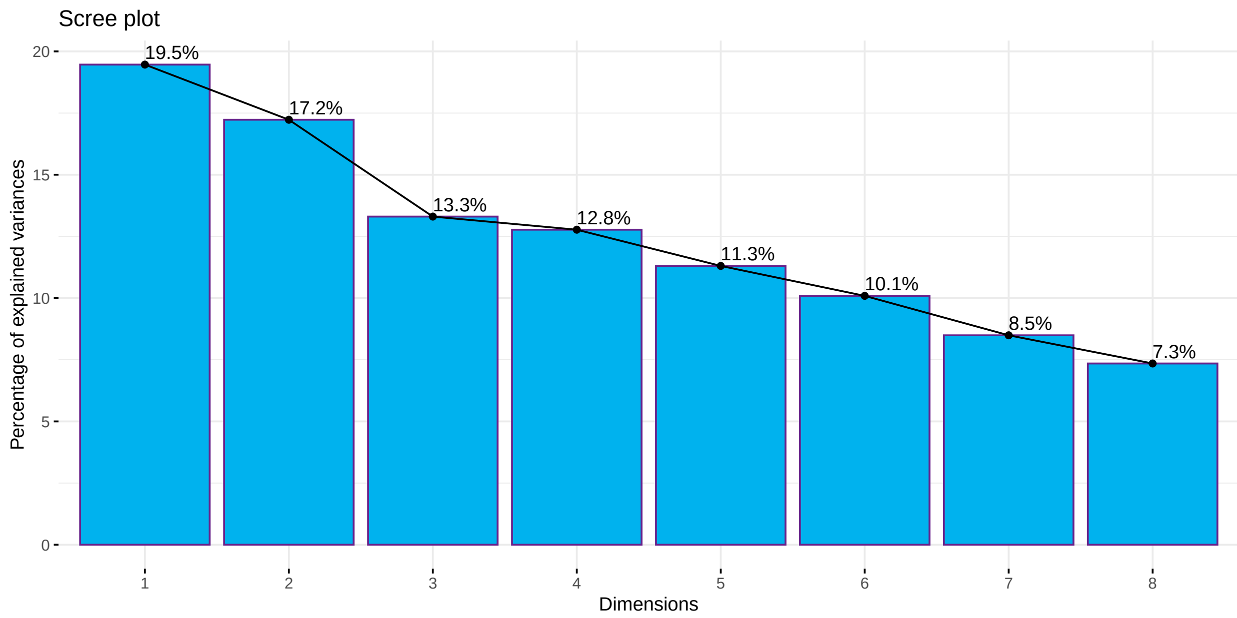

Figure 35: Eigenvalues of the sample correlation matrix

PCA by Category

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Important. For these plot the groups are now female (blue) and male (gray)

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figure 39…

figure 40…

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Figures 42 and 43 …

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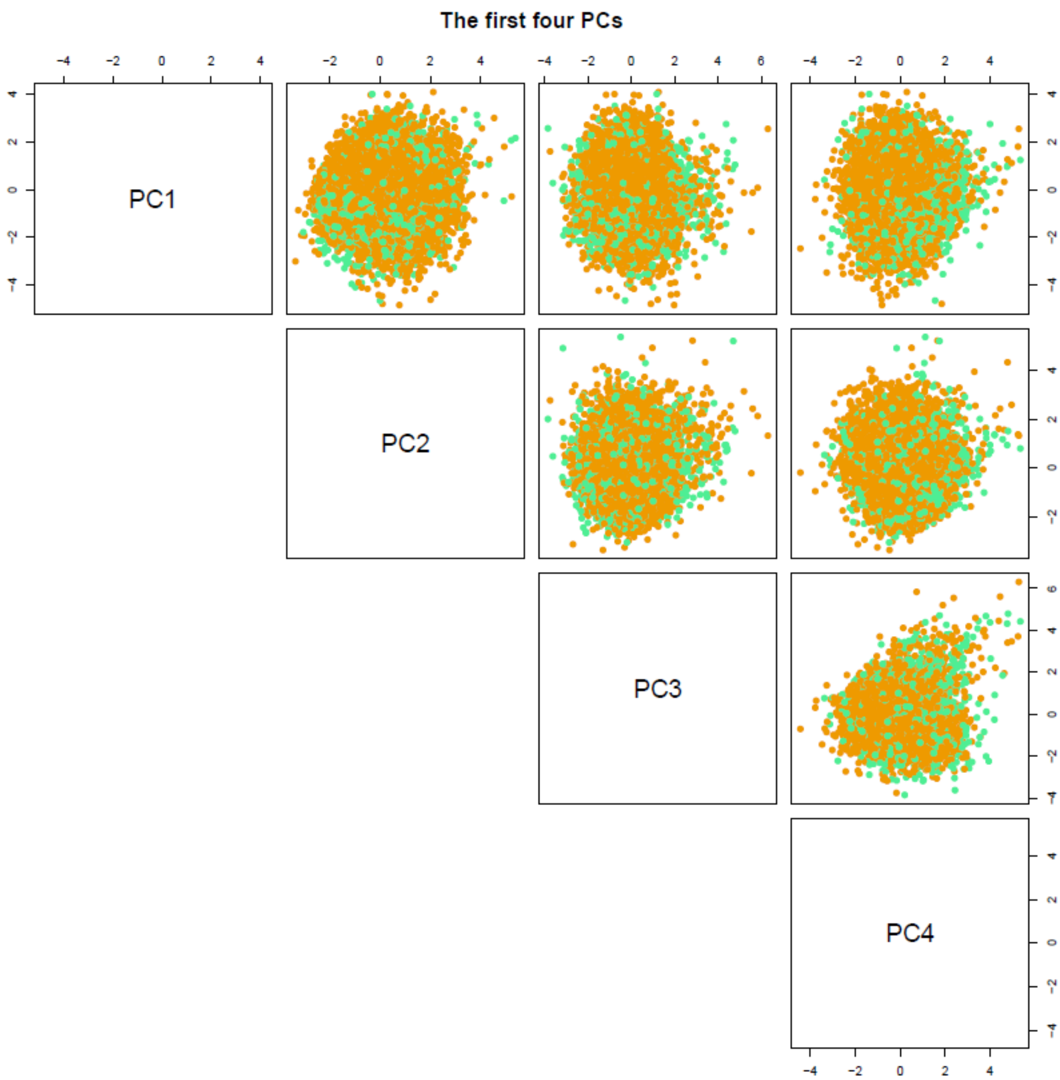

Figure 36: PC Scores and PC loading all population

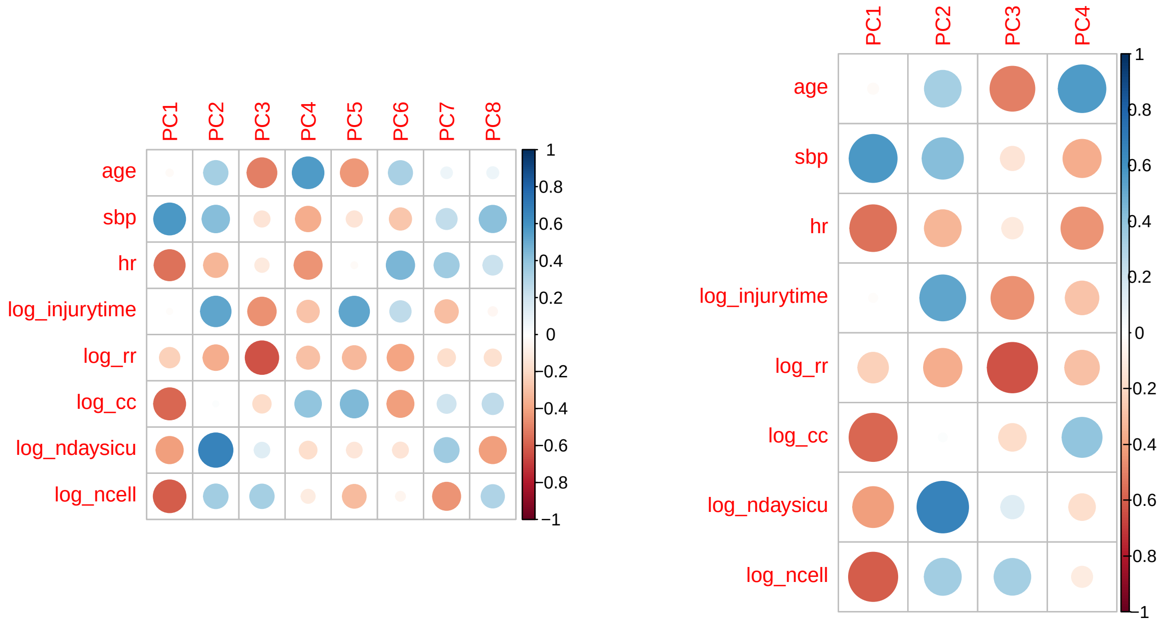

Figure 37: Correlation between dataset and all PC and the first 4 PCs

commodo consequat.

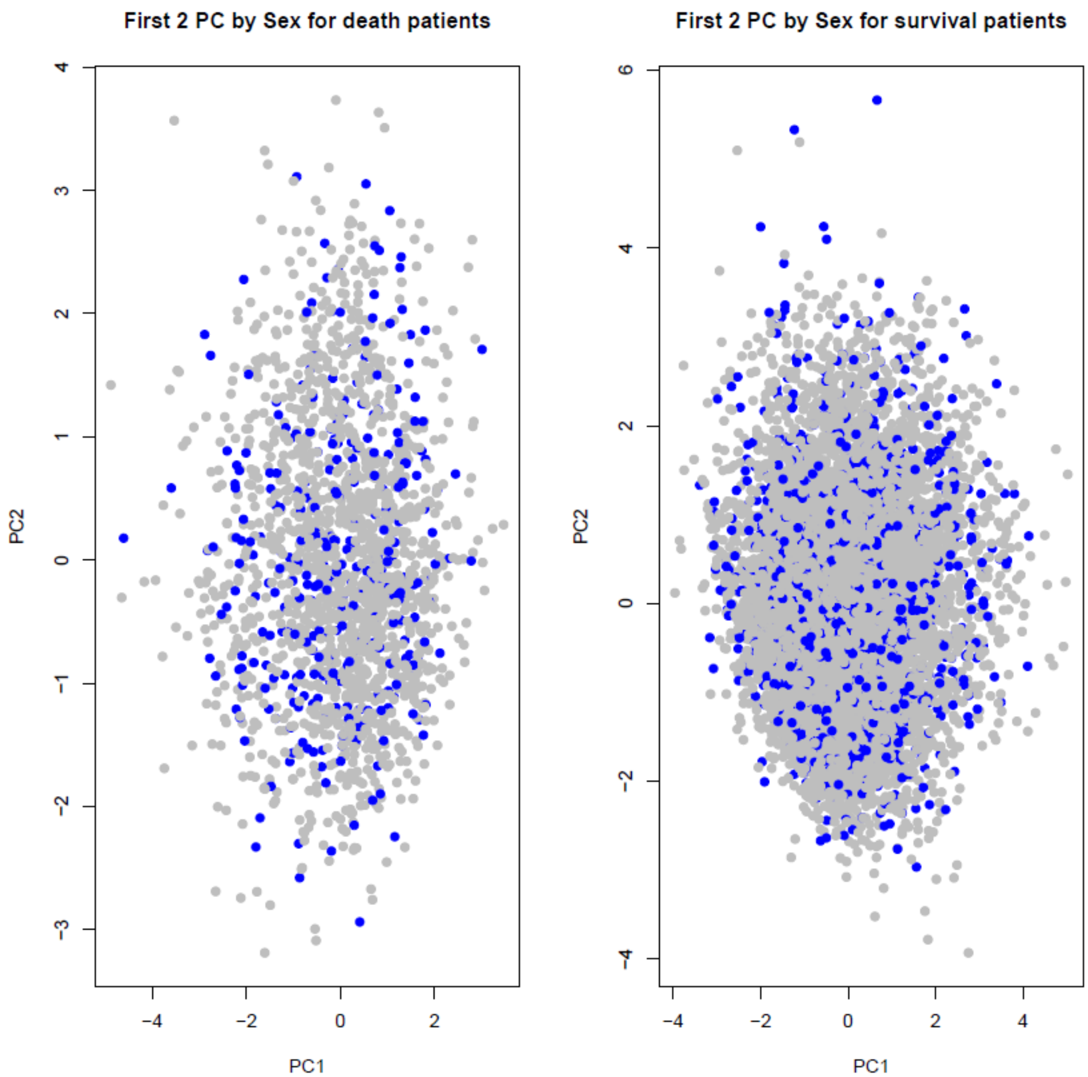

Figure 38: PC Scores and PC loading all population

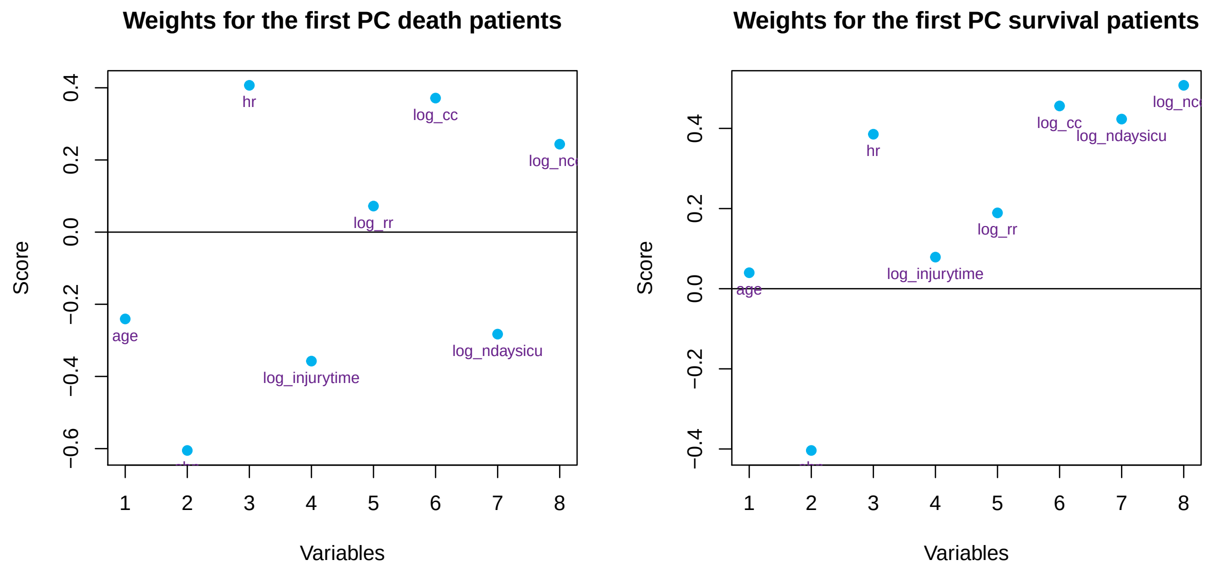

Figure 39: Loadings for the first PC by death and survival patients

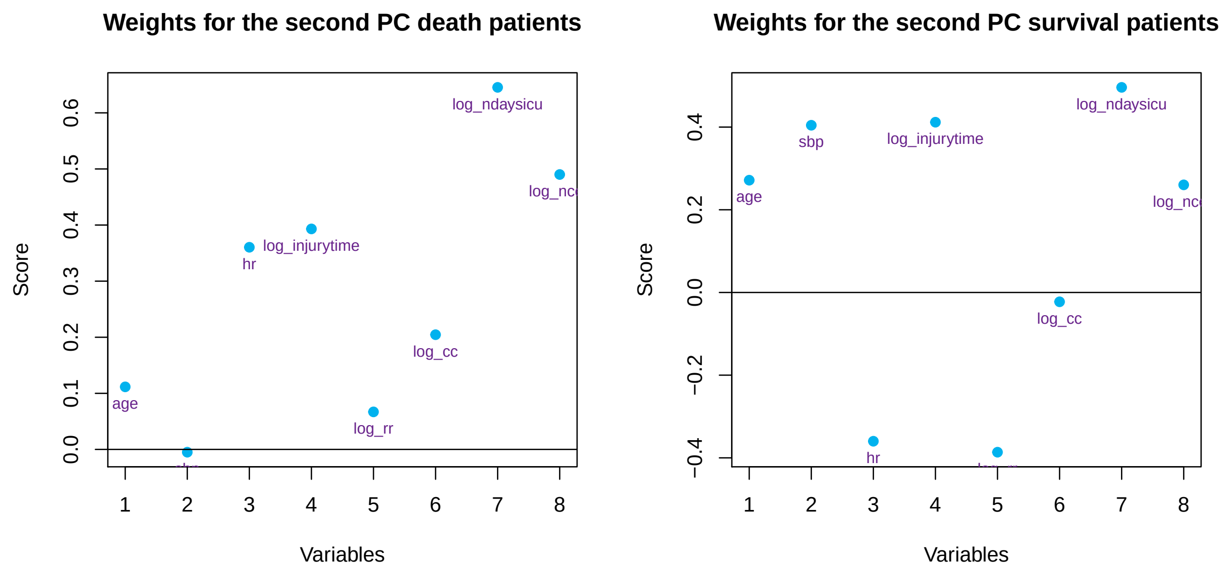

Figure 40: Loadings for the second PC by death and survival patients

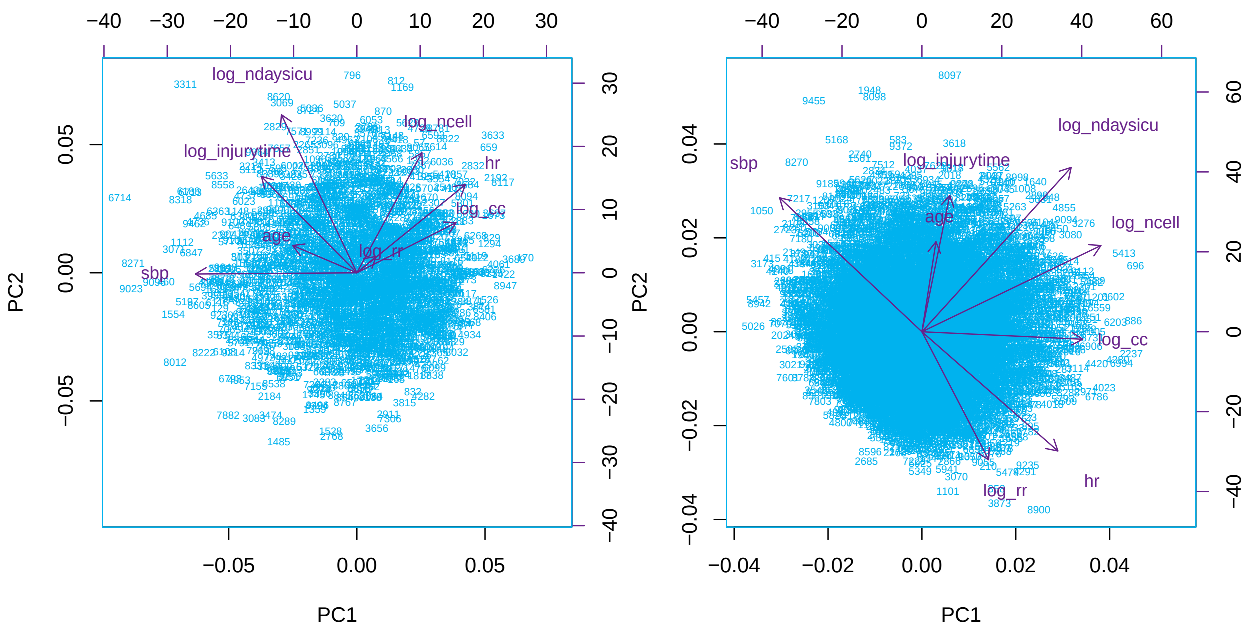

Figure 41: PC Scores and PC loading all population for death and survival patiens

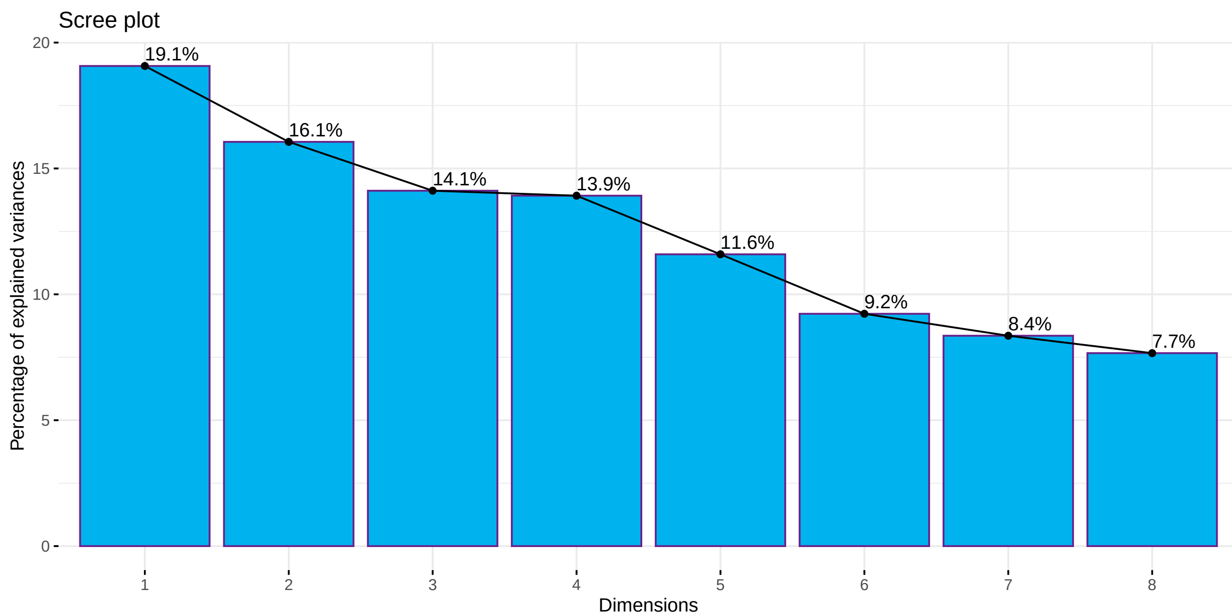

Figure 42: Eigenvalues of the sample correlation matrix for death patients

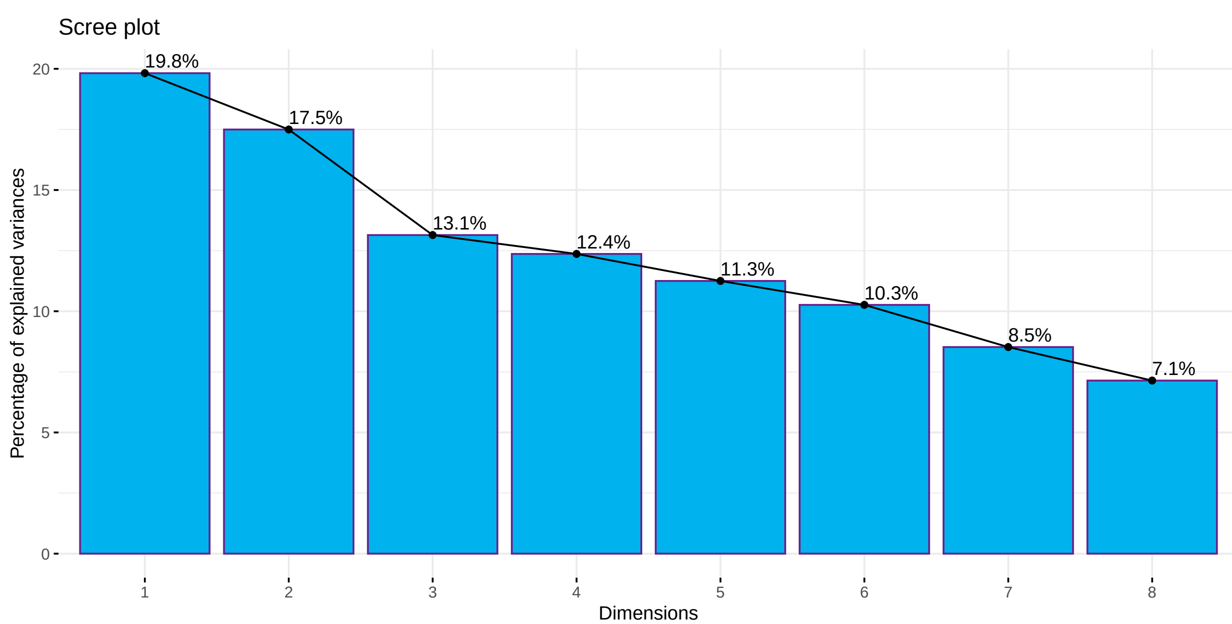

Figure 43: Eigenvalues of the sample correlation matrix for survival patients

Conclusions

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