HW2_sliang

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Problem 3

Personlly, as a researcher on data science, I tune the hyperparameter all the time, and it's had to ensure every tuning can improve the model. So, version control is important for me to recall to any previous conditions of my code.

Problem 4

Load package

```
library(data.table)
library('magrittr')
library('tidyverse')
                                                  ----- tidyverse 1.3.0 --
## -- Attaching packages -----
## v ggplot2 3.2.1
                     v purrr
                              0.3.3
## v tibble 2.1.3
                     v dplyr
                              0.8.3
          1.0.2
## v tidvr
                     v stringr 1.4.0
           1.3.1
## v readr
                     v forcats 0.5.0
## -- Conflicts ----- tidyverse conflicts() --
## x dplyr::between()
                     masks data.table::between()
## x tidyr::extract() masks magrittr::extract()
## x dplyr::filter()
                     masks stats::filter()
## x dplyr::first()
                     masks data.table::first()
## x dplyr::lag()
                      masks stats::lag()
## x dplyr::last()
                      masks data.table::last()
## x purrr::set_names() masks magrittr::set_names()
## x purrr::transpose() masks data.table::transpose()
```

Qusetion a

```
url_sen<-"https://www2.isye.gatech.edu/~jeffwu/wuhamadabook/data/Sensory.dat"
df_sen<-read.table(url_sen, skip=1, fill=TRUE, header=TRUE)</pre>
```

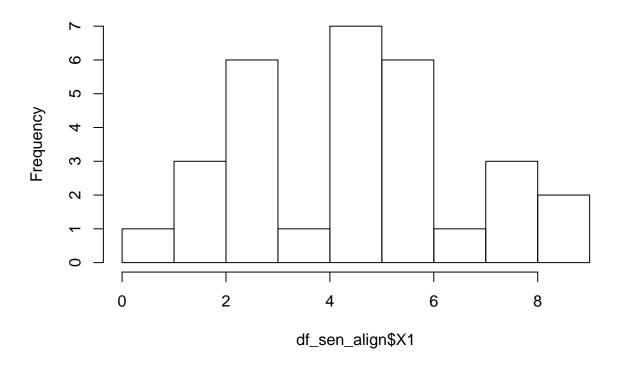
The issues in this data is the location of missing value is incorrect, so we use this function to align the data.

```
align_data <- function(row_){
   if (is.na(row_['X5'])){
     row_[2:6] <- row_[1:5]</pre>
```

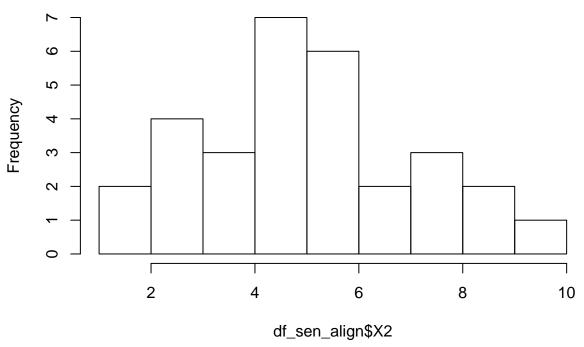
```
row_[1] <- NA
}
return(row_)
}
df_sen_align<-data.table(t(apply(df_sen, 1, align_data)))</pre>
```

Result

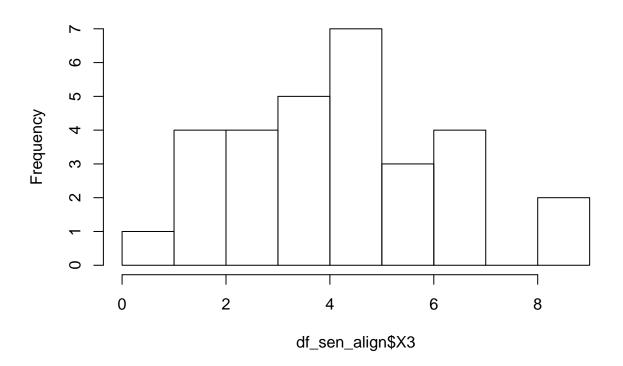
Histogram of df_sen_align\$X1



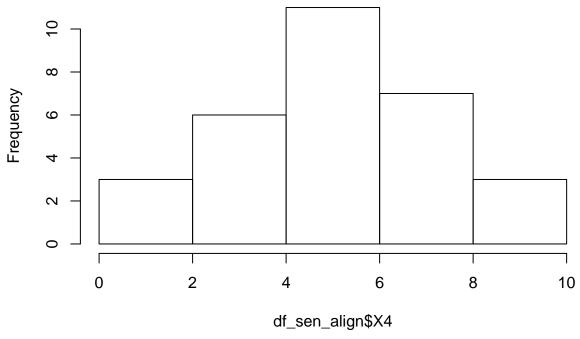
Histogram of df_sen_align\$X2



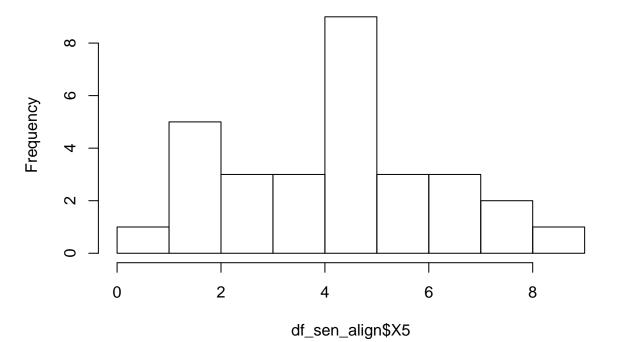
Histogram of df_sen_align\$X3



Histogram of df_sen_align\$X4



Histogram of df_sen_align\$X5



_					
	X1	X2	X3	X4	X5
	Min. :0.900	Min. :1.500	Min. :0.800	Min. :0.900	Min. :0.700
	1st Qu.:2.850 Median :4.550	1st Qu.:3.450 Median :4.950	1st Qu.:2.650 Median :4.150	1st Qu.:3.925 Median :5.400	1st Qu.:2.250 Median :4.600

X1	X2	X3	X4	X5
Mean :4.593	Mean :5.063	Mean :4.167	Mean :5.193	Mean :4.267
3rd Qu.:5.950	3rd Qu.:6.225	3rd Qu.:5.400	3rd Qu.:6.275	3rd Qu.:5.800
Max. :9.000	Max. :9.200	Max. :9.000	Max. :9.400	Max. :8.800

Question b

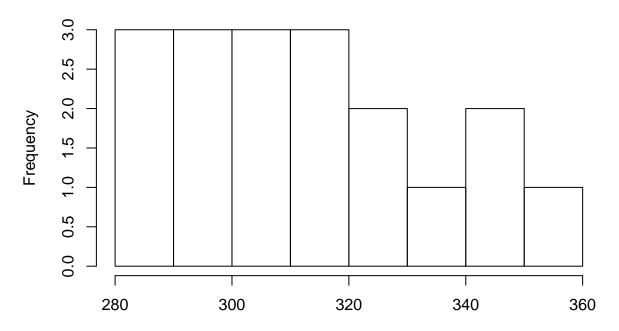
```
url_gm<-"https://www2.isye.gatech.edu/~jeffwu/wuhamadabook/data/LongJumpData.dat"
df_gm<-read.table(url_gm, skip=1, fill=TRUE, header=TRUE)</pre>
```

The issue in this data is that the columns are messy, so we should reorganized it.

```
colnames(df_gm)<- NA
df_gm_organized <- rbind(df_gm[,1:2],df_gm[,3:4],df_gm[,5:6],df_gm[,7:8])
colnames(df_gm_organized) <- c('Year', 'Long_jump_data')</pre>
```

Result

Histogram of df_gm_organized\$Long_jump_data



df_gm_organized\$Long_jump_data

 Long_jump_data
Min. :281.5
1st Qu.:298.3
Median :312.7
Mean :313.3
3rd Qu.:327.5
Max. $:350.5$
NA's :2

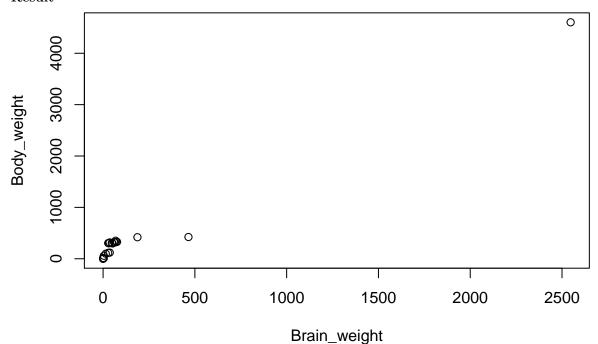
Question c

```
url_bw<-"https://www2.isye.gatech.edu/~jeffwu/wuhamadabook/data/BrainandBodyWeight.dat"
df_bw<-read.table(url_bw, skip=1, fill=TRUE, header=TRUE)</pre>
```

The issue in this data is similar to question b.

```
colnames(df_bw)<- NA
df_bw_organized <- rbind(df_bw[,1:2],df_gm[,3:4],df_gm[,5:6])
colnames(df_bw_organized) <- c('Brain_weight', 'Body_weight')</pre>
```

Result



Brain_weight	Body_weight
Min. : 0.005	Min. : 0.10
1st Qu.: 1.010	1st Qu.: 6.45
Median: 21.245	Median: 106.60
Mean: 127.691	Mean: 302.94
3rd Qu.: 58.000	3rd Qu.: 317.64
Max. $:2547.000$	Max. $:4603.00$

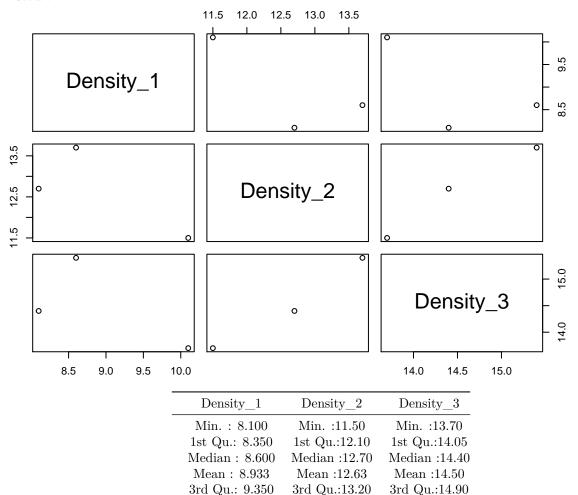
Question d

```
url_to<-"https://www2.isye.gatech.edu/~jeffwu/wuhamadabook/data/tomato.dat"
df_to<-read.table(url_to, skip=1, fill=TRUE, header=TRUE)</pre>
```

The issue in this data is they put three densities together and splited by comma, so we first convert them to vectors then to data table.

```
get_density <- function(element){
   return(as.numeric(unlist(strsplit(element,split=","))))
}
df_to_row <- data.table(apply(df_to[2,],2,get_density))
colnames(df_to_row) <- c('Density_1', 'Density_2', 'Density_3')</pre>
```

Result



Max. :10.100

Max. :13.70

Max. :15.40