Bioinformatics III

Seventh Assignment

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Exercise 7.1: Missing Data Imputation

(a) The script

```
Listing 1: Missing data imputation script
o print ("Assignment_7_-_Schmitt_Schowing")
   \#setwd("C:/Users/thsch/Desktop/Bioinformatics3/Assignments/Assignment7/Scripts
5 data = read.table("ms_toy.txt", header=TRUE)
   summary (data)
   # class(data) -> data.frame
10 # Takes a dataframe and the col name and calculate the values for the missing
        data
   # as if they were data "under the detection threshold"
   impute_missing_data <- function(data, colname, QUANTILE_VALUE, FRACTION,
        replace = FALSE) {
     \begin{array}{lll} current\_sd & <& \mathbf{apply}(\,\mathrm{data}\,[\,\mathrm{colname}\,]\,\,,\,\,2\,,\,\,\mathrm{sd}\,,\,\,\mathrm{na.rm}\,=\,\mathrm{TRUE})\\ current\_mean & <&- \mathbf{apply}(\,\mathrm{data}\,[\,\mathrm{colname}\,]\,,\,\,2\,,\,\,\mathrm{mean}\,,\,\,\mathrm{na.rm}\,=\,\mathrm{TRUE}) \end{array}
     str = sprintf("Current_sd:_%f", current_sd)
     print(str)
     str = sprintf("Current_mean: _%f", current_mean)
     print(str)
     \# Chosing the mean of the new distribution -\!\!> the 5% quantile for instance
           ?!?!?!
     #QUANTILE_VALUE <- 5
25
     quant <- quantile (data [colname], QUANTILE_VALUE/100, na.rm = TRUE)
     str = sprintf("Current_%d_percent_quantile:_%f", QUANTILE.VALUE, quant)
     print(str)
     # New mean equals the above quantile
     new_mean = quant
     # New sd -> fraction of the current sd
     #FRACTION <- 1/3
     new_sd <- FRACTION * current_sd
     print(paste0("New_mean:_", new_mean))
     print(paste0("New_sd:_", new_sd))
```

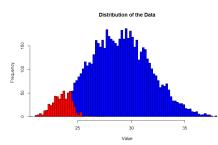
100

```
40
      # Display the distribution with the chosen mean
      hist(as.matrix(data[colname]), main = "Distribution_of_the_Data", xlab = "
            Value")
      abline (v=quant, col="red")
      # We have sd and mean -> rnorm(nb, sd, mean) will generage numbers in the
45
            distribution
      # Numbers of NA in the column (nb of data to generate):
      nb_na <- sum(is.na(data[colname]))
      generated_data = rnorm(nb_na, mean = new_mean, sd = new_sd)
      # Trying to make the plots nice but will work for ONE distribution
      p1 <- hist(generated_data, breaks = 30, freq = TRUE)
55
      p2 <- hist(as.matrix(data[colname]), breaks = 60, freq = TRUE)
      \# Acceptable graphs for any distribution
      \#p1 < -hist(generated\_data, freq = TRUE) \\ \#p2 < -hist(as.matrix(data[colname]), freq = TRUE)
      plot\left(p2\,,\ main\ =\ "Distribution\_of\_the\_Data"\,,\ xlab\ =\ "Value"\,,\ col\ =\ "blue"\,\right)
      plot(pl, main = "Distribution_of_the_Data", xlab = "Value", col = "red", add
            =T
65
      # Replace the data a copy of the original dataframe if the parameter "
            replace" is set to TRUE.
      # Also print the information of the data (copy) before and after the
            replacement
      # By default it's FALSE.
70
      if (replace){
         data.replace <- data
         print(summary(data.replace))
         data.replace[[colname]][which(is.na(data[colname]))] <- generated_data
75
         \# data \, [\lceil colname \, \rceil] \, [\lceil which \, (is \, . \, na \, (data \, [\lceil colname \, \rceil) \,) \,] \, < - \, \, generated \, \_data \, |
         p < - \ hist (as.matrix (data.replace [colname]) , breaks = 60 , \ freq = TRUE)
         plot(p, main = "Distribution_of_the_data_after_replacement", xlab = "Value
                  col = "blue")
         print(summary(data.replace))
80
   }
   # Playing with the new mean and sd values
impute_missing_data(data = data, 'ctrl.1', 1, 1/3)
impute_missing_data(data = data, 'ctrl.1', 5, 1/3)
impute_missing_data(data = data, 'ctrl.1', 5, 1/3)
impute_missing_data(data = data, 'ctrl.1', 10, 1/3)
impute_missing_data(data = data, 'ctrl.1', 15, 1/3)
impute_missing_data(data = data, 'ctrl.1', 20, 1/3)
   # 5 looks like the best new mean
impute_missing_data (data = data, 'ctrl.1', 5, 1)
impute_missing_data (data = data, 'ctrl.1', 5, 1/2)
impute_missing_data (data = data, 'ctrl.1', 5, 1/3)
impute_missing_data (data = data, 'ctrl.1', 5, 1/4)
impute_missing_data (data = data, 'ctrl.1', 5, 1/5)
```

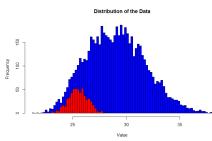
```
\# We chose the 5% quartile for the mean and 1 third of the standard deviation
       for the new sd.
   impute_missing_data(data = data, 'ctrl.1', 5, 1/3)
105 # Plot and information of the distribution after data replacement.
   impute_missing_data(data = data, 'ctrl.1', 5, 1/3, TRUE)
   # As we have to chose one of the 6, the plot are adapted especially for the
       first column of data.
    If you want to have good looking plots for any data, switch the two
       commented lines
110 \# in the impute-missing-data function.
   \# impute\_missing\_data(data = data,
                                       'ctrl.3')
   \# impute\_missing\_data(data = data,
   \# impute\_missing\_data(data = data)
                                       'kout.1')
                                       'kout.2')
\# impute\_missing\_data(data = data)
   \# impute\_missing\_data(data = data, `kout.3')
```

(b) Playing with the new mean and new standard deviation

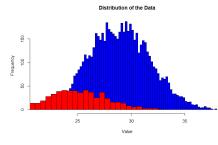
In figure 1 we vary the mean and and standard deviation of the generated data distribution. We can see the generated data distribution sliding to the right when we change the new mean to a higher quartile and getting thinner when we take a smaller fraction of the original SD. The nicest fit is with one third of the original standard deviation and the 5% quartile as the new mean (figure 1b). In figure 2 we show the distribution of the data after having replaced the NAs with the values generated with the same parameters as in figure 1b.



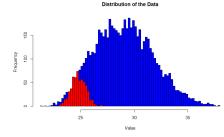
(a) New mean is the 1% quartile of the data distribution and SD is 1/3 of the original SD



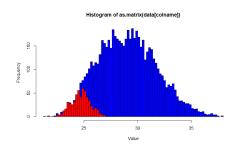
(c) New mean is the 10% quartile of the data distribution and SD is 1/3 of the original SD



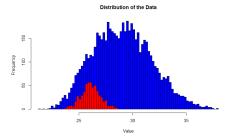
(e) New mean is the 5% quartile of the data distribution and SD is the original SD



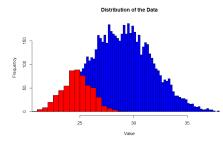
(g) New mean is the 5% quartile of the data distribution and SD is 1/4 of the original SD



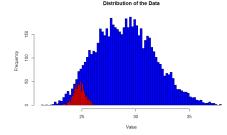
(b) New mean is the 5% quartile of the data distribution and SD is 1/3 of the original SD



(d) New mean is the 15% quartile of the data distribution and SD is 1/3 of the original SD



(f) New mean is the 5% quartile of the data distribution and SD is 1/2 of the original SD



(h) New mean is the 5% quartile of the data distribution and SD is 1/5 of the original SD

Figure 1: Variations of the new mean and the new standard deviation according to the one of the original distribution.

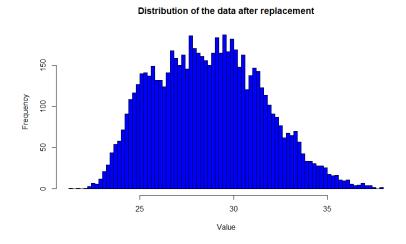


Figure 2: Data distribution after replacement of the NAs with the generated values.

Exercise 7.2: DREAM challenge

(a) NOT IMPLEMENTED