# Aufgabe1

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## Aufgabe 1

**a**)

First a get\_mode function gets defined, which will get wrapped in the impute function.

```
### get_mode function
### This function computes the mode that is defined as the the value that appears most
### often in a set of data values.
### Arguments:
\#\#\# x: A numeric vector. The vector for which the mode should be
### computed.
### na.rm: A logical of length 1. Should NA's be removed when the mode is
### Returns: A numeric of length one: the computed mode
get_mode <- function(x, na.rm= FALSE) {</pre>
 if(na.rm == TRUE){
   x <- na.omit(x)
  unique_values <- unique(x)
   unique_values[which.max(tabulate(match(x, unique_values)))]
}
### impute function
### This function imputes missing values in a dataset. For metric variables the
### median will replace the missing values, while for categorical variables the mode
### will replace the missing values. This function wraps the previously defined
### 'get_mode' function.
###
### Arguments:
### data: A data frame The dataset for which the missing values should get
### imputed.
###
### Returns: A data frame: The imputed dataset.
impute <- function(data){</pre>
 # Input checks
 if(!is.data.frame(data)){
   stop("The Arguments data has to be a data.frame object")
```

```
# Iteration over columns and rows
for (column in colnames(data)){
  # Two cases for imputation
    # Metric variable
    if(is.numeric(data[[column]]) && !(all(na.omit(data[[column]]) %in% 0:1))){
      na index <- which(is.na(data[[column]]))</pre>
      col_median <- median(data[[column]], na.rm = TRUE)</pre>
      data[[column]] <- replace(data[[column]], na_index, col_median)</pre>
    # Categorical variable
    if(any(is.factor(data[[column]]),is.character(data[[column]]),
         is.logical(data[[column]]), all(na.omit(data[[column]]) %in% 0:1))){
    na_index <- which(is.na(data[[column]]))</pre>
    col_mode <- get_mode(data[[column]], na.rm = TRUE)</pre>
    data[[column]] <- replace(data[[column]], na_index, col_mode)</pre>
  }
}
return(data)
```

#### b)

```
### pairwise_correlation function
###
### This function computes the pairwise correlation (Pearson) for all metric
### variables in the data set and crosstables for all combinations
### of categorical variables.
### Pairs of metric and categorical variables are ignored.
### The function returns a list with two elements:
        1) Correlation Matrix of metric variables in the dataset
###
###
        2) A list of crosstables of all categorical variables in the dataset
###
### Arguments:
### data: A data frame The dataset for which the missing values should get
### imputed.
### Returns: A list of two elements: The list contains 1) a correlation matrix
###
                                       all metric variables and 2) a list of
###
                                       crosstables of all categorical variables
###
                                       in the dataset.
pairwise_correlation <- function(data){</pre>
  # Input checks
  if(!is.data.frame(data)){
    stop("The Argument data has to be a data.frame object")
  }
  # Declaration of metric and categorical variables
  metric_cols <- vector()</pre>
  catego_cols <- vector()</pre>
  for(column in 1:ncol(data)){
```

```
# Metric Variables
    if(is.numeric(data[[column]])&& !all(na.omit(data[[column]]) %in% 0:1)){
      metric cols <- c(metric cols,colnames(data[column]))</pre>
    }
  # Categorical Variables
    if(any(is.factor(data[[column]]),
         is.character(data[[column]]),
         is.logical(data[[column]]),
         all(na.omit(data[[column]]) %in% 0:1))){
    catego_cols <- c(catego_cols,names(data[column]))</pre>
  }
  # Subset of metric and
  metric_data <- data[metric_cols]</pre>
  catego_data <- data[catego_cols]</pre>
  # Output element metric variables
    # Correlation calculation for metric variables
  cor_matrix <- as.matrix(cor(data[metric_cols],</pre>
                               use= "complete.obs", method ="pearson"))
  # Output element categorical variables
   # Creating Matrix with all combinations for crosstable
  catego_combn <- combn(colnames(catego_data), 2)</pre>
   # Creating list with crosstables for each combination,
  data_list <- list()</pre>
  for(i in 1:ncol(catego_combn)){
  data_list[[i]] <- prop.table(table(data[[catego_combn[1,i]]]),</pre>
                                       data[[catego_combn[2,i]]]), margin = 2)
  names(data_list)[i] <- paste(catego_combn[1,i], "vs.", catego_combn[2,i])</pre>
  # Output (list) of function
  output <- list("correlation_metric_variables"= cor_matrix,</pre>
                 "crosstables_catego_variables"= data_list)
 return(output)
}
c)
data("patient", package = "pammtools")
str(patient)
## 'data.frame':
                    2000 obs. of 12 variables:
                   : Factor w/ 4 levels "2007", "2008", ...: 1 1 1 1 1 1 1 1 1 1 ...
## $ CombinedicuID: Factor w/ 456 levels "21","24","25",...: 355 355 355 191 112 112 270 270 270 270 ...
## $ CombinedID : int 1110 1111 1116 1316 1410 1414 1507 1509 1518 1520 ...
## $ Survdays
                   : num 30.1 30.1 9.8 30.1 30.1 30.1 30.1 30.1 9 30.1 ...
## $ PatientDied : num 0 0 1 0 0 0 0 1 0 ...
                  : num 30.1 30.1 9.8 30.1 30.1 5.4 30.1 6.4 8 30.1 ...
## $ survhosp
## $ Gender
                   : Factor w/ 2 levels "Female", "Male": 2 1 1 2 2 2 1 2 2 2 ...
                   : int 68 57 68 47 69 47 36 58 36 57 ...
## $ Age
```

```
## $ AdmCatID : Factor w/ 3 levels "Medical", "Surgical Elective",..: 2 1 2 1 2 1 1 1 3 1 ...
## $ ApacheIIScore: int 20 22 25 16 20 21 18 26 14 16 ...
                   : num 31.6 24.3 18 33.7 38.8 ...
## $ BMI
## $ DiagID2
                   : Factor w/ 9 levels "Gastrointestinal",..: 2 8 2 6 8 8 9 4 1 1 ...
# Imputation
patient <- impute(patient)</pre>
# Checking for NA values
na_matrix <- matrix(data = NA, nrow = ncol(patient), ncol= 1)</pre>
for (i in 1:ncol(patient)){
  na_matrix[i,1] <- sum(is.na(patient[i]))</pre>
}
na_matrix
##
         [,1]
    [1,]
##
  [2,]
## [3,]
            0
## [4,]
            0
## [5,]
            0
## [6,]
            0
## [7,]
            0
## [8.]
## [9,]
            0
## [10,]
            0
## [11,]
            0
## [12,]
            0
# Pairwise correlation
patient_no_id <- patient[!(colnames(patient) %in% c("CombinedicuID", "CombinedID"))]</pre>
pairwise_correlation(patient_no_id)
## $correlation_metric_variables
##
                                                    Age ApacheIIScore
                                                                                 BMI
                    Survdays
                                  survhosp
## Survdays
                  1.00000000 0.4864554699 -0.14846658
                                                          -0.21911575 0.0304426441
## survhosp
                  0.48645547 1.0000000000 -0.09531638
                                                          -0.04456446 -0.0002456162
                 -0.14846658 -0.0953163830 1.00000000
                                                           0.24559797 -0.0245339739
## ApacheIIScore -0.21911575 -0.0445644628 0.24559797
                                                         1.00000000 0.0238129354
## BMI
                  0.03044264 -0.0002456162 -0.02453397
                                                           0.02381294 1.0000000000
##
## $crosstables_catego_variables
## $crosstables_catego_variables$`Year vs. PatientDied`
##
##
     2007 0.2246835 0.2666667
##
##
     2008 0.2215190 0.2380952
     2009 0.2500000 0.2142857
     2011 0.3037975 0.2809524
##
##
## $crosstables_catego_variables$`Year vs. Gender`
##
##
             Female
                         Male
##
     2007 0.2358247 0.2320261
##
     2008 0.2177835 0.2295752
     2009 0.2487113 0.2385621
##
```

```
##
    2011 0.2976804 0.2998366
##
## $crosstables_catego_variables$`Year vs. AdmCatID`
##
##
           Medical Surgical Elective Surgical Emergency
##
    2007 0.2155887
                           0.2442244
                                             0.2708758
##
    2008 0.2247098
                           0.2244224
                                             0.2260692
    2009 0.2537313
##
                           0.2145215
                                             0.2321792
##
    2011 0.3059701
                           0.3168317
                                             0.2708758
##
## $crosstables_catego_variables$`Year vs. DiagID2`
##
         Gastrointestinal Cardio-Vascular
##
                                             Other Metabolic Neurologic
##
    2007
                0.2364672 0.3306452 0.1666667 0.2164179
##
    2008
                0.2419929
##
    2009
                0.2028470
                                0.1965812 0.2580645 0.2916667 0.2313433
##
    2011
                0.3167260
                                0.3133903 0.2741935 0.2083333 0.3320896
##
##
         Orthopedic/Trauma
                               Renal Respiratory
                                                    Sepsis
                                      0.2474849 0.1896552
##
    2007
                 0.2564103 0.1304348
##
    2008
                 0.2051282 0.3913043
                                      0.2072435 0.1839080
##
    2009
                 0.2777778 0.1739130
                                      0.2575453 0.3103448
    2011
                 0.2606838 0.3043478 0.2877264 0.3160920
##
##
  $crosstables_catego_variables$`PatientDied vs. Gender`
##
                     Male
##
         Female
##
    0 0.7938144 0.7875817
    1 0.2061856 0.2124183
##
##
## $crosstables_catego_variables$`PatientDied vs. AdmCatID`
##
        Medical Surgical Elective Surgical Emergency
##
##
    0 0.7470978
                        0.8679868
                                          0.8472505
    1 0.2529022
                        0.1320132
                                           0.1527495
##
##
## $crosstables_catego_variables$`PatientDied vs. DiagID2`
##
##
      Gastrointestinal Cardio-Vascular
                                         Other Metabolic Neurologic
             0.8256228
                             0.7635328 0.7903226 0.8750000 0.7873134
##
    0
                             0.2364672 0.2096774 0.1250000 0.2126866
##
             0.1743772
##
##
      Orthopedic/Trauma
                            Renal Respiratory
                                                 Sepsis
##
              0.8888889 0.7391304
                                  0.7625755 0.7183908
##
              0.1111111 0.2608696
                                   0.2374245 0.2816092
##
  $crosstables_catego_variables$`Gender vs. AdmCatID`
##
##
             Medical Surgical Elective Surgical Emergency
    Female 0.3963516
                            0.4224422
                                                0.3462322
##
    Male 0.6036484
                             0.5775578
##
                                                0.6537678
##
## $crosstables_catego_variables$`Gender vs. DiagID2`
##
```

```
##
            Gastrointestinal Cardio-Vascular
                                                  Other Metabolic Neurologic
##
                   0.3736655
                                    0.3646724 0.4112903 0.4166667
                                                                   0.4216418
     Female
                                    0.6353276 0.5887097 0.5833333 0.5783582
##
     Male
                   0.6263345
##
##
            Orthopedic/Trauma
                                   Renal Respiratory
                                                         Sepsis
                    0.2435897 0.4347826
                                           0.4426559 0.4137931
##
     Female
     Male
                    0.7564103 0.5652174
                                           0.5573441 0.5862069
##
##
##
   $crosstables_catego_variables$`AdmCatID vs. DiagID2`
##
##
                        Gastrointestinal Cardio-Vascular
                                                                Other Metabolic
                                               0.48148148 0.45967742 1.00000000
##
     Medical
                               0.22419929
                                               0.36467236 0.25000000 0.00000000
##
     Surgical Elective
                               0.25266904
                                               0.15384615 0.29032258 0.00000000
##
     Surgical Emergency
                               0.52313167
##
##
                        Neurologic Orthopedic/Trauma
                                                            Renal Respiratory
##
     Medical
                        0.56716418
                                           0.31196581 0.52173913
                                                                   0.92152918
##
     Surgical Elective 0.11194030
                                           0.02564103 0.34782609
                                                                   0.05835010
##
     Surgical Emergency 0.32089552
                                           0.66239316 0.13043478 0.02012072
##
##
                            Sepsis
##
     Medical
                        1.0000000
##
     Surgical Elective 0.00000000
##
     Surgical Emergency 0.00000000
```

d)

##

1

The correlation between the age of the patients (age) and the time until they were released from the hospital (survhosp) almost no correlation. The Pearson correlation coefficient is with -0.0953 slightly negative.

Taking a look at the crosstable of the diagnose (DiagID2) and the categorical variable if or if no the patient survived (PatientDied). One can see that (relatively) always more people survived than died for each diagnosis. The highest probability to survive is for an Orthopedic and Metabolic emergencies, the lowest probability to survive is for a Sepsis diagnose.

```
# Age and survhosp
pairwise_correlation(patient_no_id)[[1]][3,2, drop= FALSE]
##
          survhosp
## Age -0.09531638
# DiagID2 and PatientDied
pairwise_correlation(patient_no_id)[[2]][7]
## $`PatientDied vs. DiagID2`
##
##
       Gastrointestinal Cardio-Vascular
                                             Other Metabolic Neurologic
                               0.7635328 0.7903226 0.8750000
##
     0
              0.8256228
                                                              0.7873134
##
     1
              0.1743772
                               0.2364672 0.2096774 0.1250000
                                                               0.2126866
##
##
       Orthopedic/Trauma
                              Renal Respiratory
                                                   Sepsis
               0.8888889 0.7391304
                                      0.7625755 0.7183908
##
     0
```

0.2374245 0.2816092

0.1111111 0.2608696

### **Session Info**

```
sessionInfo()
## R version 4.0.2 (2020-06-22)
## Platform: x86_64-apple-darwin17.0 (64-bit)
## Running under: macOS 10.16
##
## Matrix products: default
           /Library/Frameworks/R.framework/Versions/4.0/Resources/lib/libRblas.dylib
## LAPACK: /Library/Frameworks/R.framework/Versions/4.0/Resources/lib/libRlapack.dylib
##
## locale:
## [1] de_DE.UTF-8/de_DE.UTF-8/de_DE.UTF-8/C/de_DE.UTF-8/de_DE.UTF-8
## attached base packages:
## [1] stats
                 graphics grDevices utils
                                                datasets methods
                                                                    base
##
## other attached packages:
## [1] forcats_0.5.1
                        stringr_1.4.0
                                         dplyr_1.0.7
                                                         purrr_0.3.4
                        tidyr_1.1.3
##
   [5] readr_1.4.0
                                         tibble_3.1.1
                                                         ggplot2_3.3.3
  [9] tidyverse_1.3.1 pammtools_0.5.7
##
## loaded via a namespace (and not attached):
## [1] Rcpp_1.0.6
                            lubridate_1.7.10
                                                 mvtnorm_1.1-1
## [4] lattice_0.20-44
                            assertthat_0.2.1
                                                 digest_0.6.27
## [7] foreach_1.5.1
                            utf8_1.2.1
                                                 R6_2.5.0
## [10] cellranger_1.1.0
                            backports_1.2.1
                                                 reprex_2.0.0
## [13] evaluate_0.14
                            httr_1.4.2
                                                 pillar_1.6.0
## [16] rlang_0.4.11
                                                 readxl 1.3.1
                            lazyeval 0.2.2
## [19] rstudioapi_0.13
                            Matrix_1.3-3
                                                 checkmate_2.0.0
## [22] rmarkdown_2.8
                            splines_4.0.2
                                                 munsell_0.5.0
## [25] broom_0.7.6
                                                 numDeriv_2016.8-1.1
                            compiler_4.0.2
## [28] modelr_0.1.8
                            xfun_0.22
                                                 pkgconfig_2.0.3
## [31] mgcv 1.8-35
                            htmltools 0.5.1.1
                                                 tidyselect 1.1.1
## [34] prodlim_2019.11.13
                                                 fansi_0.4.2
                            codetools_0.2-18
## [37] withr_2.4.2
                            crayon_1.4.1
                                                 dbplyr_2.1.1
## [40] timereg_2.0.0
                            grid_4.0.2
                                                 nlme_3.1-152
## [43] jsonlite_1.7.2
                            gtable_0.3.0
                                                 lifecycle_1.0.0
## [46] DBI_1.1.1
                            magrittr_2.0.1
                                                 scales_1.1.1
## [49] cli_2.5.0
                            stringi_1.6.1
                                                 fs_{1.5.0}
## [52] xml2_1.3.2
                            ellipsis_0.3.2
                                                 generics_0.1.0
## [55] vctrs_0.3.8
                            Formula_1.2-4
                                                 lava_1.6.9
## [58] iterators_1.0.13
                            tools_4.0.2
                                                 glue_1.4.2
## [61] hms_1.0.0
                            pec_2020.11.17
                                                 survival_3.2-11
                            colorspace_2.0-1
                                                 rvest_1.0.0
## [64] yaml_2.2.1
## [67] knitr 1.33
                            haven 2.4.1
```