

# Processing

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## Imports

```
library(tidyverse)
library(readxl)
library(yaml)
library(lubridate)
```

## Loading files

```
df <- read_csv("../dataset/BD Marcapasso_11jul22.csv", show_col_types = FALSE) %>%
  select(-record_id)
df_names <- read_excel("../dataset/Dicionario_dados_BD Marcapasso_11jul22.xlsx")
```

## Fixing data dictionary

```
names(df_names) <- make.names(names(df_names), unique=TRUE) %>% tolower

df_names <- df_names %>%
  mutate(variable.name = case_when(variable.name == 'icu_days_t0' ~ 'icu_t0',
                                    variable.name == 'dialysis_days_t0' ~ 'dialysis_t0',
                                    TRUE ~ variable.name), # mismatch with dataset column name
         variable.name = str_replace(variable.name, "\\[", ""),
         variable.name = str_replace(variable.name, "\\]", ""),
         field.label = str_replace(field.label, "\\+", "com"),
         field.label = str_replace_all(field.label, "_", " "), # add spaces
         field.label = str_replace_all(field.label, "[\\r\\n]", ""),
         abbrev.field.label = str_replace(field.label, " \\s*\\\[^\"]+\\]", ""),
         abbrev.field.label = if_else(variable.name %in% c('admission_posop_count',
                                                         'admission_pre_t0_count'),
                                     str_replace(field.label, "de episódios", ""),
                                     abbrev.field.label)) %>%
  rename(momento.aquisicao = momento.da.aquisição.do.dado..admissão.t0.ou.pós.t0.)
```

## Separating columns by type

```
outcome_columns <- df_names %>%
  filter(str_detect(momento.aquisicao, 'Desfecho')) %>%
  .$variable.name

categorical_columns <- df_names %>%
  filter(stringr::str_detect(options..definition, '\\|')) %>%
  .$variable.name %>%
  setdiff(outcome_columns)

date_columns <- df_names %>%
```

```

filter(options..definition == 'data') %>%
  .$variable.name

location_columns <- c('zipcode', 'patient_city')

other_columns <- c('record_id')

numerical_columns <- setdiff(names(df),
                             c(categorical_columns, date_columns,
                               location_columns, other_columns))

df[categorical_columns] <- lapply(df[categorical_columns],
                                 as.character)
df[outcome_columns] <- lapply(df[outcome_columns],
                              as.numeric)
df[date_columns] <- lapply(df[date_columns],
                          ymd)

columns_list <- list('categorical_columns' = categorical_columns,
                    'numerical_columns' = numerical_columns,
                    'date_columns' = date_columns,
                    'location_columns' = location_columns,
                    'outcome_columns' = outcome_columns)

con <- file('./auxiliar/columns_list.yaml', "w")
write_yaml(columns_list, con)
close(con)

```

## Filling missing values on death outcomes

```

df <- df %>%
  tidyr::replace_na(list(death_30days = 0,
                        death_180days = 0,
                        death_1year = 0,
                        death_2year = 0,
                        death_3year = 0))

# df %>%
#   select(death_30days, death_180days, death_1year, death_2year, death_3year) %>%
#   mutate(death_sum = death_30days + death_180days + death_1year + death_2year + death_3year) %>%
#   .$death_sum %>%
#   unique

```

## Filtering eligible patients

```

df <- df %>%
  filter(disch_outcomes_t0 == 0)

df %>% dim

## [1] 15766 239

```

## Recalculating outcome columns for modeling

```

df <- df %>%
  mutate(readmission_1year = readmission_30d + readmission_60d + readmission_180d + readmission_1year,
         readmission_180d = readmission_30d + readmission_60d + readmission_180d,

```

```
readmission_60d = readmission_30d + readmission_60d,  
death_3year = death_30days + death_180days + death_1year + death_2year + death_3year,  
death_2year = death_30days + death_180days + death_1year + death_2year,  
death_1year = death_30days + death_180days + death_1year,  
death_180days = death_30days + death_180days)
```

## Saving processed data

```
saveRDS(df, "../dataset/processed_data.rds")  
saveRDS(df_names, "../dataset/processed_dictionary.rds")  
  
save(df, file = "../dataset/processed_data.RData")  
save(df_names, file = "../dataset/processed_dictionary.RData")
```