Breast Cancer Prediction

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```
library(tidyverse)
## -- Attaching packages ----- tidyverse 1.2.1 --
## v ggplot2 3.2.1
                         v purrr
                                   0.3.2
## v tibble 2.1.3
                         v dplyr
                                   0.8.0.1
## v tidyr
           0.8.3
                        v stringr 1.4.0
## v readr
            1.3.1
                         v forcats 0.4.0
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                     masks stats::lag()
library(ggplot2)
library(magrittr)
##
## Attaching package: 'magrittr'
## The following object is masked from 'package:purrr':
##
##
       set_names
## The following object is masked from 'package:tidyr':
##
##
       extract
library(dplyr)
Import data
data <- read.csv("../data/data.csv")</pre>
dim(data)
## [1] 569 33
print(colnames(data))
  [1] "id"
                                  "diagnosis"
  [3] "radius_mean"
                                  "texture_mean"
## [5] "perimeter_mean"
                                  "area_mean"
## [7] "smoothness_mean"
                                  "compactness_mean"
## [9] "concavity_mean"
                                  "concave.points_mean"
## [11] "symmetry_mean"
                                  "fractal_dimension_mean"
## [13] "radius_se"
                                  "texture_se"
## [15] "perimeter_se"
                                  "area_se"
## [17] "smoothness_se"
                                  "compactness_se"
## [19] "concavity_se"
                                  "concave.points_se"
```

```
## [21] "symmetry_se"
                               "fractal_dimension_se"
## [23] "radius_worst"
                               "texture_worst"
## [25] "perimeter_worst"
                               "area worst"
## [27] "smoothness_worst"
                               "compactness_worst"
## [29] "concavity_worst"
                               "concave.points_worst"
## [31] "symmetry_worst"
                               "fractal_dimension_worst"
## [33] "X"
Check for NAs
data %>%
summarise_all(funs(sum(is.na(.))))
## Warning: funs() is soft deprecated as of dplyr 0.8.0
## please use list() instead
##
## # Before:
## funs(name = f(.)
##
## # After:
## list(name = \sim f(.))
## This warning is displayed once per session.
    id diagnosis radius_mean texture_mean perimeter_mean area_mean
                  0
                                         0
## 1 0 0
                             0
  smoothness_mean compactness_mean concavity_mean concave.points_mean
## symmetry_mean fractal_dimension_mean radius_se texture_se perimeter_se
                                   0 0 0
## 1
## area_se smoothness_se compactness_se concavity_se concave.points_se
              0
## 1 0
                                   0
                                              0
## symmetry_se fractal_dimension_se radius_worst texture_worst perimeter_worst
                                0
                                   0
## area_worst smoothness_worst compactness_worst concavity_worst
                          0
## concave.points_worst symmetry_worst fractal_dimension_worst X
Remove id and X
data %<>%
dplyr::select(-c(id, X))
```