

Project Report G9

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1 Introduction

2 Methods

2.1 Analysis protocol

2.2 Data cleaning and preparation

```
filename <- "armpit.txt"
armpit <- read.table(filename)
summary(armpit)
```

```
## Corynebacterium.1 Corynebacterium.2 Corynebacterium.3 Corynebacterium.4
## Min. : 0.0000 Min. : 0.0000 Min. : 0.0000 Min. : 0.0000
## 1st Qu.: 0.0000 1st Qu.: 0.0000 1st Qu.: 0.0000 1st Qu.: 0.0000
## Median : 0.2019 Median : 0.3458 Median : 0.0000 Median : 0.0000
## Mean : 6.5259 Mean : 11.4825 Mean : 1.9110 Mean : 1.2008
## 3rd Qu.: 1.6563 3rd Qu.: 2.7544 3rd Qu.: 0.7368 3rd Qu.: 0.2539
## Max. : 96.1716 Max. : 87.1904 Max. : 30.2198 Max. : 18.3894
##
## Staphylococcus.1 Staphylococcus.2 Staphylococcus.3 Staphylococcus.4
## Min. : 1.203 Min. : 0.0000 Min. : 0.0000 Min. : 0.0000
## 1st Qu.: 42.155 1st Qu.: 0.0000 1st Qu.: 0.0000 1st Qu.: 0.0000
## Median : 93.504 Median : 0.0000 Median : 0.0000 Median : 0.0000
## Mean : 71.173 Mean : 6.2755 Mean : 0.4602 Mean : 0.97152
## 3rd Qu.: 98.961 3rd Qu.: 0.2506 3rd Qu.: 0.0801 3rd Qu.: 0.07669
## Max. : 100.000 Max. : 89.7331 Max. : 6.6667 Max. : 21.56593
```

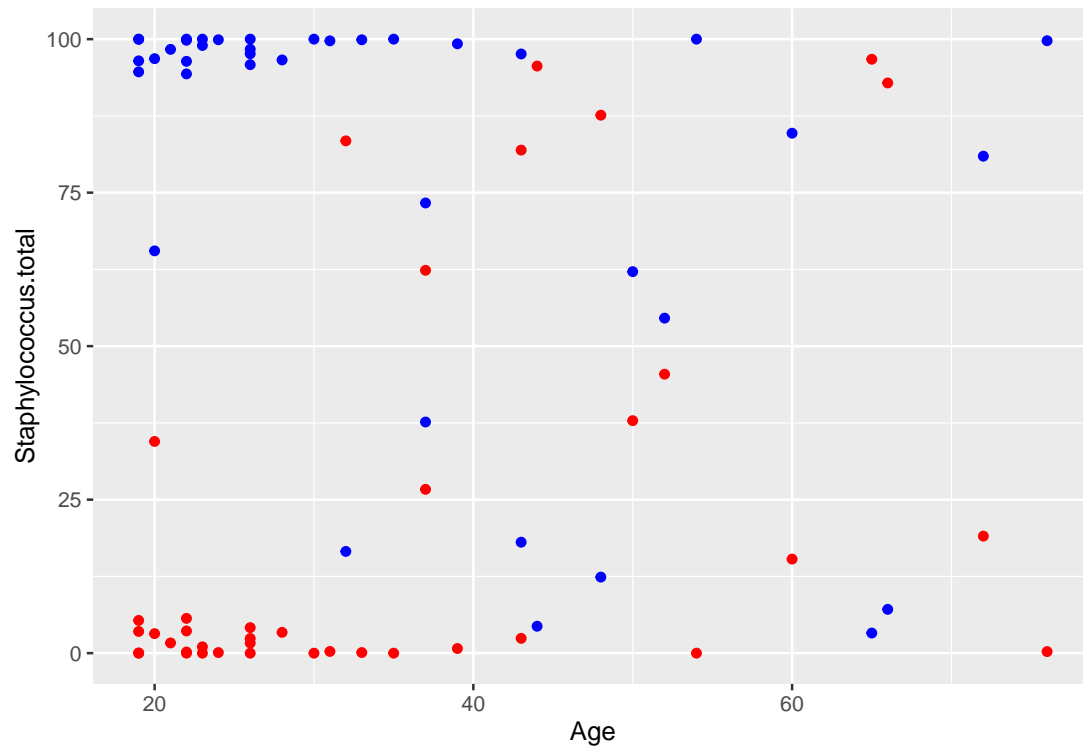
```
##
##      Age      BMI      Gender
## Min.   :19.00   Min.   :0.000   : 1
## 1st Qu.:22.00   1st Qu.:0.000   F :23
## Median :30.00   Median :0.000   F : 1
## Mean   :35.23   Mean   :0.325   M :14
## 3rd Qu.:43.50   3rd Qu.:1.000   M : 1
## Max.   :76.00   Max.   :1.000
## NA's    :1

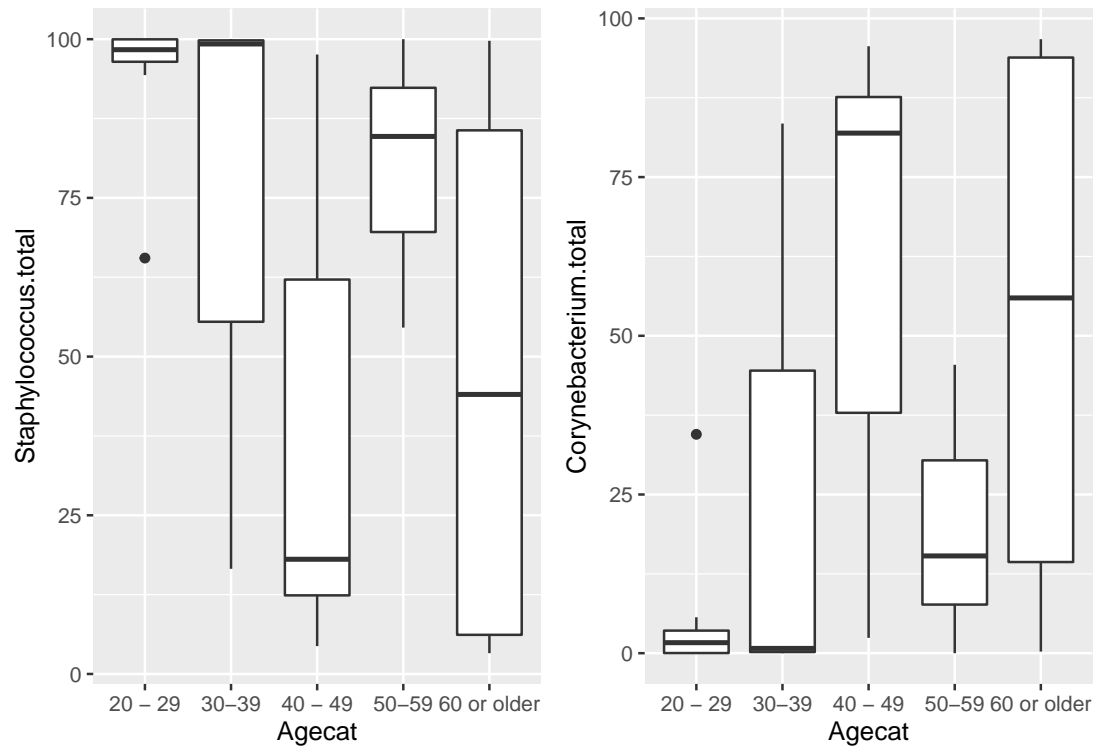
# Step 1: assure we have correct gender labels
# convert the gender 'F ' to 'F' and ' M' to 'M'
armpit[grepl('F', armpit$Gender), 'Gender'] <- 'F'
armpit[grepl('M', armpit$Gender), 'Gender'] <- 'M'
# drop the observation where gender and age were not filled out --> TODO : is this acceptable?
armpit <- armpit[armpit$Gender != '',]
armpit$Gender <- factor(armpit$Gender)
armpit <- armpit[!is.na(armpit$Age),]
summary(armpit)

## Corynebacterium.1 Corynebacterium.2 Corynebacterium.3 Corynebacterium.4
## Min.   : 0.0000   Min.   : 0.0000   Min.   : 0.000   Min.   : 0.0000
## 1st Qu.: 0.0000   1st Qu.: 0.0000   1st Qu.: 0.000   1st Qu.: 0.0000
## Median : 0.1968   Median : 0.2011   Median : 0.000   Median : 0.0000
## Mean   : 6.6481   Mean   :11.7397   Mean   : 1.396   Mean   : 1.2183
## 3rd Qu.: 1.6086   3rd Qu.: 3.0679   3rd Qu.: 0.540   3rd Qu.: 0.1478
## Max.   :96.1716   Max.   :87.1904   Max.   :30.220   Max.   :18.3894
## Staphylococcus.1 Staphylococcus.2 Staphylococcus.3 Staphylococcus.4
## Min.   : 1.203   Min.   : 0.0000   Min.   :0.00000   Min.   : 0.00000
## 1st Qu.: 40.355   1st Qu.: 0.0000   1st Qu.:0.00000   1st Qu.: 0.00000
## Median : 93.675   Median : 0.0000   Median :0.00000   Median : 0.00000
## Mean   : 71.213   Mean   : 6.3247   Mean   :0.46404   Mean   : 0.99643
## 3rd Qu.: 99.022   3rd Qu.: 0.1605   3rd Qu.:0.06568   3rd Qu.: 0.08635
## Max.   :100.000   Max.   :89.7331   Max.   :6.66667   Max.   :21.56593
##      Age      BMI      Gender
## Min.   :19.00   Min.   :0.0000   F:24
## 1st Qu.:22.00   1st Qu.:0.0000   M:15
## Median :30.00   Median :0.0000
## Mean   :35.23   Mean   :0.3333
## 3rd Qu.:43.50   3rd Qu.:1.0000
## Max.   :76.00   Max.   :1.0000
```

3 Results

3.1 Correlation between bacteria and age





4 Conclusion