BDA - Assignment 1

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<pre># To install aaltobda, see the General information in the assignment. remotes::install_github("avehtari/BDA_course_Aalto", subdir = "rpackage", upgrade")</pre>	= "never")
## Skipping install of 'aaltobda' from a github remote, the SHA1 (38f34d35) has no	ot changed since last

Exercise 1 (Basic probability theory notation and terms):

- **Probability:** Based on *known* parameters, how likely an specific event occurs.
- Probability mass: For discrete variables, how likely a series of events (could be a single event) occurs.
- **Probability density** For continuous variables, how likely an interval of events (probability at a single point is zero) occurs.
- **Probability mass function (pmf):** Related to *Probability mass*, it is the function that maps the event to the probability that it occurs.
- **Probability density function (pdf):** Related to *Probability density*, it is the function that maps the interval of events to the probability that it occurs.
- Probability distribution: The mass/density assigned by the pmf/pdf to each outcome occurrence.
- Discrete probability distribution: The mass assigned by the pmf to each outcome occurrence.
- Continuous probability distribution: The density assigned by the pdf to each outcome occurrence.
- Cumulative distribution function: The summed of the mass/density from all outcomes occurrence as they are arranged from left to right.
- Likelihood: The probability of observing the data given the parameters.

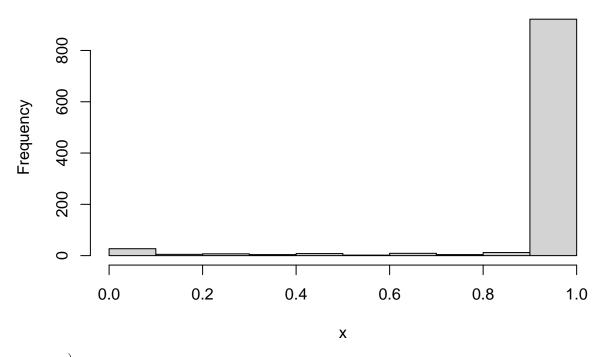
Exercise 2 (Basic computer skills):

• a)

```
p <- seq(0, 1, length = 100)</pre>
mu <- 0.2
sd <- .01
ps <- dbeta(p, mu, sd)
plot(p, ps, type = "l")</pre>
      0.8
      9.0
bs
      0.4
      0.0
              0.0
                               0.2
                                                                 0.6
                                                                                  8.0
                                                                                                   1.0
                                                0.4
                                                          p
                                                                                                          - b)
n <- 1000
x <- rbeta(n, mu, sd)
```

hist(x)

Histogram of x



```
c)
cat("This is the mean:", mean(x))
## This is the mean: 0.9492692
cat("\n")
cat("This is the variance:", var(x))
## This is the variance: 0.03726147
quantile(x, prob = seq(0.1, 1, by = .05), CI.type = "two.sided")
                                                                        40%
##
         10%
                   15%
                              20%
                                        25%
                                                   30%
                                                             35%
                                                                                  45%
## 0.9937794 0.9999824 0.9999999 1.0000000 1.0000000 1.0000000 1.0000000 1.0000000
                                        65%
                                                             75%
                                                                        80%
##
         50%
                   55%
                              60%
                                                   70%
                                                                                  85%
## 1.0000000 1.0000000 1.0000000 1.0000000 1.0000000 1.0000000 1.0000000 1.0000000
         90%
                   95%
                             100%
##
## 1.0000000 1.0000000 1.0000000
```

Exercise 3 (Bayes' theorem):

According from my notes for the exercise required in the pre-requisites, I found that P(cancer does not get detected) = P(cancer | negative result) = 0.000002, and P(positive test but no cancer) = P(cancer | positive result) = 0.97. My conclusion was that while P(C|N) is low, I am concerned about P(C|P) being close to 1. I would pass on investing on this company.

Exercise 4 (Bayes' theorem):

```
boxes \leftarrow matrix(c(2,4,1,5,1,3),
                 ncol = 2,
                 dimnames = list(c("A", "B", "C"),
                                   c("red", "white")))
cat("The following is the boxes matrix")
## The following is the boxes matrix
boxes
##
     red white
## A
       2
## B
       4
              1
## C
       1
              3
p_red <- function(boxes) {</pre>
  Pr_boxes_balls <- boxes/rowSums(boxes)</pre>
  P_R_box <- matrix(Pr_boxes_balls[,1], ncol = 1)</pre>
  Pr_boxes \leftarrow matrix(c(.4,.1,.5), ncol = 3)
  p_red <- Pr_boxes %*% P_R_box</pre>
  return(p_red)
pred <- p_red(boxes)</pre>
cat("The probability of a red ball is P(R):", pred)
## The probability of a red ball is P(R): 0.3192857
p_box <- function(boxes) {</pre>
  Pr_boxes_balls <- boxes/rowSums(boxes)</pre>
  P_R_box <- matrix(Pr_boxes_balls[,1], ncol = 1)</pre>
  Pr_boxes \leftarrow matrix(c(.4,.1,.5), ncol = 3)
  P_box_R_list <- list()</pre>
  for(i in 1:3) {
    P_box_R <- (P_R_box[i]*Pr_boxes[i])/pred</pre>
    P_box_R_list[i] <- P_box_R</pre>
  return(P_box_R_list)
}
cat("These are the probabilities of each box being picked given grabbing a red ball")
## These are the probabilities of each box being picked given grabbing a red ball
p_box(boxes)
## [[1]]
## [1] 0.3579418
## [[2]]
## [1] 0.2505593
##
## [[3]]
## [1] 0.3914989
```

```
cat("\n")
cat("The most likely box to be picked if we observed a red ball is c")
```

The most likely box to be picked if we observed a red ball is c

Exercise (Bayes' theorem):

First thing to notice is that probabilities here dependent on Elvis and the other facts that we know. That is P(B|I, Elvis) = 1 because we know that Elvis was male. For sake of exposition, I dropped the Elvis from the notation.

Our bayes theorem formula is:

$$P(I|B) = \frac{P(B|I)P(I)}{P(B)} = \frac{P(B|I)P(I)}{P(B|I)P(I) + P(B|F)P(F)}$$

where I stands for identical twin, B is for fraternal. Let us assume the following (no strict) assumption holds:

$$P(B|F) = \frac{P(B,F)}{P(F)} = \frac{P(B)P(F)}{P(F)} = P(B)$$

The latter simply assumes that the probability of Elvis having a fraternal brother is idependent of the probability of the brother being a boy.

Therefore, P(I|B) is equal to:

```
fraternal_prob <- 1/150
identical_prob <- 1/400
p_identifical_twin <- function() {
  identical_prob/(identical_prob + .5*fraternal_prob)
}
p_identifical_twin()</pre>
```

[1] 0.4285714