流行病學與生物統計計算

Homework 4

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Homework4

#Q1: Calculate factorial(10), using while() and for() respectively

```
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#Q1 : Calculate factorial(10), using while() and for() respectively
   # while(){}
      fa <- function(x) {</pre>
          if (x != round(x) || x < 0) {
             x <- "error, x should be nature number or 0"
          }else if (x == 0 \mid | x == 1)  {
             x <- 1
          }else {
             i < -x - 1
             while (i != 1) {
                   x \leftarrow x * i
                    i <- i - 1
                 }
          }
          return(x)
      }
      fa(10) # 10! = 3628800
```

```
# for(){}
fac <- function(x) {
    if (x != round(x) || x < 0) {
        x <- "error, x should be nature number or 0"
    }else if (x == 0 || x == 1) {
        x <- 1
    }else {
        for (i in 1:(x - 1)) {
            x <- x * i
        }
    }
    return(x)
}</pre>
```

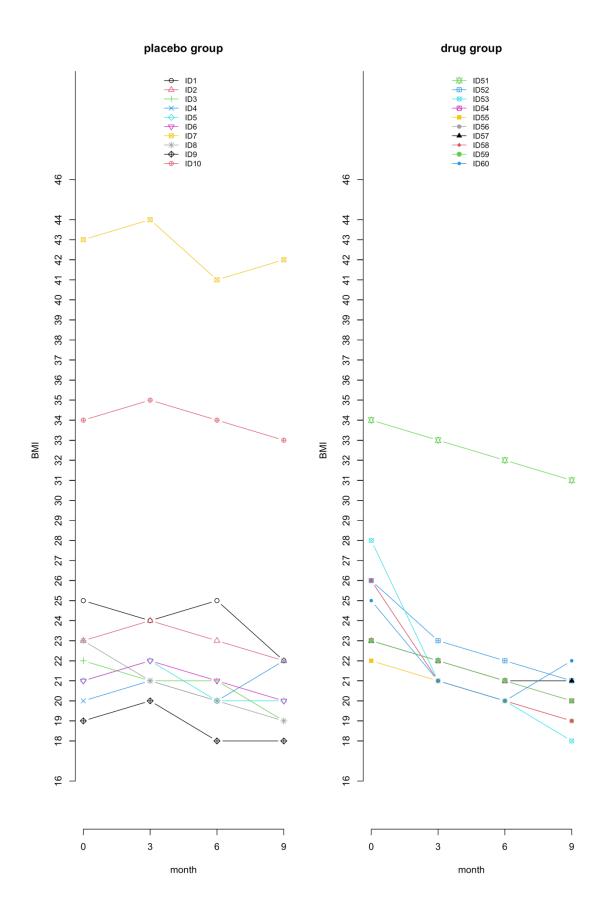
Q2 use while(){} and cat() to find missing value

```
# Q2 use while(){} and cat() to find missing value
   find_na <- function(x) {</pre>
       done <- FALSE
       i <- 1
       while (!done) {
          if (is.na(x[i] == 1)) {
              done <- TRUE
          }else if (i == (length(x))) {
              i <- "NULL"
              done <- TRUE
          }else {
              i <- i + 1
          }
       return(cat(i, "\n"))
   }
   x \leftarrow c(3600, 5000, 12000, NA, 1000, 2000, 600, 7500, 1800, 9000)
   find_na(x)
```

Q2 terminal

Q3 : BMI curve ID51 - ID60

```
# Q3 : BMI curve ID51 - ID60
   BMI <- read.csv("/Users/raymond/Desktop/R/course/data/BMIrepeated.csv")</pre>
   x \leftarrow seq(0, 9, by = 3)
   y <- cbind(BMI$BMI0, BMI$BMI1, BMI$BMI2, BMI$BMI3)
   png(filename = "hw4.3.png", width = 2000, height = 3000, res = 200)
   par(mfrow = c(1, 2))
   plot(x, y[1, ], type = "b",
          lwd = 1, col = 1, lty = 1, pch = 1,
          ylim = c(15, 50),
          axes = FALSE,
          xlab = "month", ylab = "BMI", main = "placebo group")
   for (i in 2:10) {
      lines(x, y[i, ], lty = 1, lwd = 1, col = i, type = "b", pch = i)
   axis(1, at = x, labels = seq(0, 9, by = 3))
   axis(2, at = y)
   legend("top", bty = "n",
   c("ID1", "ID2", "ID3", "ID4", "ID5", "ID6", "ID7", "ID8", "ID9", "ID10"),
   lty = 1, col = (1:10), lwd = 1, pch = (1:10), cex = 0.8, pt.cex = 1.0)
   plot(x, y[51, ], type = "b",
          lwd = 1, col = 11, lty = 1, pch = 11,
          ylim = c(15, 50),
          axes = FALSE,
          xlab = "month", ylab = "BMI", main = "drug group")
   for (k in 52:60) {
      lines(x, y[k, ], lty = 1, lwd = 1, col = k - 40, type = "b", pch = k-40)
   }
   axis(1, at = x, labels = seq(0, 9, by = 3))
   axis(2, at = y)
   legend("top", bty = "n",
   c("ID51", "ID52", "ID53", "ID54", "ID55", "ID56", "ID57", "ID58", "ID59",
"ID60"),
   lty = 1, col = (11:20), lwd = 1, pch = (11:20), cex = 0.8, pt.cex = 1.0)
   dev.off()
```



Q4: 3D pie chart with percentage

```
# Q4 : 3D PIE with percentage
    library(plotrix)

par(mfrow = c(1, 1))
subject <- c(10, 12, 4, 16, 8)
lbls <- c("US", "UK", "Australia", "Germany", "France")
percent <- round((subject / sum(subject)) * 100)
lbls <- paste(lbls, percent)
lbls <- paste(lbls, "%", sep = "")
png(filename = "hw4.4.png", width = 3000, height = 2000, res = 200)
pie3D(subject, labels = lbls, explode = 0.1, col = rainbow(length(lbls))
    , main = "pie chart of countries")
dev.off()</pre>
```

pie chart of countries

