Practical Exercises for Day 3

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Exercise 8

Apply the summary statistics to the perulung_ems and ToothGrowth data set.

```
# Read in .csv data
lung <- read.csv("C:\\Users\\Exercises\\data\\perulung_ems.csv", sep = ";")</pre>
head(lung)
str(lung)
summary(lung)
lung$sex <- factor(lung$sex, levels = c("0", "1"))</pre>
levels(lung$sex) <- c("female", "male")</pre>
lung$respsymptoms <- factor(lung$respsymptoms, levels = c("0", "1"))</pre>
# Continuous and factor
tapply(lung$height, lung$sex, mean)
tapply(lung$height, lung$respsymptoms, mean)
# Factor and factor
table(lung$respsymptoms, lung$sex)
prop.table(table(lung$respsymptoms, lung$sex))
# Continuous and factor
tapply(lung$age, lung$sex, mean)
tapply(lung$age, lung$respsymptoms, mean)
# Continuous and factor
tapply(lung$fev1, lung$sex, mean)
tapply(lung$fev1, lung$respsymptoms, mean)
# Continuous and continuous
pairs(lung)
cor.test(lung$fev1, lung$age, method = "pearson")
cor.test(lung$fev1, lung$height, method = "pearson")
# ToothGrowth
```

```
summary(ToothGrowth)
table(ToothGrowth$supp)
tapply(ToothGrowth$len, ToothGrowth$supp, mean)
tapply(ToothGrowth$len, ToothGrowth$supp, median)
tapply(ToothGrowth$len, ToothGrowth$supp, sd)
table(ToothGrowth$dose)
tapply(ToothGrowth$len, ToothGrowth$dose, mean)
tapply(ToothGrowth$len, ToothGrowth$dose, median)
tapply(ToothGrowth$len, ToothGrowth$dose, sd)
```

Exercise 9A: Plausibility Checks

- What can go wrong?
- Identify different strategies for spotting these potential errors.
 - Logical errors
 - Spelling mistakes
- Import the data set bacteria_plausibility_check.csv to R.

• Detect the **six** errors in the imported data set bacteria_plausibility_check.csv in R.

```
str(bac)
table(bac$y) # We have wrong factor levels: 0, 1
table(bac$ap)
table(bac$hilo) # We have a spelling mistake: Hi.
table(bac$week) # There's only ONE observation in week 20.
table(bac$ID)
table(bac$trt) # We have wrong factor levels: drug++, penicillin+
summary(bac$child_weight) # child weight of 302.8 kg is impossible --> comma
```

• Find possible solutions in R how to handle these challenges.

```
bac$y[which(bac$y == 0] <- "n"

# bac$y[bac$y == 0] <- "y"

# Delete the unused levels with the function droplevels(...)

bac$y <- droplevels(bac$y)

bac$hilo[bac$hilo == "Hi"] <- "hi"

bac$hilo[which(bac$hilo == "Hi")] <- "hi"

levels(bac$hilo) <- c("hi", "hi", "lo")

summary(bac)

bac <- bac[-which(bac$week == 20), ] # dim(bac)

bac$trt[bac$trt == "drug++"] <- "drug+"

bac$trt[bac$trt == "penicillin+"] <- "drug+"

table(bac$trt) # We have wrong factor levels: drug++, penicillin+

bac$child_weight[bac$child_weight == 302.8] <- 30.28

summary(bac)</pre>
```

• Do all variables have the correct data type (numeric, integer, factor)? - If not, do correct / define them.

```
bac$y <- factor(bac$y, levels = c("n", "y"))
bac$hilo[bac$hilo == "Hi"] <- "hi"
bac$ID <- factor(bac$ID)
bac$trt <- factor(bac$trt)</pre>
```

Exercise 9B: Missing Values

Check out the difference between the different missing values

```
y1 <- c(2, 4, 3, NA, 6, 1)
y2 <- c("diseased", "healthy", NA, "NA")
y3 <- c(1, "NA", 0, 1, NaN)

is.na(y1)
which(is.na(y1))
is.na(y2)
which(is.na(y2))
is.na(y3)
which(is.na(y3))
is.nan(y3)</pre>
```

• Create a vector with missing values and determine the mean and median

```
myvector <- c(1:3,NA,NA,1:3)
mean(myvector)
mean(myvector,na.rm=TRUE) # calculates c(1, 2, 3, 1, 2, 3)
median(myvector,na.rm=TRUE)</pre>
```

• If x = c (22,3,7,NA,NA,67) what will be the output for the R statement length(x)?

```
x <- c (22,3,7,NA,NA,67)

length(x)
```

• If x = c(NA,3,14,NA,33,17,NA,41) which line of R code removes all occurrences of NA in x.

```
x <- c(NA,3,14,NA,33,17,NA,41)
x[!is.na(x)]
x[is.na(x)]
x[which(is.na(x))] <- 0</pre>
```

• If y = c(1,3,12,NA,33,7,NA,21) what R statement will replace all occurrences of NA with 11?

```
y <- c(1,3,12,NA,33,7,NA,21)
y[y=="NA"] <- 11
y[is.na(y)] <- 11
y[y==11] <- NA
```

• If x = c(34,33,65,37,89,NA,43,NA,11,NA,23,NA) then what will count the number of occurrences of NA in x?

```
x <- c(34,33,65,37,89,NA,43,NA,11,NA,23,NA)
sum(x=="NA")
sum(x == "NA", is.na(x))
sum(is.na(x))</pre>
```

• Create a vector and find the number of missing values and their position

```
x1 <- c(rnorm(10,5,2),NA,5:12,NA,6,7.5,NA)
is.na(x1)
summary(x1)
sum(is.na(x1))
which(is.na(x1))</pre>
```

• Now, create the vector x2 and assess the difference to x1

```
x2 <- c(rnorm(10,5,2),NA,5:12,NA,6,7.5,NA,log(-2))
x2
```

- What is the meaning of "NA" versus "NaN"?
- Replace the missing values in x1 with a 0, and check that no NAs are present try two different commands to coerce the NAs into 0

```
x1[is.na(x1)] <- 0
is.na(x1)
# or
ifelse(is.na(x1),0,x1)</pre>
```

Exercise 10

- Import the data set water_errors.csv to R: A data frame with 61 observations on the following 6 variables.
 - location: a factor with levels North and South indicating whether the town is as north as Derby.
 - town: the name of the town.

- mortality: averaged annual mortality per 100.000 male inhabitants.
- hardness: calcium concentration (in parts per million).
- **smoker**: If there are any smokers living in town.
- num.of.cig: In case, smokers live in town, what number of cigarettes do they smoke per day.

```
# H20_err <- read_csv("C:\\Users\\admin\\Dropbox\\data\\water_errors.csv")
# str(H20_err)
# H20_err <- data.frame(H20_err)
# str(H20_err)
# BEST SOLUTION how to read it in:
# Try to use the "read.csv(...)" function to read data in!
# use the separator sep=";" or sep="," - which ever works better.
H20_err <- read.csv("C:\\Users\\admin\\Dropbox\\data\\water_errors.csv", sep=",")
str(H20_err)
# H20_err <- read.csv("~/Dropbox/201710_Makerere/03_Exercises/data/water_errors.csv")
H20_err <- read.csv("~/Dropbox/data/water_errors.csv", sep=",")
H20_err <- data.frame(H20_err)
str(H20_err)
head(H20_err)</pre>
```

• Detect the errors in the imported data set water_errors.csv in R.

```
str(H20_err)
table(H20_err$location) # Only one N and only one West observation.
table(H20_err$town) # LIVERPOOL is in capital letter.
summary(H20_err$mortality)
summary(H20_err$hardness) # hardness of -2 does not make sense, two NA's
table(H20_err$num.of.cig) # only one "zero" observation (wrong coding / level)
table(H20_err$smoker, H20_err$num.of.cig) # non-smokers who smoke more than 20?
```

• Find possible solutions in R how to handle these challenges.

```
str(H20_err)
which(H20_err$location == "N") # 6th row
which(H20_err$location == "West") # 9th row
```

```
H2O_err$location[H2O_err$location == "N"] <- "North"
H2O_err$location[H2O_err$location == "West"] <- NA # Option 1: Set to NA.
dim(H20_err)
H20_err <- H20_err[-which(H20_err$location == "West"), ] # Option 2: Remove from data.
dim(H20_err)
# H20_err$town[H20_err$town == "LIVERPOOL"] <- "Liverpool"</pre>
# H2O_err <- H2O_err$town[-which(H2O_err$town == "LIVERPOOL"), ]
which(is.na(H20_err$hardness))
H2O_err$hardness[which(is.na(H2O_err$hardness))] <- NA
H2O_err$hardness[which(H2O_err$hardness == -2)] <- NA
# H20_err$hardness[which(H20_err$hardness == -2)] <- 2</pre>
summary(H20_err$hardness)
# Check levels of varibale num.of.cig
levels(H20_err$num.of.cig)
table(H2O_err$num.of.cig)
# Change the zero level to none
H2O_err$num.of.cig[H2O_err$num.of.cig == "zero"] <- "none"
# Drop unused levels
H20_err$num.of.cig <- droplevels(H20_err$num.of.cig)</pre>
# levels(droplevels(H20_err$num.of.cig))
table(H20_err$num.of.cig)
which.F.morethan20 <- which(H20_err$smoker == FALSE & H20_err$num.of.cig == "more than 20")
H20_err[which.F.morethan20, ]
# OPTION 1:
H2O_err$num.of.cig[which.F.morethan20] <- NA
# OPTION 2:
H20_err$smoker[which.F.morethan20] <- TRUE</pre>
# check again, that we corrected it right
H20_err[which.F.morethan20, ]
table(H2O_err$smoker, H2O_err$num.of.cig) # check again!
which(H20_err$smoker == FALSE & H20_err$num.of.cig == "more than 20")
which.T.none <- which(H2O_err$smoker == TRUE & H2O_err$num.of.cig == "none")
H20_err[which.T.none, ]
H20_err$smoker[which.T.none] <- FALSE</pre>
```

```
table(H2O_err$smoker, H2O_err$num.of.cig)
```

• Do all variables have the correct data type (numeric, integer, factor)? - If not, do correct / define them.