

# BIOST 2039

## Lab 1

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## Instructions

Please **change the author** of this file to your name instead of mine in the header above.

Type your answer below each question. For questions that require the use of R, you can type both plain text (like this) and R commands by including an R chunk (see example below).

```
# This is an R chunk. Everything in this grey box is R code, not regular text.  
print("Hello, world!")
```

```
## [1] "Hello, world!"
```

You can run R chunks individually by clicking the green arrow in the top right corner of an R chunk.

When typing plain text in an R Markdown, leave two spaces at the end of the line when you want a paragraph break. Otherwise, paragraphs are typeset together, like this! For additional help with R Markdown, check out this reference sheet <https://www.rstudio.com/wp-content/uploads/2015/02/rmarkdown-cheatsheet.pdf>.

When you are finished, click **Knit** or **Knit to PDF** at the top of the Rmd editor. This will create an PDF file of that same name. Please submit the **PDF file** through Canvas.

```
set.seed(09042019) # Do not change this!
```

## Problem 1 (4 points)

When asked to convert a vector, do not store it as a new vector unless asked to do so.

Consider the vector:

```
num <- c("1.5", "2", "8")
```

A. Convert `num` into a numeric vector using `as.numeric()`.

```
num <- c("1.5", "2", "8")
num <- as.numeric(num)
print(num)
```

```
## [1] 1.5 2.0 8.0
```

B. Convert `num` into a factor using `factor`. Store this in a new vector `num_fac`.

```
num <- c("1.5", "2", "8")
num_fac <- factor(num)
print(num_fac)
```

```
## [1] 1.5 2 8
## Levels: 1.5 2 8
```

C. Convert `num_fac` into a numeric using `as.numeric()`.

```
num_fac <- as.numeric(num_fac)
print(num_fac)
```

```
## [1] 1 2 3
```

D. Convert `num_fac` into a numeric using `as.numeric(as.character())`.

```
num_fac <- as.numeric(as.character(num_fac))
print(num_fac)
```

```
## [1] 1 2 3
```

## Problem 2 (4 points)

When asked to convert a vector, do not store it as a new vector unless asked to do so.

Consider the vector:

```
num <- c(0, 1, 1, 0)
```

A. Append `TRUE` to the `num` vector using `c()`. Store the new vector as `num`.

```
num <- c(num, 'TRUE')
print(num)
```

```
## [1] "0" "1" "1" "0" "TRUE"
```

B. Check the class of `num` using `class()`.

```
num <- c(0, 1, 1, 0)
class(num)
```

```
## [1] "numeric"
```

C. Convert `num` into a logical vector using `as.logical()`.

```
num <- c(0, 1, 1, 0)
num <- as.logical(num)
print(num)
```

```
## [1] FALSE TRUE TRUE FALSE
```

D. Convert `num` into a logical vector where the result is `TRUE` if `num` is 0, using the `==` operator.

```
num <- c(0, 1, 1, 0)
num <- num == 0
print(num)
```

```
## [1] TRUE FALSE FALSE TRUE
```

### Problem 3 (4 points)

Consider the data set `data_bmi` which is contained in the `bmi_age.txt` file. Download the file from Canvas and store it in the same folder that you have this `*.Rmd` file stored.

```
file_name = "bmi_age.txt"
data_bmi = read.table(file = file_name, header = TRUE, stringsAsFactors = FALSE)
```

Perform the following operations:

A. Extract the column names of `data_bmi` using the `colnames()` function.

```
cat('\014')
```

```
colnames(data_bmi)
```

```
## [1] "PID" "BMI" "SEX" "AGE"
```

B. Extract the `AGE` column using the `[,]` operator.

```
cat('\014')
```

```
data_bmi[, 'AGE']
```

```
## [1] 45 57 66 49 33 40 65 59 65 42
```

C. Extract the `AGE` column using the `$` operator.

```
cat('\014')
```

```
data_bmi$AGE
```

```
## [1] 45 57 66 49 33 40 65 59 65 42
```

D. Extract the `AGE` and `BMI` columns using brackets and the `c()` function.

```
cat('\014')
```

```
data_bmi[, c('AGE', 'BMI')]
```

```
##      AGE BMI
## 1     45  22
## 2     57  27
## 3     66  31
## 4     49  24
## 5     33  23
## 6     40  18
## 7     65  21
## 8     59  26
## 9     65  34
## 10    42  20
```

## Problem 4 (3 points)

Here we will work with sequences and lengths:

A. Create a sequence from 1 to 4.5 by 0.24. Call this `run_num`.

```
cat('\014')
```

```
run_num <- seq(from = 1, to = 4.5, by = 0.24 )
print(run_num)
```

```
## [1] 1.00 1.24 1.48 1.72 1.96 2.20 2.44 2.68 2.92 3.16 3.40 3.64 3.88 4.12
## [15] 4.36
```

B. Use the `length()` function to find the length of `run_num`?

```
cat('\014')
```

```
length(run_num)
```

```
## [1] 15
```

C. Extract the fifth element of `run_num` using brackets.

```
cat('\014')
```

```
run_num[5]
```

```
## [1] 1.96
```

## Problem 5 (6 points)

Lets create a tibble called `df`:

```
df = dplyr::tibble(x = rnorm(10), y = rnorm(10), z = rnorm(10))
```

A. Extract the column `x` using the `$`.

```
cat('\014')
```

```
df$x
```

```
## [1] 0.5786590 -0.3981434 -0.4739019 0.2420337 0.4469141 0.1046046
## [7] 0.3013089 -1.3459991 -0.2647004 1.0165752
```

B. Extract the column `x` using the `[,]` notation.

```
cat('\014')
```

```
df['x']
```

```
## # A tibble: 10 x 1
##       x
##   <dbl>
## 1 0.579
## 2 -0.398
## 3 -0.474
## 4 0.242
## 5 0.447
## 6 0.105
## 7 0.301
## 8 -1.35
```

```
## 9 -0.265
## 10 1.02
```

C. Extract columns `x` and `z`.

```
cat('\014')
```

```
df[c('x','z')]
```

```
## # A tibble: 10 x 2
##       x      z
##   <dbl> <dbl>
## 1  0.579  0.371
## 2 -0.398  0.206
## 3 -0.474  0.0109
## 4  0.242  0.728
## 5  0.447 -0.280
## 6  0.105 -0.0368
## 7  0.301 -0.283
## 8 -1.35   0.900
## 9 -0.265  1.31
## 10 1.02   0.0730
```

D. Extract the third and fifth rows of `df` and columns `z` and `y`.

```
cat('\014')
```

```
df[c(3,5),c('z','y')]
```

```
## # A tibble: 2 x 2
##       z      y
##   <dbl> <dbl>
## 1  0.0109 -0.219
## 2 -0.280 -0.107
```

E. Get the mean of the column `x` using the `$` operator and `mean()`.

```
cat('\014')
```

```
mean(df$x)
```

```
## [1] 0.02073507
```

F. Pipe (`%>%`) the column `x` into the `mean()` function.

```
library(magrittr)
```

```
cat('\014')
```

```
df$x %>% mean()
```

```
## [1] 0.02073507
```

## Problem 6 (10 points)

Consider the data set `data_bmi` using the BMI data, but read in using `readr`:

```
file_name = "bmi_age.txt"
data_bmi = readr::read_table2(file = file_name)
```

```
## Warning: Missing column names filled in: 'X5' [5]
```

```
## Parsed with column specification:
## cols(
##   PID = col_double(),
##   BMI = col_double(),
##   SEX = col_double(),
##   AGE = col_double(),
##   X5 = col_logical()
## )

## Warning: 2 parsing failures.
## row col expected actual file
## 2 -- 5 columns 4 columns 'bmi_age.txt'
## 3 -- 5 columns 4 columns 'bmi_age.txt'
```

```
data_bmi
```

```
## # A tibble: 10 x 5
##   PID BMI SEX AGE X5
##   <dbl> <dbl> <dbl> <dbl> <lgl>
## 1 1 22 1 45 NA
## 2 2 27 0 57 NA
## 3 3 31 1 66 NA
## 4 4 24 1 49 NA
## 5 5 23 0 33 NA
## 6 6 18 0 40 NA
## 7 7 21 0 65 NA
## 8 8 26 1 59 NA
## 9 9 34 1 65 NA
## 10 10 20 0 42 NA
```

A. What is the class of data\_bmi?

```
cat('\014')
```

```
class(data_bmi)
```

```
## [1] "spec_tbl_df" "tbl_df"      "tbl"         "data.frame"
```

B. What is the class of data\_bmi[, "AGE"]?

```
cat('\014')
```

```
class(data_bmi[, 'AGE'])
```

```
## [1] "tbl_df"      "tbl"         "data.frame"
```

C. What is the class of data\_bmi\$AGE?

```
cat('\014')
```

```
class(data_bmi$AGE)
```

```
## [1] "numeric"
```

D. What is the mean of the AGE column?

```
cat('\014')
```

```
data_bmi$AGE %>% mean()
```

```
## [1] 52.1
```

E. Set the 3rd element of `data_bmi$AGE` to be 42.

```
cat('\014')
```

```
data_bmi$AGE[3]=42
```

F. What is the mean of the AGE column now?

```
cat('\014')
```

```
data_bmi$AGE %>% mean()
```

```
## [1] 49.7
```

G. Remove the X5 column using `data_bmi$X5 = NULL`.

```
cat('\014')
```

```
data_bmi$X5 = NULL
data_bmi
```

```
## # A tibble: 10 x 4
##       PID   BMI SEX  AGE
##   <dbl> <dbl> <dbl> <dbl>
## 1     1    22   1    45
## 2     2    27   0    57
## 3     3    31   1    42
## 4     4    24   1    49
## 5     5    23   0    33
## 6     6    18   0    40
## 7     7    21   0    65
## 8     8    26   1    59
## 9     9    34   1    65
## 10    10    20   0    42
```

H. Create `mat`, which is `data_bmi` as a matrix, using `as.matrix`.

```
cat('\014')
```

```
mat <- data_bmi %>% as.matrix()
mat
```

```
##       PID BMI SEX AGE
## [1,]    1  22   1  45
## [2,]    2  27   0  57
## [3,]    3  31   1  42
## [4,]    4  24   1  49
## [5,]    5  23   0  33
## [6,]    6  18   0  40
## [7,]    7  21   0  65
## [8,]    8  26   1  59
## [9,]    9  34   1  65
## [10,]  10  20   0  42
```

I. Try to extract AGE from `mat` using the `$`. What happened?

```
cat('\014')
```

```
mat$AGE
```

```
## Error in mat$AGE: $ operator is invalid for atomic vectors
```

J. Extract AGE from mat using the [,] notation.

```
cat('\014')
```

```
mat[, 'AGE']
```

```
## [1] 45 57 42 49 33 40 65 59 65 42
```

## Problem 7 (5 points)

Use the roll2() function that you created during the lab for this problem. You will need to re-define the function here by copying/pasting the code into the R chunk below.

```
# put your roll2() function here
roll2 <- function(){
  die <- seq(1:6)
  dice <- sample(die, 2, replace = TRUE)
  return(sum(dice))
}
```

A. Create a numeric vector called sim\_num which has the numbers 1 through 10.

```
cat('\014')
```

```
sim_num <- c(1,2,3,4,5,6,7,8,9,10)
print(sim_num)
```

```
## [1] 1 2 3 4 5 6 7 8 9 10
```

B. Create a character vector called roll which has the results of 10 runs of the roll2() function.

```
cat('\014')
```

```
roll <- as.character(sample(roll2(), size = 10, replace = TRUE))
print(roll)
```

```
## [1] "6" "6" "5" "3" "1" "5" "3" "7" "6" "6"
```

C. Create a data frame consisting of two columns using the two vectors you created above. Call this data frame myrolls.

```
cat('\014')
```

```
myrolls <- data.frame(sim_num, roll)
myrolls
```

```
##      sim_num roll
## 1         1    6
## 2         2    6
## 3         3    5
## 4         4    3
## 5         5    1
## 6         6    5
## 7         7    3
## 8         8    7
## 9         9    6
## 10        10    6
```

D. Convert myrolls into a tibble.



```
library(dplyr)
```

```
##  
## Attaching package: 'dplyr'  
## The following objects are masked from 'package:stats':  
##  
##   filter, lag  
## The following objects are masked from 'package:base':  
##  
##   intersect, setdiff, setequal, union
```

```
cat('\014')
```

```
myrolls <- as.tbl(myrolls)  
myrolls
```

```
## # A tibble: 10 x 2  
##   sim_num roll  
##   <dbl> <fct>  
## 1      1  1 6  
## 2      2  2 6  
## 3      3  3 5  
## 4      4  4 3  
## 5      5  5 1  
## 6      6  6 5  
## 7      7  7 3  
## 8      8  8 7  
## 9      9  9 6  
## 10     10 10 6
```

E. Using the data frame `myrolls` and the `table` function, create a table of the 10 roll results.

```
cat('\014')
```

```
table(myrolls)
```

```
##      roll  
## sim_num 1 3 5 6 7  
##      1 0 0 0 1 0  
##      2 0 0 0 1 0  
##      3 0 0 1 0 0  
##      4 0 1 0 0 0  
##      5 1 0 0 0 0  
##      6 0 0 1 0 0  
##      7 0 1 0 0 0  
##      8 0 0 0 0 1  
##      9 0 0 0 1 0  
##     10 0 0 0 1 0
```

## Session Info

```
sessionInfo()
```

```
## R version 3.5.3 (2019-03-11)  
## Platform: x86_64-w64-mingw32/x64 (64-bit)
```

```

## Running under: Windows 7 x64 (build 7601) Service Pack 1
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=English_United States.1252
## [2] LC_CTYPE=English_United States.1252
## [3] LC_MONETARY=English_United States.1252
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United States.1252
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] dplyr_0.8.3  magrittr_1.5
##
## loaded via a namespace (and not attached):
## [1] Rcpp_1.0.1      knitr_1.22      hms_0.5.1       tidyselect_0.2.5
## [5] R6_2.4.0        rlang_0.4.0     fansi_0.4.0     stringr_1.4.0
## [9] tools_3.5.3     xfun_0.6        utf8_1.1.4      cli_1.1.0
## [13] htmltools_0.3.6 yaml_2.2.0      assertthat_0.2.1 digest_0.6.18
## [17] tibble_2.1.3    crayon_1.3.4    readr_1.3.1     purrr_0.3.2
## [21] vctrs_0.2.0     zeallot_0.1.0   glue_1.3.1      evaluate_0.13
## [25] rmarkdown_1.12 stringi_1.4.3    compiler_3.5.3  pillar_1.4.2
## [29] backports_1.1.3 pkgconfig_2.0.2

```