# Undergrad Biostatistics - R Training - Week IV

#### Ege Ulgen

### R tips

- If you can't figure out how to solve an issue, Google is your friend. e.g., "how to calculate mode r"
- If you need help with the usage of a function, type ?function\_name. e.g. ?quantile
- If you get an error, and cannot fix it. C/P the error into Google. Someone else most likely had a similar problem
- Some resources for learning the basic syntax of R;
  - Codecademy https://www.codecademy.com/learn/learn-r
  - RStudio Cloud Primers https://rstudio.cloud/learn/primers
  - Dataquest https://www.dataquest.io/course/introduction-to-data-analysis-in-r/
  - R for Data Science https://r4ds.had.co.nz/index.html
- Interesting read: https://www.dataquest.io/blog/learn-r-for-data-science/

## Confidence Interval Example

### Prepare data

We'll read in the AIDS data as follows:

```
aids_df <- read.delim("../data/aids_dataset.txt", sep = " ")</pre>
```

We can take a look at the first 6 rows via head():

```
head(aids_df)
```

```
age gender week_1 cd4_1 week_2 cd4_2
##
     id treatment
             trt2 36.43
                           male
                                      0
                                            23
                                                 7.57
              trt4 47.85
                                      0
                                            21
                                                 8.00
                                                          49
                           male
             trt3 36.60
                           male
                                      0
                                                 7.14
                                                          61
     5
                                      0
                                                 8.00
             trt1 35.95
                                            36
                                                          31
                           male
     6
             trt2 38.40
                                      0
                                            11
                                                 7.29
                           male
                                                          11
             trt2 45.08
                                                 9.00
                           male
                                            11
                                                          41
```

We will use the first 10 patients (first 10 rows):

```
sub_df <- aids_df[1:10, ]</pre>
```

We'll define perc\_benefit and add it to the data frame:

##		id	${\tt treatment}$	age	gender	week_1	cd4_1	week_2	cd4_2	<pre>perc_benefit</pre>
##	1	1	trt2	36.43	male	0	23	7.57	21	-1.14870
##	2	2	trt4	47.85	male	0	21	8.00	49	16.66667
##	3	4	trt3	36.60	male	0	61	7.14	61	0.00000
##	4	5	trt1	35.95	male	0	36	8.00	31	-1.73611
##	5	6	trt2	38.40	male	0	11	7.29	11	0.00000
##	6	7	trt2	45.08	male	0	11	9.00	41	30.30303
##	7	8	trt3	37.20	male	0	16	7.71	11	-4.05318
##	8	11	trt2	42.25	male	0	16	4.14	21	7.54831
##	9	12	trt4	31.46	male	0	46	16.14	51	0.67346
##	10	13	trt4	41.86	male	0	1	17.00	1	0.00000

#### Calculate necessary values

Recall that the general formula for the confidence interval of the mean is:

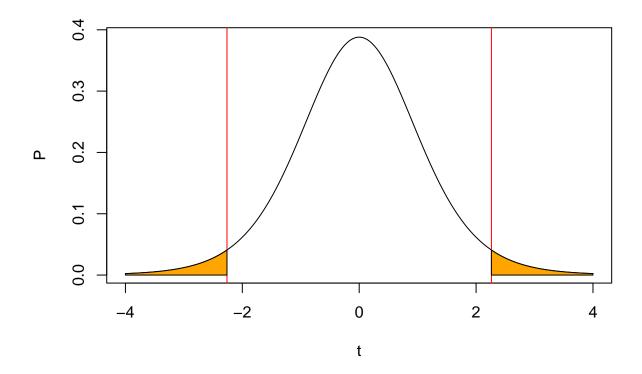
$$(1 - \alpha)\% \ CI = [\bar{X} - t^* \frac{s}{\sqrt{n}}, \bar{X} + t^* \frac{s}{\sqrt{n}}]$$

where  $\bar{X}$  is the sample mean, s is the sample standard deviation, n is the number of samples, and  $t^*$  is the critical value (that is dependent on the confidence level).

Let's first calculate the sample mean (stored in  $x_bar$ ) and the sample standard deviation (stored in  $s_x$ ):

```
x_bar <- mean(sub_df$perc_benefit)
s_x <- sd(sub_df$perc_benefit)</pre>
```

Next, we want to find the critical value  $t^*$ , such that 95% (our confidence level) of the area below the corresponding t distribution (displayed below) lies between  $-t^*$  and  $+t^*$ . The degrees-of-freedom of the corresponding t distribution (displayed below) is n-1=10-1=9



In the above plot of  $t_9$ , the white area is 95%. The left critical value is the 2.5th percentile ((1-.95)/2 = 0.05/2 = 0.025):

```
qt(0.025, df = 10 - 1)
```

## [1] -2.2622

### The 95% Confidence Interval

Recall that the general formula for the confidence interval of the mean is:

$$(1 - \alpha)\% \ CI = [\bar{X} - t^* \frac{s}{\sqrt{n}}, \bar{X} + t^* \frac{s}{\sqrt{n}}]$$

```
x_bar - 2.2622 * s_x / sqrt(10)

## [1] -2.8698

x_bar + 2.2622 * s_x / sqrt(10)

## [1] 12.521
```

#### via the built-in t.test() function'

```
t.test(sub_df$perc_benefit, conf.level = .95)
```

##
## One Sample t-test

```
##
## data: sub_df$perc_benefit
## t = 1.42, df = 9, p-value = 0.19
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## -2.8697 12.5204
## sample estimates:
## mean of x
## 4.8253

res <- t.test(sub_df$perc_benefit, conf.level = .90)
res$conf.int

## [1] -1.4102 11.0609
## attr(,"conf.level")
## [1] 0.9</pre>
```