

# Undergrad Biostatistics - R Training - Week IV

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## 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 = " ")
```

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

```
head(aids_df)
```

```
##   id treatment   age gender week_1 cd4_1 week_2 cd4_2
## 1  1      trt2 36.43  male      0    23   7.57    21
## 2  2      trt4 47.85  male      0    21   8.00    49
## 3  4      trt3 36.60  male      0    61   7.14    61
## 4  5      trt1 35.95  male      0    36   8.00    31
## 5  6      trt2 38.40  male      0    11   7.29    11
## 6  7      trt2 45.08  male      0    11   9.00    41
```

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

```
sub_df <- aids_df[1:10, ]
```

We'll define `perc_benefit` and add it to the data frame:

```
sub_df$perc_benefit <- (sub_df$cd4_2 - sub_df$cd4_1) / sub_df$cd4_1 / (sub_df$week_2 - sub_df$week_1) *
sub_df
```

##	id	treatment	age	gender	week_1	cd4_1	week_2	cd4_2	perc_benefit
## 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:

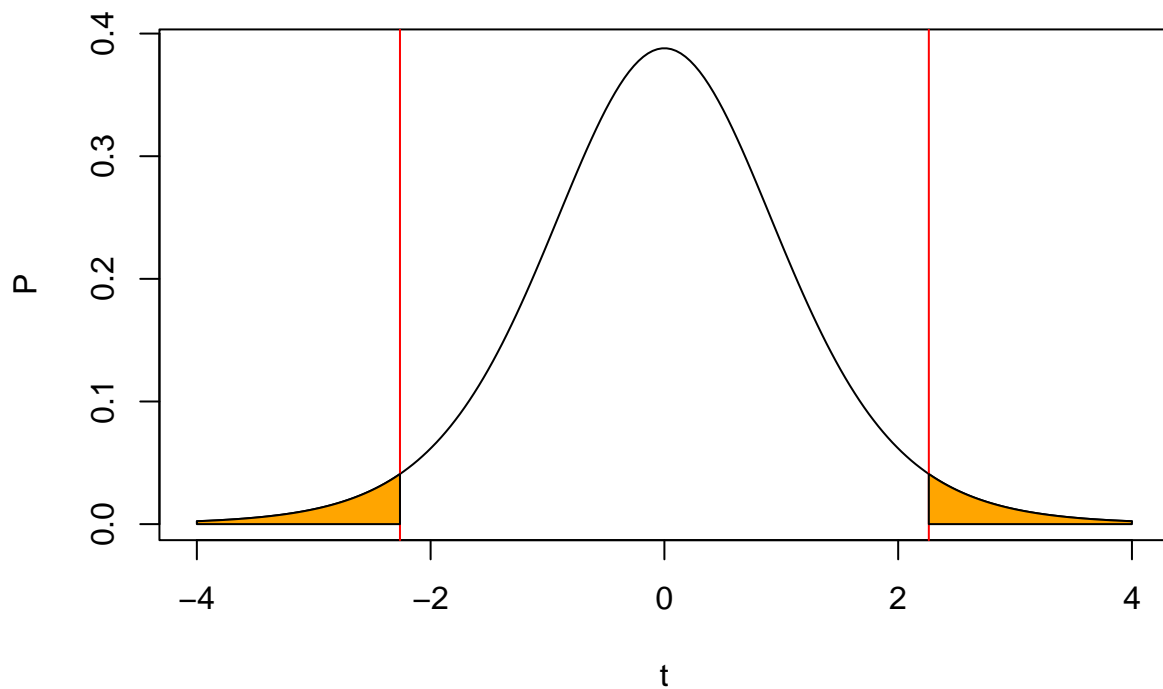
$$(1 - \alpha)\% \text{ CI} = [\bar{X} - t^* \frac{s}{\sqrt{n}}, \bar{X} + t^* \frac{s}{\sqrt{n}}]$$

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)
```

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
```