

# MTH142 Intro to Stats:: CHEAT SHEET

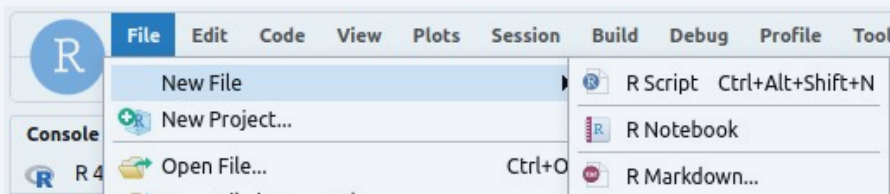
## Getting started

Signing into RStudio

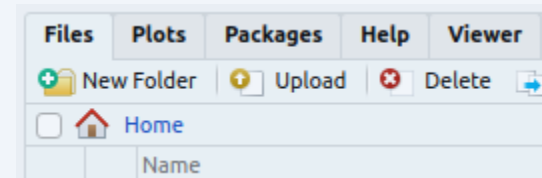
<http://hcc-rstudiosrv1.hcc.edu:8787/>

Username: Your HCC email with @hcc.edu  
Password: Your HCC password

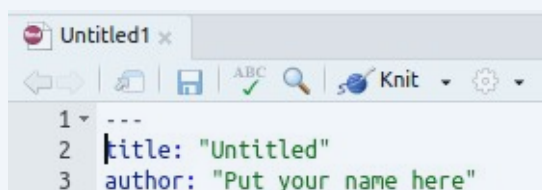
## .Rmd files



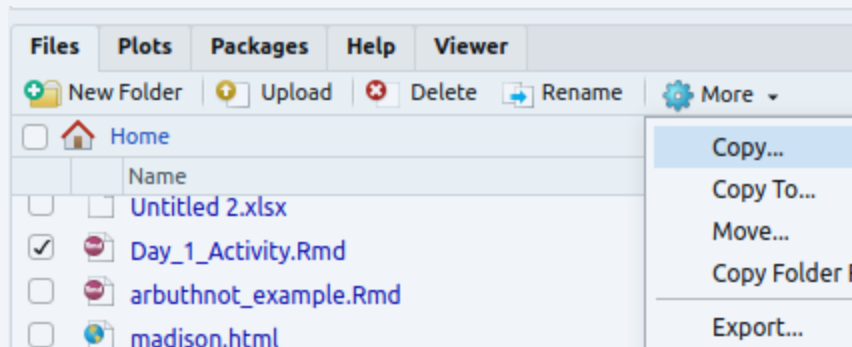
**Uploading a file** Lower Right quadrant click upload



**Knitting a file** click the knit button and save.



**Exporting a file select file** in lower right quadrant click **more** then **Export...**



## Screen Shots

**Mac:** Shift + Command + 3

**Windows:**  Shift + S

**Chromebook:** Shift + Ctrl + 

## Libraries

```
library(openintro)
library(tidyverse)
library(broom)
```

To get more info on a dataset,  
load a library then use ?  
library(openintro)  
?cars93

## View Data Sets

**cars93** is the data set  
**glimpse(data = cars93)**  
**head(cars93)**  
First six rows (tail() works too)  
**view(cars93)**  
makes window for all data  
**names(cars93)**  
show variable names only

## Summary Statistics

**fivenum(cars93\$price)**

Just the five number summary

**summary(cars93)**

summarizes all the variables

**tally(cars93)**

counts the observations

**mean(cars93\$weight, na.rm = TRUE)**

calculates the mean of the weights

and removes missing values.

**sd(cars93\$weight)**

calculates the standard deviation

## Variables

**a\_variable <- c(1,2,3)**

takes the vector 1,2,3 and  
saves it as a variable with the  
assignment operator

## Math Type

Use \$\$ to enclose math type.

$\mu$	$\sigma$
$\bar{x}$	$\hat{p}$
$H_o$	$\alpha$
$\neq$	$\approx$
$\sim$	

## Confidence Intervals

**For a mean from a vector of data**

**t.test(cars93\$weight)**

produces a 95% confidence interval for the mean  
weights of cars sold in 1993.

**For a proportion from Statistics**

**prop.test(x = 20, n = 60, conf.level=0.90,correct=FALSE)**

produces a 90% confidence interval for the population  
proportion given 20 successes in 60 trials.

**For mean use formula below**

For the confidence interval of the mean from statistics we use the  
following formula:

$$\bar{x} \pm t * SE$$

## Graphs

**box plots**

```
ggplot(data = cars93,
       mapping = aes(x = weight, fill = type)) +
  geom_boxplot()
```

**histograms**

```
ggplot( cars93, aes(x = weight)) +
  geom_histogram()
```

**bar**

```
ggplot(cars93, aex(x= type )) +
  geom_bar()
```

**scatter plots**

```
ggplot(data=cars93, aes(x = weight, y = mpg_city)) +
  geom_point()+
  geom_smooth(method = "lm", se=FALSE)
```

**titles and labels**

```
ggplot(cars93, aes(x= type )) +
  geom_bar()+
  labs(x= 'WEIGHTS', title = 'Cars from 93')
```

## List of Standard Error formulas

$$SE = \sqrt{\frac{p(1-p)}{n}}$$

SE for a single proportion

$$SE = \frac{\sigma}{\sqrt{n}} \approx \frac{s}{\sqrt{n}}$$

SE for a single mean or paired  
means.

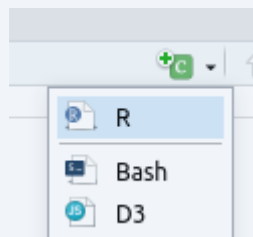
$$SE = \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$$

SE for two means (not paired)

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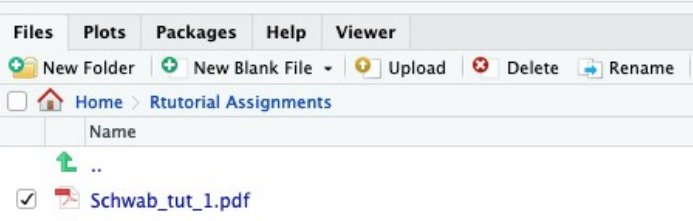
## R Markdown cont.

**New chunk** click the green c+ and select r



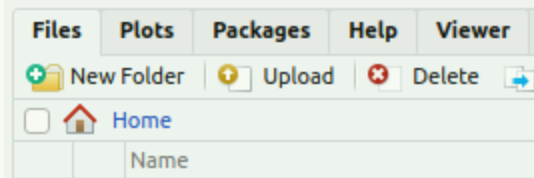
### Rename a file

Click the box next to the file and chose rename.

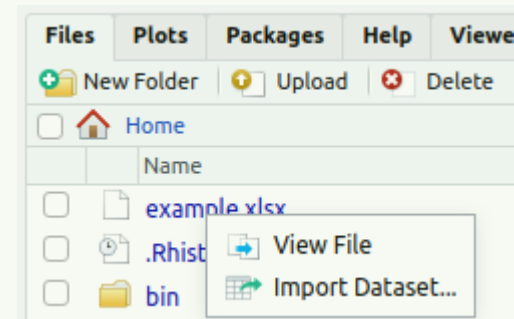


## Uploading Data

1. Copy and paste data into a spreadsheet.
2. Variables are column headers (no spaces in variable name)
3. Save as an .xls file to your computer (no spaces in file names)
4. Upload .xls file



5. Click on the file you uploaded > Import Dataset



## normal distribution

**Pnorm( q= 1, mean=0 ,sd=1)**

Outputs the probability if z=1 from a standard normal dist.

**qnorm(p = 0.01,mean=0, sd=1)**

Outputs the z-score if probability=0.01

$$z^* = \frac{\hat{p} - p}{SE}$$

This is the formula for a z\*-score for a single proportion.

## simulation

library(infer)

# 1. Make a vector to mimic data

```
more_than_1_relationship <- c(
  rep(TRUE, 152),
  rep(FALSE, 51)
)
```

# 2. change the vector to a data frame

```
more_than_1_relationship <- as.data.frame(more_than_1_relationship)
```

# 3 run infer chain

```
set.seed(2024)
```

```
null_distn_one_prop <- more_than_1_relationship |>
specify(response = more_than_1_relationship, success = "TRUE") |>
hypothesize(null = "point", p = 0.5) |>
generate(reps = 10000, type = "draw") |>
calculate(stat = "prop")
```

# 4 Take a look at the distribution

```
visualise(data = null_distn_one_prop)
```

```
get_p_value(x = null_distn_one_prop,
  obs_stat = 0.749,
  direction = 'greater')
```

## student t distribution

**pt(1,df=11, lower.tail = FALSE)**

outputs the p-value if t = 1 and 11 degrees of freedom on the upper tail of the distribution

**qt(0.01,df=11)**

outputs the t-score if probability = 0.01

$$t^* = \frac{\bar{x} - \mu}{SE}$$

This is the formula for a t\*-score for a single mean.

## Hypothesis Tests

### Proportions

**prop.test(x=20, n=60, p=0.5, correct=FALSE)**

Outputs the results of hypothesis test with 20 successes in 60 trials

### Single Mean

**t.test(cars93\$weight, mu=2000, alternative="g")**

Outputs of a right tailed hypothesis test with parameter mu=2000.

### Difference of Means

**t.test(time\_hrs~division,data=nyc\_marathon, alternative="t")**

Computes the difference of men and women winning times in NYC marathon

### ANOVA

**cars\_anova<-aov(weight~type,data=cars93)**

performs the analysis of variance and saves it as a variable “cars\_anova”

**anova(cars\_anova)**

outputs the summary of the analysis

### Linear Regression

**cars\_linear<-lm(price~weight,data=cars93)**

performs the linear regression and saves it as a variable “cars\_linear”

**summary(cars\_linear)**

outputs the linear summary of the regression