**Workshop: Dealing with Data in R** 

Getting help in R

After this workshop



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# First things first

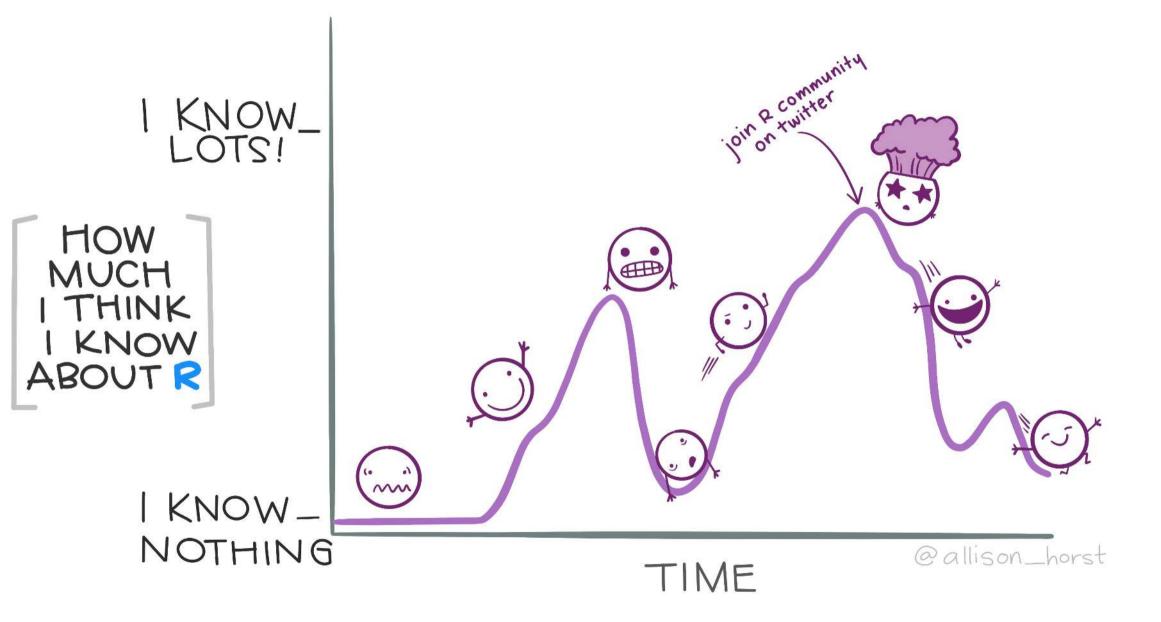
Save previous script
Open New File

(make sure you're in the RStudio Project)

Add library(tidyverse) to the top

Save this new script

consider names like troubleshoting.R or 5\_getting\_help.R



# **Troubleshooting**

## Line by line

- R is sequential
- If you skip lines, you're not running that part

```
#library(tidyverse)
count(mtcars, am)
```

## Error in count(mtcars, am): could not find function "count"

## Line by line

## 2 1 13

- R is sequential
- If you skip lines, you're not running that part

```
#library(tidyverse)
count(mtcars, am)
## Error in count(mtcars, am): could not find function "count"
```

• Error? Start at the beginning and go line by line

```
library(tidyverse)
count(mtcars, am)

## am n
## 1 0 19
```

## Line by line

Especially important if loading and modifying data

```
# Load Data
size <- read_csv("./data/grain_size2.csv")</pre>
# First modification
size <- mutate(size,</pre>
               total sand = coarse sand + medium sand + fine sand,
               total silt = coarse silt + medium silt + fine silt)
# Second modification
size <- size %>%
                                                                   Can't run 1st modification
  group_by(plot) %>%
                                                                     after 2nd modification
  summarize(n = n(),
            total_sand = sum(total_sand),
            mean_sand = mean(total_sand),
            sd sand = sd(total sand),
            se_sand = sd_sand / sqrt(n))
```

## **Section by section**

```
## Error: Problem with `summarise()` column `mean_sand`.
## i `mean_sand = mean(totall_sand)`.
## x object 'totall_sand' not found
## i The error occurred in group 1: plot = "CSP01".
```

## **Section by section**

```
size <- read_csv("./data/grain_size2.csv")</pre>
```

No error

## **Section by section**

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### No error

### No error

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## **Section by section**

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## i `mean_sand = mean(totall_sand)`.
## x object 'totall_sand' not found
## i The error occurred in group 1: plot = "CSP01".
```

### Ah ha!

## **Applies to error messages too**

- First, don't panic!
- Look at the error bit by bit

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## Error: Problem with `summarise()` column `mean_sand`.
## i `mean_sand = mean(totall_sand)`.
## x object 'totall_sand' not found
## i The error occurred in group 1: plot = "CSP01".
```

## **Applies to error messages too**

```
Error: Problem with 'summarise()' column 'mean_sand'`
```

Okay, we know the problem is in the **summarize()** part and then **mean\_sand** part of that

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```
i 'mean_sand = mean(totall_sand)'
x object 'totall_sand' not found
```

Looks like this is the line with the problem.

And the problem is object 'totall\_sand' not found. Ooops! Typo!

## **Applies to error messages too**

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Error: Problem with 'summarise()' column 'mean_sand'`
```

Okay, we know the problem is in the **summarize()** part and then **mean\_sand** part of that

```
i 'mean_sand = mean(totall_sand)'
x object 'totall_sand' not found
```

Looks like this is the line with the problem.

And the problem is object 'totall\_sand' not found. Ocops! Typo!

```
i The error occurred in group 1: plot = "CSP01".
```

Lastly, it's telling us that the problem was when working with this group of data. (This can be useful when troubleshooting, because you can **filter()** your data and take a look)

# debugging



I got this.



Huh. Really thought that was it.





4. Fine. Restarting.



5. OH WTF.



b.. Zombie meltdown







A NEW HOPE!



9. [insert awesome] theme song



I V CODING!

# R is never wrong

# R is never wrong

Just sometimes unhelpful!

# **Getting Help**

## **Cheat Sheets**

## **RStudio Menu**

- Help
  - Cheatsheets

Take a look yourself

## **Cheat Sheets**

### RStudio Menu

- Help
  - Cheatsheets

Take a look yourself

## Data Visualization with ggplot2:: CHEAT SHEET

#### Basics

**ggplot2** is based on the **grammar of graphics**, the idea that you can build every graph from the same components: a data set, a coordinate system. and geoms-visual marks that represent data points.



To display values, map variables in the data to visual properties of the geom (aesthetics) like size, color, and x and y locations.



Complete the template below to build a graph.



ggplot(data = mpg, aes(x = cty, y = hwy)) Begins a plot that you finish by adding layers to. Add one geom function per layer.

aesthetic mappings | data | geom

qplot(x = cty, y = hwy, data = mpg, geom = "point") Creates a complete plot with given data, geom, and mappings. Supplies many useful defaults.

last plot() Returns the last plot

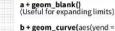
ggsave("plot.png", width = 5, height = 5) Saves last plot as 5' x 5' file named "plot.png" in working directory. Matches file type to file extension.

#### Geoms

Use a geom function to represent data points, use the geom's aesthetic properties to represent variables. Each function returns a layer.

#### GRAPHICAL PRIMITIVES

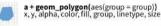
a <- ggplot(economics, aes(date, unemploy)) b <- ggplot(seals, aes(x = long, y = lat))



b + geom\_curve(aes(yend = lat + 1, xend=long+1),curvature=1) - x, xend, y, yend, alpha, angle, color, curvature, linetype, size



a + geom\_path(lineend="butt", linejoin="round" x, y, alpha, color, group, linetype, size



b + geom rect(aes(xmin = long, vmin=lat, xmax= long + 1, ymax = lat + 1)) - xmax, xmin, ymax, ymin, alpha, color, fill, linetype, size



a + geom\_ribbon(aes(ymin=unemploy - 900, ymax=unemploy + 900)) - x, ymax, ymin, alpha, color, fill, group, linetype, size

#### LINE SEGMENTS

common aesthetics: x, y, alpha, color, linetype, size



b + geom abline(aes(intercept=0, slope=1))

b + geom\_hline(aes(yintercept = lat)) b + geom\_vline(aes(xintercept = long))

b + geom\_segment(aes(vend=lat+1, xend=long+1)) b + geom spoke(aes(angle = 1:1155, radius = 1))

#### ONE VARIABLE continuous

c <- ggplot(mpg, aes(hwy)); c2 <- ggplot(mpg)



c + geom\_area(stat = "bin") x, y, alpha, color, fill, linetype, size



c + geom\_density (kernel = "gaussian") x, y, alpha, color, fill, group, linetype, size, weight



c + geom\_dotplot() x, y, alpha, color, fill



c + geom\_freqpoly() x, y, alpha, color, group,



c + geom\_histogram(binwidth = 5) x, y, alpha, color, fill, linetype, size, weight



c2 + geom\_qq(aes(sample = hwy)) x, y, alpha, color, fill, linetype, size, weight

#### discrete

d <- ggplot(mpg, aes(fl))



d + geom bar() x, alpha, color, fill, linetype, size, weight

#### TWO VARIABLES

continuous x, continuous v e <- ggplot(mpg, aes(cty, hwy))



e + geom\_label(aes(label = cty), nudge\_x = 1, nudge\_y = 1, check\_overlap = TRUE) x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust



e + geom\_jitter(height = 2, width = 2) x, y, alpha, color, fill, shape, size



e + geom\_point(), x, y, alpha, color, fill, shape, e + geom\_quantile(), x, y, alpha, color, group, linetype, size, weight



e + geom\_rug(sides = "bl"), x, y, alpha, color, linetype, size



e + geom\_smooth(method = lm), x, y, alpha, color, fill, group, linetype, size, weigh



e + geom\_text(aes(label = cty), nudge\_x = 1 nudge\_y = 1, check\_overlap = TRUE), x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust

#### discrete x . continuous v f <- ggplot(mpg, aes(class, hwy))



f + geom\_col(), x, y, alpha, color, fill, group, linetype, size



f + geom\_boxplot(), x, y, lower, middle, upper, ymax, ymin, alpha, color, fill, group, linetype, shape, size, weight



f + geom\_dotplot(binaxis = "y", stackdir = "center"), x, y, alpha, color, fill, group



f + geom\_violin(scale = "area"), x, y, alpha, color, l, group, linetype, size, weight

#### discrete x , discrete v

g <- ggplot(diamonds, aes(cut, color))



g + geom\_count(), x, y, alpha, color, fill, shape, size, stroke

#### continuous bivariate distribution

h <- ggplot(diamonds, aes(carat, price))



h + geom\_bin2d(binwidth = c(0.25, 500)) x, y, alpha, color, fill, linetype, size, weight

ggplot2



h + geom\_density2d() x, y, alpha, colour, group, linetype, size



h + geom hex() x, y, alpha, colour, fill. size



i <- ggplot(economics, aes(date, unemploy))



i + geom area() x, y, alpha, color, fill, linetype, size



i + geom line() x, y, alpha, color, group, linetype, size



df < data.frame(grp = c("A", "B"), fit = 4:5, se = 1:2)i <- ggplot(df, aes(grp, fit, ymin = fit-se, ymax = fit+se))



j + geom\_crossbar(fatten = 2) x, y, ymax, ymin, alpha, color, fill, group, linetype,



j + geom\_errorbar(), x, ymax, ymin, alpha, color, group, linetype, size, width (also



j + geom\_linerange() x, ymin, ymax, alpha, color, group, linetype, size



j + geom\_pointrange() x, y, ymin, ymax, alpha, color, fill, group, linetype

data <- data frame/murder = USArrests\$Murder. state = tolower(rownames(USArrests))) map <- map\_data("state") k <- ggplot(data, aes(fill = murder))



k + geom\_map(aes(map\_id = state), map = map)
+ expand\_limits(x = mapSlong, y = mapSlat), map\_id, alpha, color, fill, linetype, size

#### THREE VARIABLES

seals\$z <- with(seals, sgrt(delta\_long^2 + delta\_lat^2)); l <- ggplot(seals, aes(long, lat))



l+geom contour(aes(z=z)) x, y, z, alpha, colour, group, linetype, size, weight



I + geom\_raster(aes(fill = z), hjust=0.5, vjust=0.5, interpolate=FALSE) x, y, alpha, fill



I + geom\_tile(aes(fill = z)), x, y, alpha, color, fill, linetype, size, width



# Vignettes

Many packages come with vignettes (aka, R tutorials)

## **List Vignettes**

# Vignettes

Many packages come with vignettes (aka, R tutorials)

## **List Vignettes**

## **Load Vignettes**

```
vignette("ggplot2-specs", package = "ggplot2")
```

## Try it!

## **Tutorials**

## **Vignettes also online**

• e.g., tidyverse

## **Organizations/Websites**

- Software Carpentry
- STHDA



ggplot2 is a system for declaratively creating graphics, based on The Grammar of Graphics. You provide the data, tell ggplot2 how to map variables to aesthetics, what graphical primitives to use, and it takes care of the details.

### Installation

```
# The easiest way to get ggplot2 is to install the whole tidyverse:
install.packages("tidyverse")
# Alternatively, install just ggplot2:
install.packages("ggplot2")
# Or the the development version from GitHub:
# install.packages("devtools")
devtools::install_github("tidyverse/ggplot2")
```

Download from CRAN at https://cloud.r-project.org/ package=ggplot2

Browse source code at https://github.com/tidyverse/ ggplot2

Report a bug at

https://github.com/tidyverse/ ggplot2/issues

Learn more at

http://r4ds.had.co.nz/datavisualisation.html

Extensions at

http://www.ggplot2-exts.org/ gallery/

License

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## **Books!**

### **Free Online**

- R for Data Science (read it!)
- R Graphics Cookbook (how to do X)
- ggplot2 (next level)
- <u>Data Visualization: A practical introduction</u>
- <u>Geocomputation with R</u> (spatial, GIS, maps)
- Statistical Inference via Data Science: A ModernDive into R and the tidyverse (stats)

# **Specific help**

# **Examples**

## In R

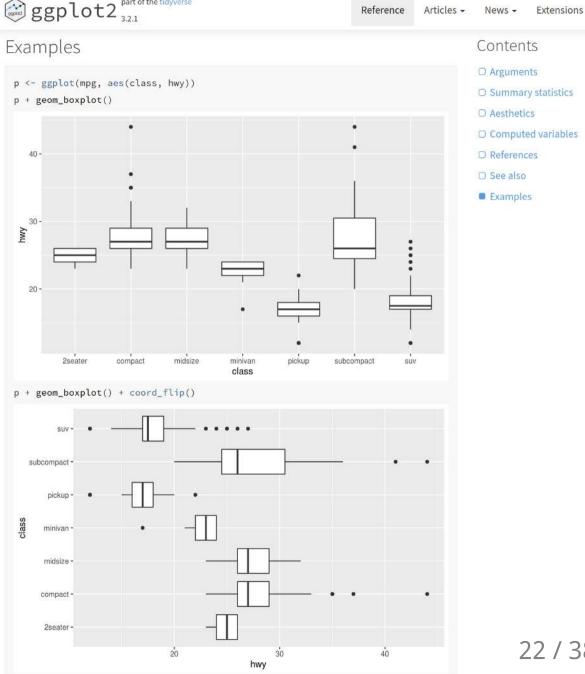
?geom\_boxplot

Copy and paste the examples into your console

# **Examples**

## On the web

- Nice to see expected output
- Helps figure out if it's your system or your code



## Web searches

- Always include "R" in the search
- Include the package name!
- Use keywords
- Some errors are very general

## Web searches

- Always include "R" in the search
- Include the package name!
  - Try "R boxplots" vs. "R boxplots ggplot2"
- Use keywords
  - Try "R boxplots ggplot2 notch"
- Some errors are very general
  - Try "R Error: object 'm' not found"

## Stackoverflow etc.

"R how to remove duplicate rows"

## Stackoverflow etc.

## Things to consider

- Date (i.e., R version, Package Version)
- Packages used (tidyverse? R base? A mix?)
- What are the example data?
  - mtcars and iris are commonly used data sets built into R base
  - msleep and diamonds are commonly used data sets built into ggplot2
- What are the example columns?
- What is actually required to answer *your* question?

# **Asking for Help**

## Not useful

- "I got an error"
- "It didn't work"

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## **Better!**

- "I got *this* error"
- "It didn't give me *this*"

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- "It didn't give me *this*"

### **Best!!**

- "I did *this* and I got *this* error"
- "I expected it to do *this*, but in fact the output was *this*"

## **Asking for Help**

#### Not useful

- "I got an error"
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#### **Better!**

- "I got *this* error"
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#### Best!!

- "I did *this* and I got *this* error"
- "I expected it to do *this*, but in fact the output was *this*"

#### **Best of the Best!!!**

• "I did this [small reproducible code, including data set] and I got this [exact error/output]"

- Minimal code and data required to reproduce the error
- Often preparing this actually helps you solve the error!
- Includes
  - o packages (library())
  - data
  - runnable code

### How do I change the order of vore?

### Not reproducible

```
## Error in ggplot(data = m, aes(x = vore, y = awake, fill = `Body Size`)): could not find function
"ggplot"
```

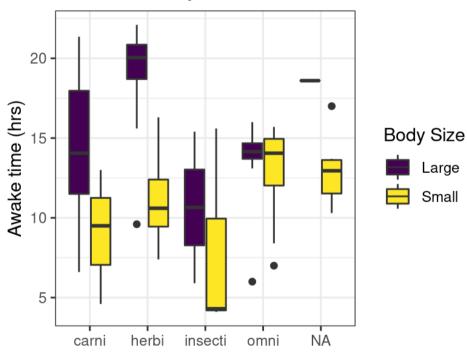
- No indication of packages
- No indication of what **m** is

### How do I change the order of vore?

### Reproducible, but not minimal

```
library(ggplot2)
m <- msleep %>%
 mutate(`Body Size` = if_else(bodywt > median(bodywt),
                               "Large", "Small"))
ggplot(m, aes(x = vore, y = awake, fill = `Body Size`)) +
  theme bw() +
  theme(axis.title.x = element_blank()) +
  geom_boxplot() +
  scale_fill_viridis_d() +
  labs(y = "Awake time (hrs)",
       title = "Awake time by Diet")
```

### Awake time by Diet

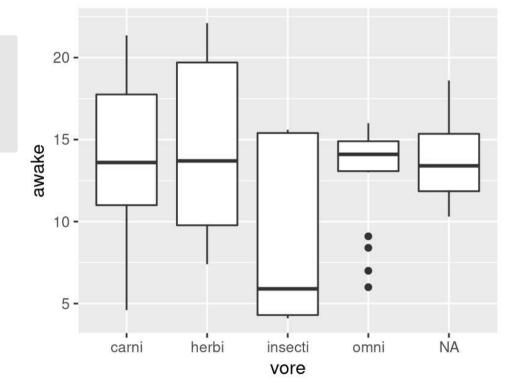


### How do I change the order of vore?

### **Reproducible AND Minimal**

```
library(ggplot2)

ggplot(msleep, aes(x = vore, y = awake)) +
  geom_boxplot()
```



# **Paying it forward**

## **Citing Software**

#### **In-line Text**

- Software name
- Version
- Programmers/authors OR Journal article releasing the software (if available)

### **Bibliography**

- Journal article releasing the program OR
- Programmers/authors
- Year of release
- Program Name
- URL

## Citing R

### **Inline**

"All statistical analyses were performed with R statistical software (v3.6.2, R Core Team 2019)."

### **Bibliography**

R Core Team (2019). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL <a href="https://www.R-project.org/">https://www.R-project.org/</a>.

## Citing R

#### **Version information**

```
R.Version()$version.string
## [1] "R version 4.1.2 (2021-11-01)"
```

#### **Citation information**

```
##
## To cite R in publications use:
##
## R Core Team (2021). R: A language and environment for statistical
## computing. R Foundation for Statistical Computing, Vienna, Austria.
## URL https://www.R-project.org/.
```

## **Citing R Packages**

#### **Inline**

"All statistical analyses were performed with R statistical software (v4.0.3, R Core Team 2020). We performed Type III ANOVAs using the 'car' package for R (v3.0.10, Fox and Weisberg)."

### **Bibliography**

John Fox and Sanford Weisberg (2019). An R Companion to Applied Regression, Third Edition. Thousand Oaks CA: Sage.

## Citing R Packages

#### **Version information**

```
packageVersion("car")
## [1] '3.0.12'
```

#### **Citation information**

```
##
## To cite the car package in publications use:
##
## John Fox and Sanford Weisberg (2019). An {R} Companion to Applied
## Regression, Third Edition. Thousand Oaks CA: Sage. URL:
## https://socialsciences.mcmaster.ca/jfox/Books/Companion/
```

See more about citing packages in my rOpenSci blog post: How to Cite R and R packages

at first I was like ... mw WWW

You made it!

Thank you!