

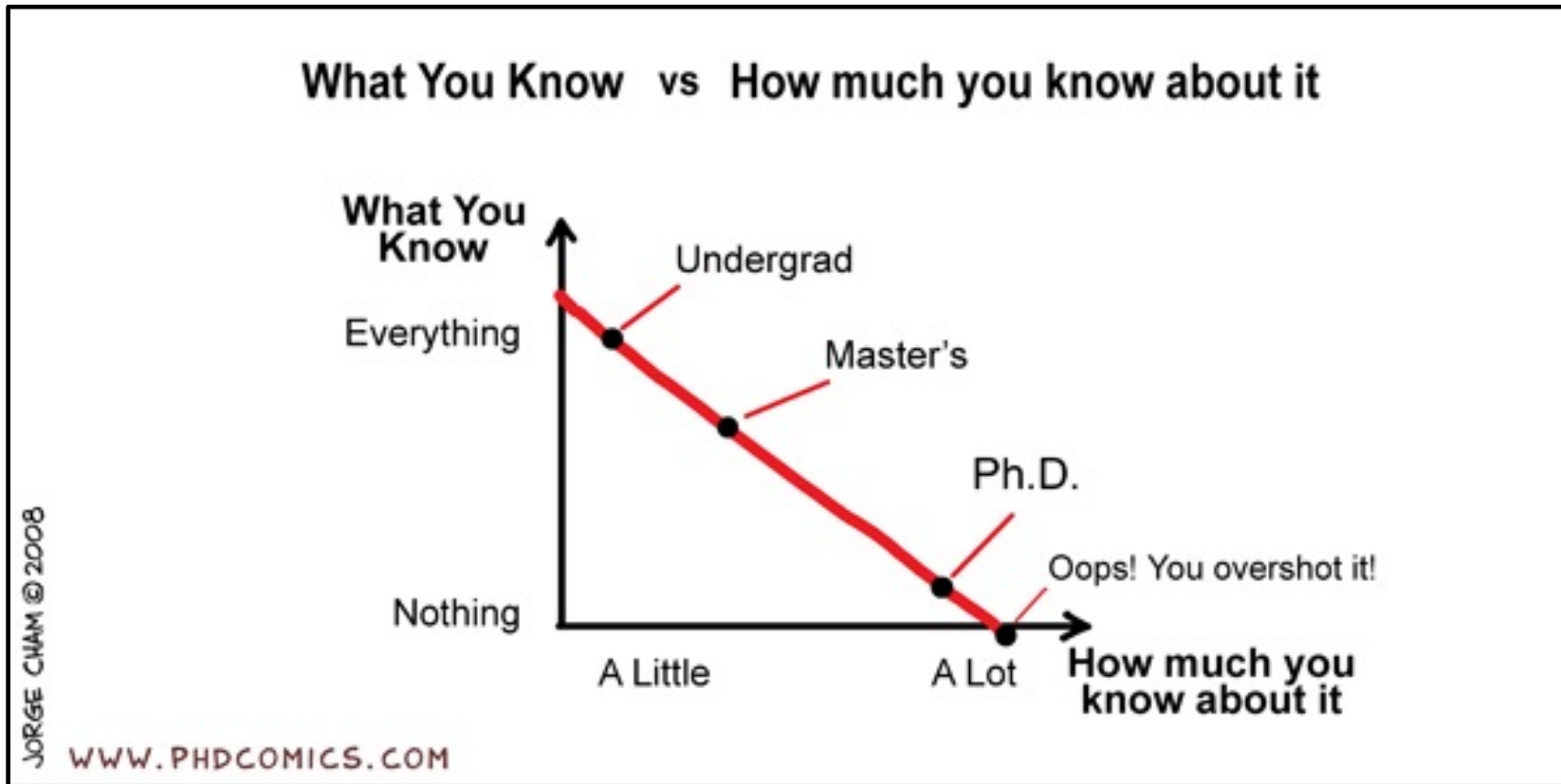
# Peer-Feedback Survey

1 survey every Tuesday (constant link)



[https://ubc.ca1.qualtrics.com/jfe/form/SV\\_bvLa6xMXdRk1j6u](https://ubc.ca1.qualtrics.com/jfe/form/SV_bvLa6xMXdRk1j6u)

# BIOL 501: Graphics



Get me out, or make a new one

**FIRST NAME**  
Preferred pronouns

# Outline

- Peer-feedback survey on Discussion Moderation and Presenting
- Why make graphs?
- 4 Principles of effective display
- Types of graphs to achieve these principles
- Why some graphs fail, and what can be done
- Tables
- Interactive Graphs
- Ggplot
- Motion (video)
- Wrap-up

# Discussion

What kind of graphs do you have experience making or do you think you will make in your thesis?

**Go to Jamboard:**

<https://jamboard.google.com/d/103Uolikn1ZhLSpJ5ururRoUIRr1JCsqoGb3ibRPfko/edit?usp=sharing>

## **Instructions to add a sticky note (anonymous)**

1. Click the 4th icon down on the left "sticky note".
2. Type in the sticky note, sentences, phrases
3. Click "SAVE" to submit



You can submit multiple sticky notes

# Recommendations for data entry

# Best practices for data entry and management

UBC Biology 501 R workshops

Biol 501 home

R Workshops ▾

R tips pages ▾

Install R on your own computer

Install packages

Work at your own speed

Get help

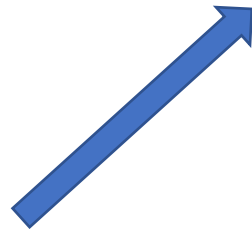
Get answers

## Biol 501 R workshops

Use the **R Workshops** dropdown menu to navigate to the R workshops.

For help, access the [R tips pages](#).

Recommendations for data entry handout [here](#). Best practices and simple guidelines for effective data management



Bring the.csv of your thesis dataset to a workshop, I'm happy to look at the headers, and layout of it **before** you import it into R

# A common mistake in organizing databases for categorical variables

- **Question:** Can heart rate predict metabolic rate and does this relationship differ between single dives vs dive bouts?
- X=heart rate (numerical)
- Y=metabolic rate (numerical)
- Treatments in this LME model are fixed effects (categorical variable)
  - Single dive
  - Dive bout
- Animal is a random effect

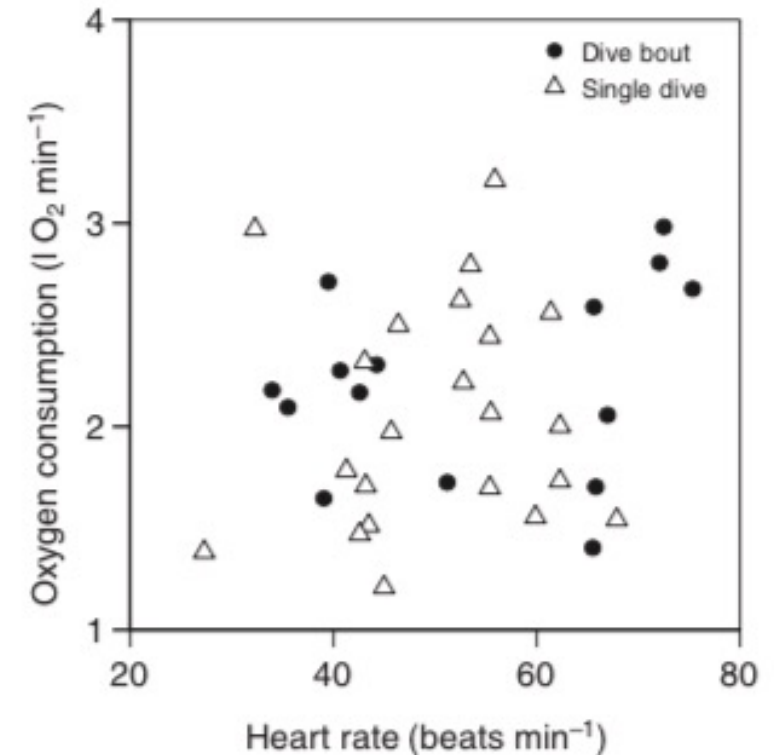


Fig. 2. The relationships between diving  $f_H$  and DMR for Steller sea lions over a dive bout or over a single dive were not significantly linear.





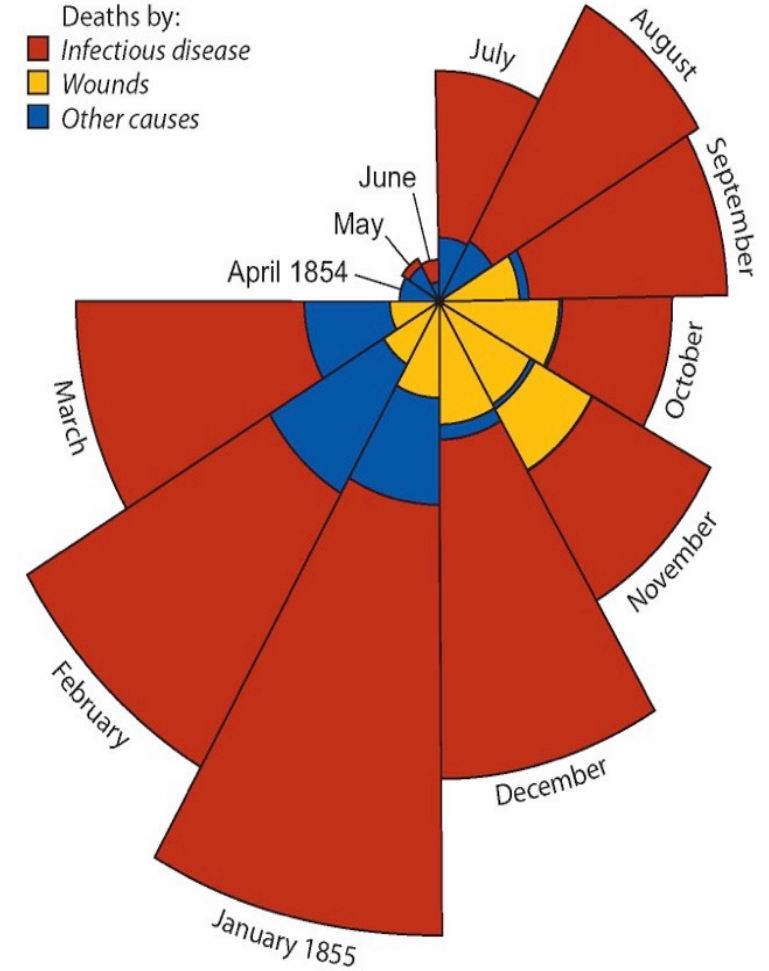
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# Why make graphs?

- Because they can change the world!
- The human eye is a pattern detector.
- Enable visual comparisons of measurements between groups and expose relationships between variables.
- Best method for communicating results to a wide audience

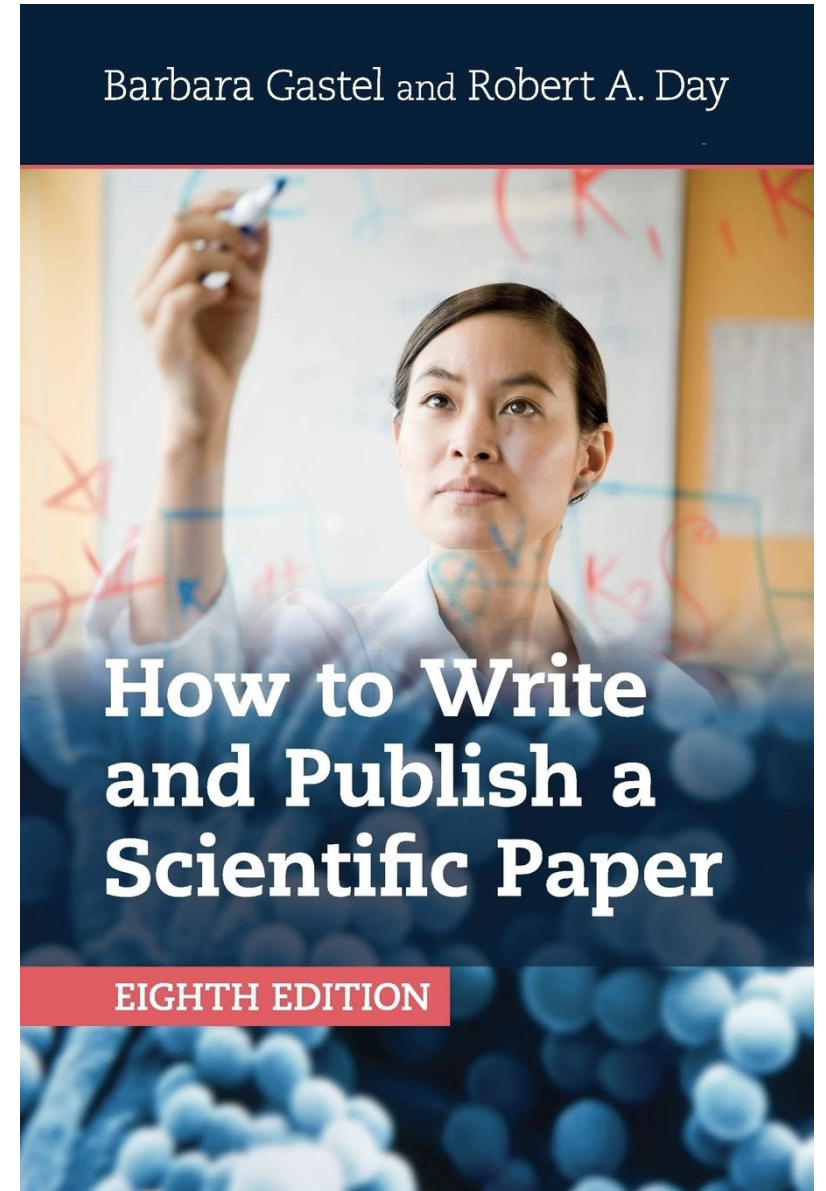
**Goal: show a pattern clearly  
& quickly**



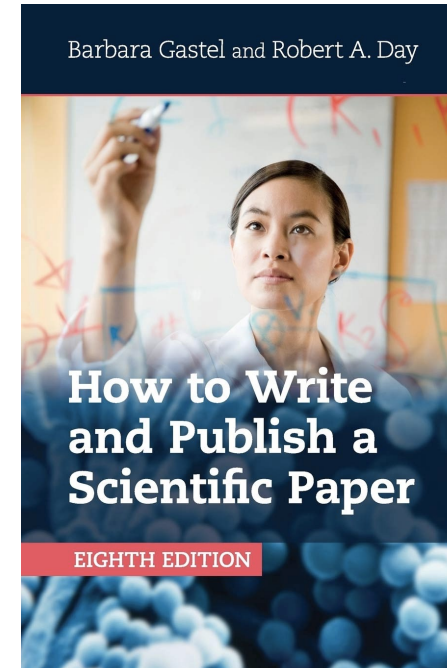
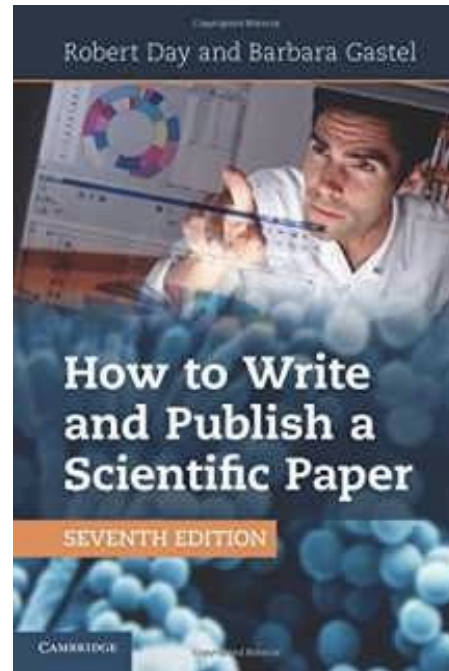
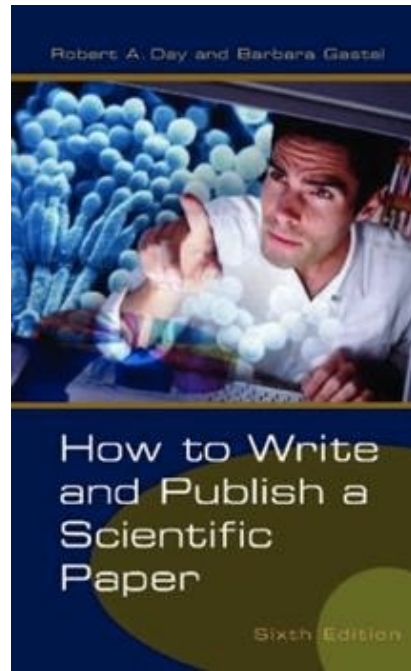
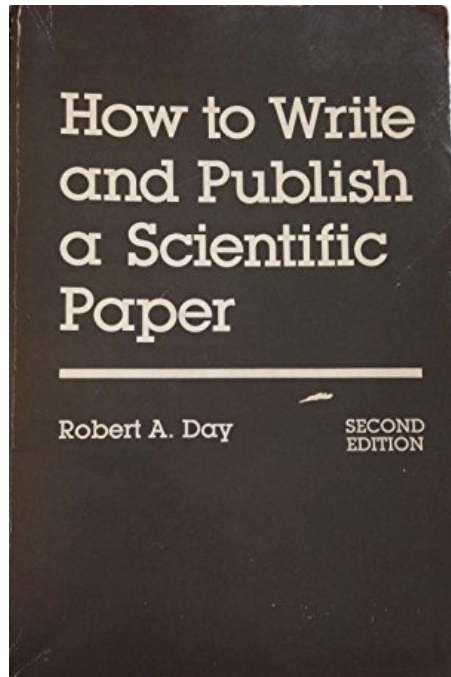
Causes of deaths in the British Army during the Crimean War (F. Nightingale 1858)  
(area of pie = number of deaths)

# Resource on writing and publishing scientific papers

- Chapter 16: Tables
- **Chapter 17: Graphs**
- Presentations
- CVs
- Writing manuscripts
- How to submit a manuscript and review process



Currently in 9<sup>th</sup> edition since ~ 1979



If you want an electronic copy of 8<sup>th</sup> edition, see me during the workshop

# Why do we use graphs?

- To show patterns in data “greatest number of ideas in shortest time”
- To show spread, shape of distribution—frequency distributions
- To show associations between variables
  - The relationship between two or more variables (linear, nonlinear, other)
  - Differences between groups

Graphs should make the viewer goes “Oh!” and not “Huh?”

# TRUISMS



A PICTURE IS WORTH A THOUSAND WORDS.

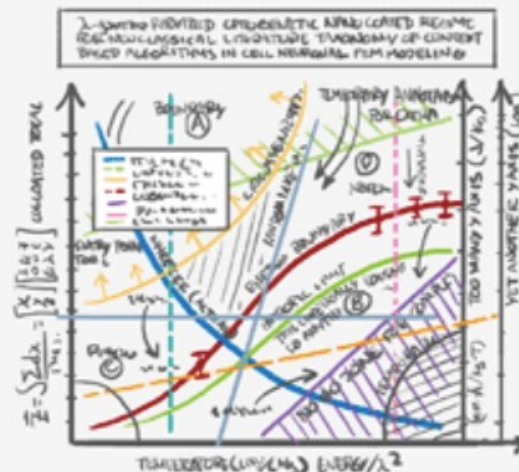


FIGURE 1.

A PICTURE WITH A THOUSAND WORDS IS USUALLY WORTH A PH.D.

WOW!

JORGE CHAM © 2017

WWW.PHDCOMICS.COM

# How to achieve graphical excellence?

*“Graphical excellence is that which gives to the viewer the **greatest number of ideas in the shortest time** with the least ink in the smallest space”*

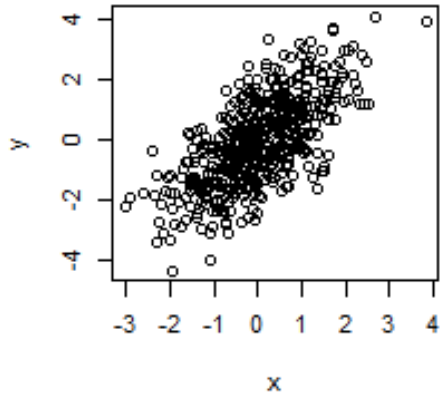
– Tufte (1983)

# Take home point

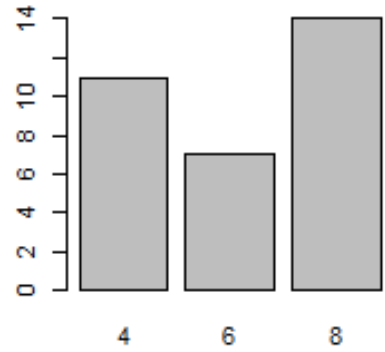
- An entire manuscript key points should be understandable with only the figures +tables+captions
  - No undefined abbreviations or unfamiliar terms
  - Include units
  - Include sample sizes



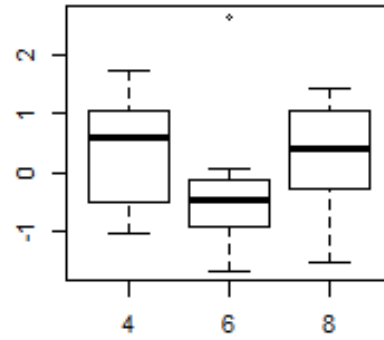
Scatterplot



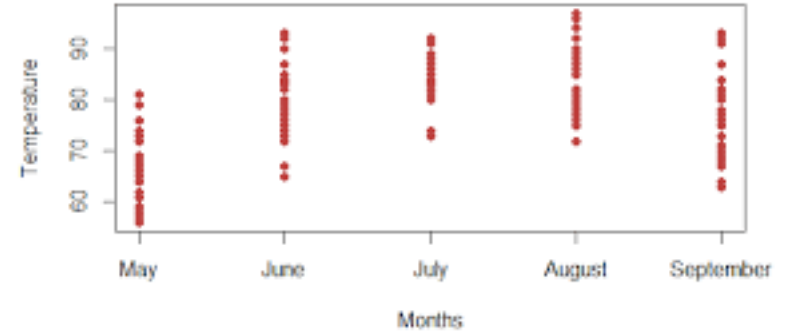
Barplot



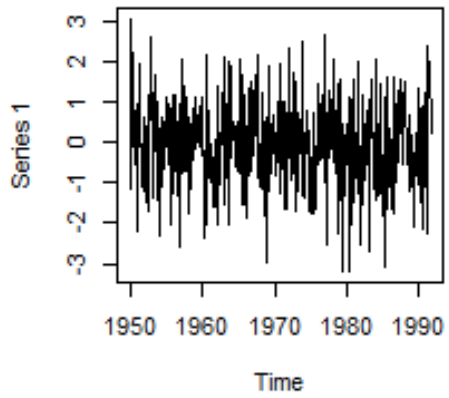
Boxplot



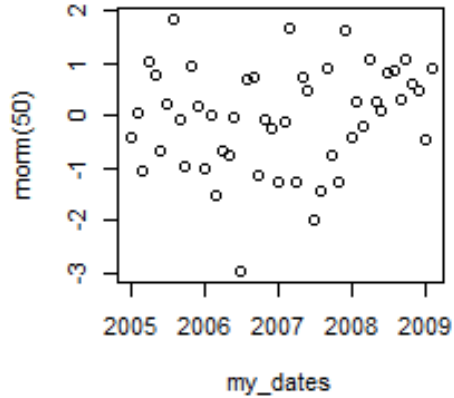
Different strip chart for each month



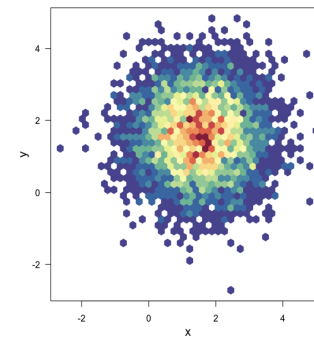
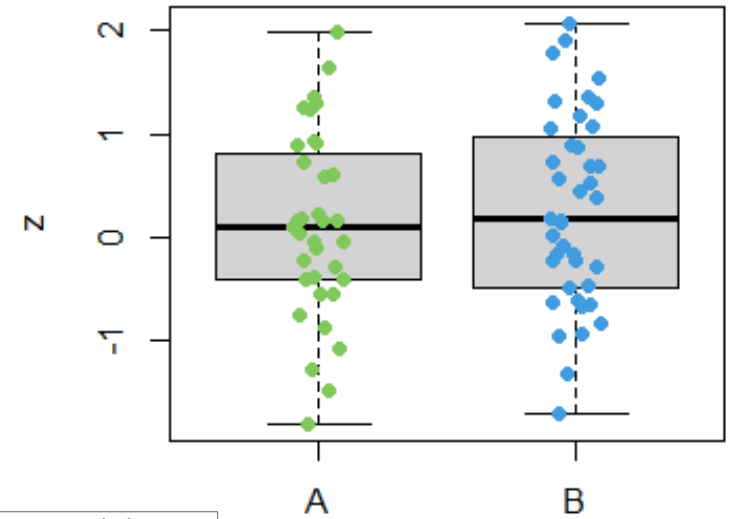
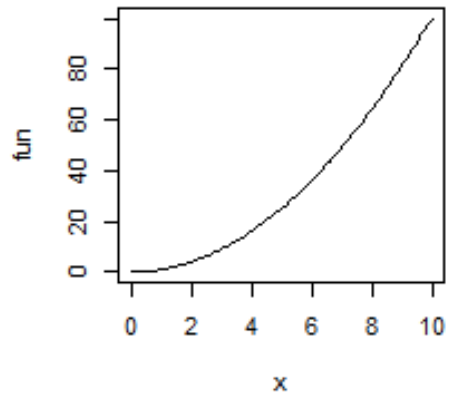
Time series



Time based plot



Plot a function



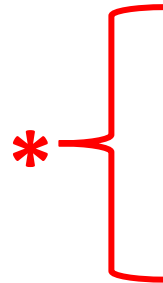
Endless types of graphs available in R

# 4 Principles to Increase Effectiveness of Graphs

1. Show the data
2. Make patterns in the data easy to see
3. Represent magnitudes honestly
4. Draw graphical elements clearly, minimizing clutter

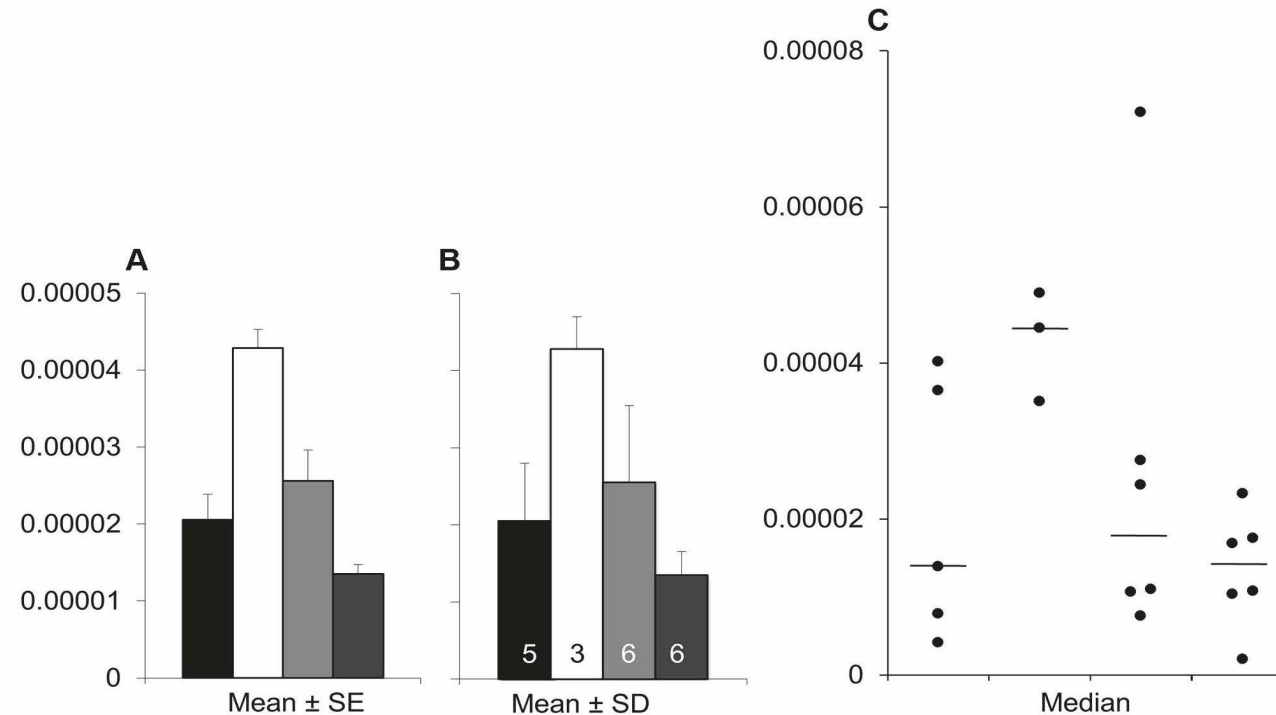
**\*Hint: The rubric for Assignment #1 specifically refers to these 4 principles**

Assignment number 1: Improve a Graph					
Full name of student: _____					
Student #: _____					
Assignment score out of 15: _____					
Written interpretation of the Bad Graph (2 marks)					
	Excellent (full mark)	Average	Poor	Missing	
Written explanation of the study (enough detail to understand graph, but no too verbose)	1		0.5		Total per section
Analyze the "bad graph". Explain what patterns it intended to show and why it was not effective at accomplishing this?	1		0.5		2
Written interpretation & Plotting the New graph (8 marks)					
	Excellent (full mark)	Average	Poor	Missing	
Written explanation explaining how your improvements display the patterns more effectively	2		1		
1. Show the data (is all the data shown accurately?)	1		0.5		
2. Make patterns in the data easy to see. Are the patterns easier to see in this NEW graph compared to the original?	2		1		
3. Represent magnitudes honestly. Is the baseline (0) reasonable, are the axis scales reasonable?	1		0.5		
4. Draw graphical elements clearly, minimizing clutter. Is the new graph more clear or less cluttered?	1		0.5		Total per section
Does the graph have a well-written caption that stands alone from text?	1		0.5		8
Quality of the R code (5 marks)					
	Excellent (full mark)	Average	Poor	Missing	
Is the code well annotated (with # comments) so that a new person could understand it?	2		1		
Are the variable names and assignments appropriate (not too long, not too short) and logical if a new person was using this code?	1		0.5		Total per section
Could the code be extended to another situation without starting again (is the code reproducible)? Is it "general" to be re-used?	2		1		5
					<b>Total Score out of 15</b>
					<b>15</b>



# 1. Show the data

*“While scatterplots prompt the reader to critically evaluate the statistical tests and the authors’ interpretation of the data, bar graphs discourage the reader from thinking about these issues”*

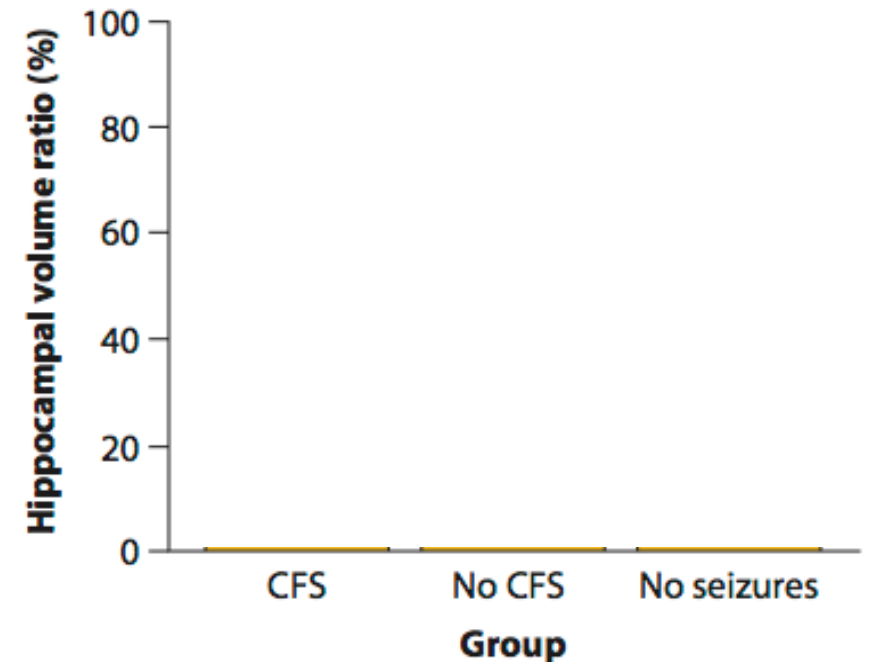
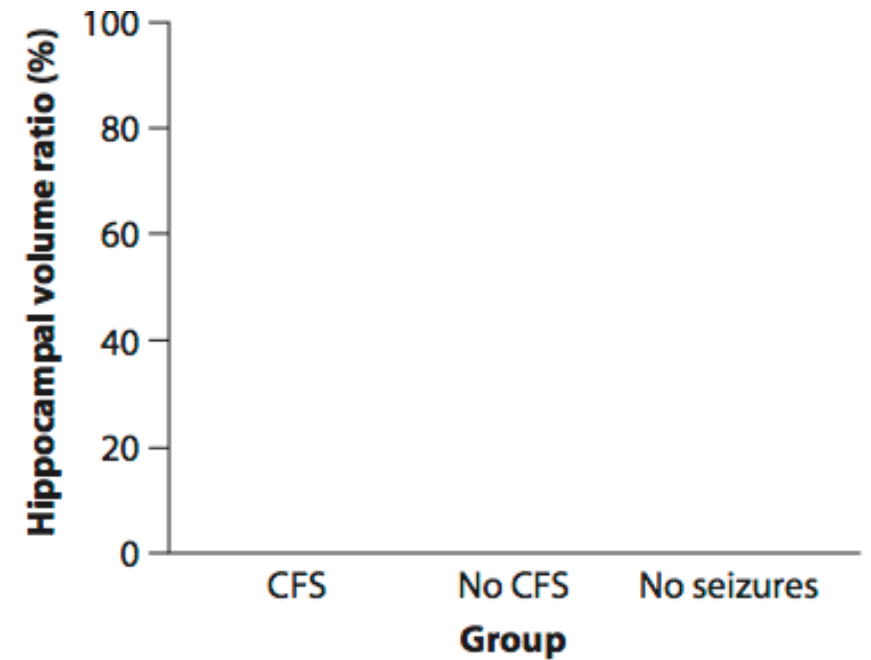


Weissgerber et al. (2015) Beyond bar and line graphs: time for a new data presentation paradigm.

*PLoS Biol.* DOI:10.1371/journal.pbio.1002128

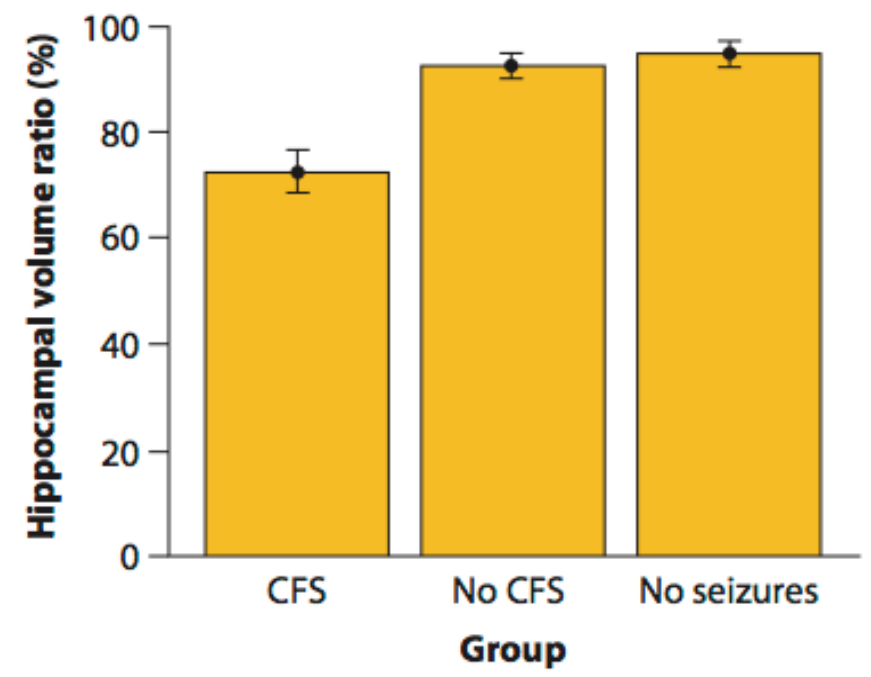
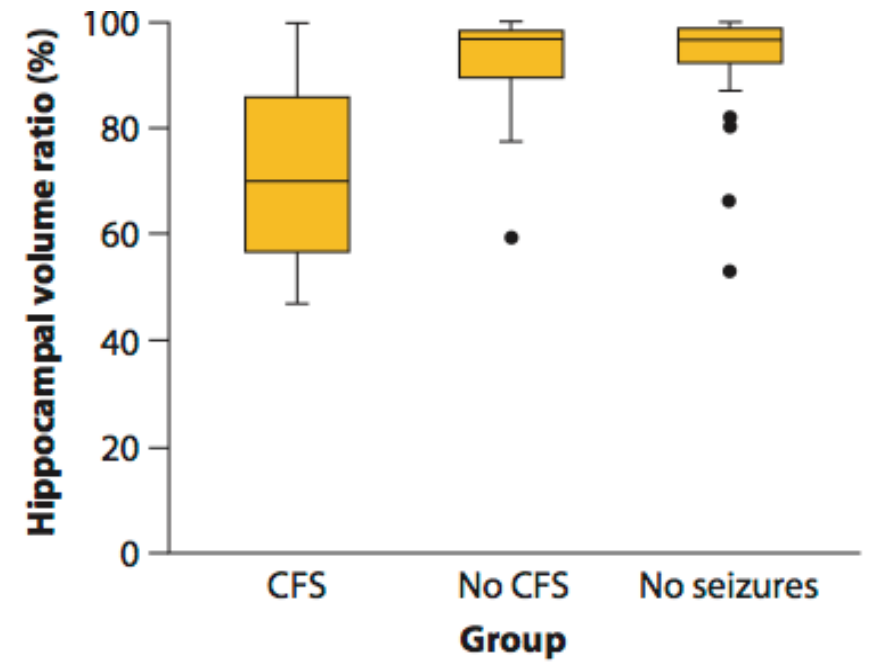
# 1. Show the data

- Hippocampal volume loss in patients with drug resistant epilepsy
  - Y-variable=numerical
- Treatments (X-variable categorical)
  - Childhood febrile seizures (CFS)
  - Childhood non-febrile seizures (no CFS)
  - No childhood seizures



# 1. Show the data

Discussion:  
These show the same data.  
Which graph is more effective ? Why?

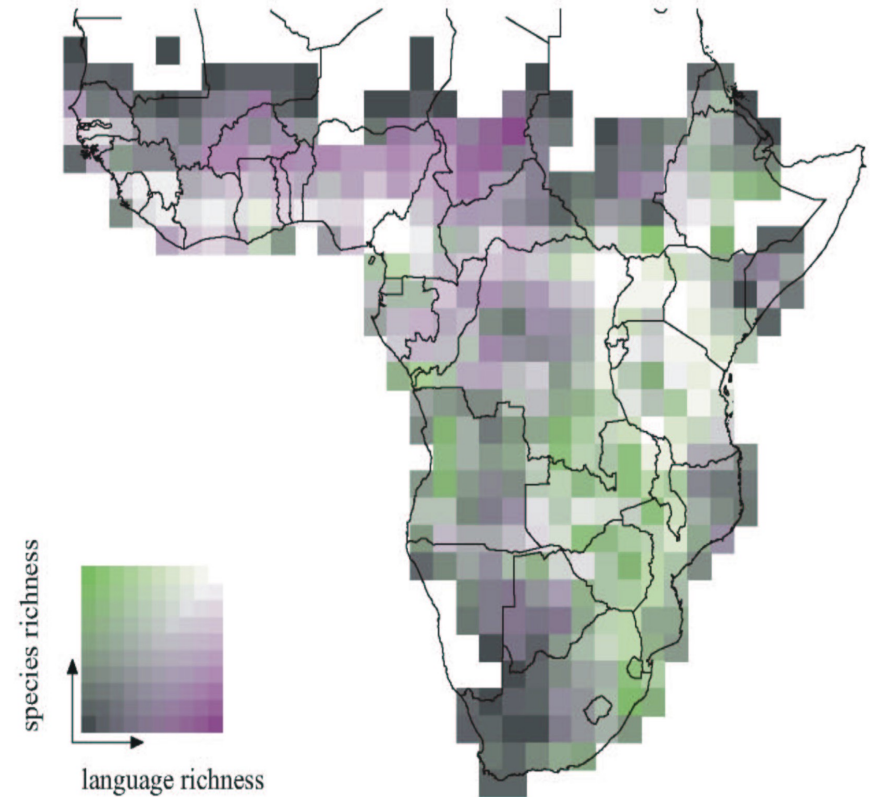


## 2. Make patterns easy to see

*“Graphical excellence consists of complex ideas communicated with clarity, precision and efficiency” – Tufte (1983)*

## 2. Make patterns easy to see

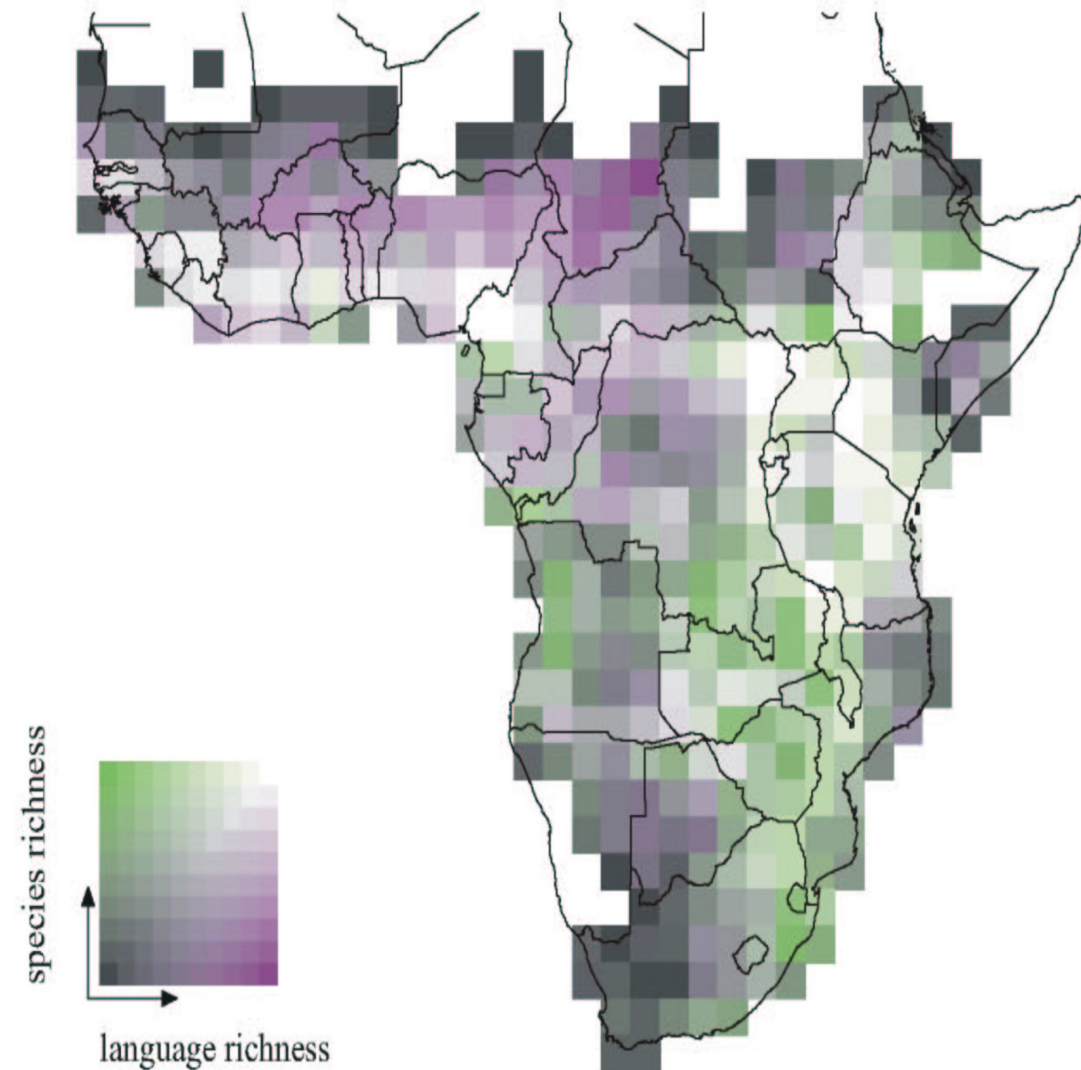
- Number of bird species and number of distinct human languages present in each square of a grid of continental Africa.
- What is the pattern in these data? How long did it take you to “see”?
- Is it easy to appreciate how strong the relationship is between the variables?





## 2. Make patterns easy to see

- Fig. 2. A comparison of language and species richness in sub-Saharan Africa. Each variable has been partitioned into 10 equal-frequency ranks. Cells of equal rank for both variables are coloured in grey scale from black (both low rank) to white (both high rank). The degree of colour saturation indicates that the ranks are different. A cell is green if the species-richness rank is high relative to language richness and blue if language richness is high relative to species richness.



## The distribution of cultural and biological diversity in Africa

Joslin L. Moore<sup>1,2\*</sup>, Lisa Manne<sup>3</sup>, Thomas Brooks<sup>1,4</sup>, Neil D. Burgess<sup>1,5</sup>, Robert Davies<sup>6</sup>, Carsten Rahbek<sup>2</sup>, Paul Williams<sup>3</sup> and Andrew Balmford<sup>1</sup>

<sup>1</sup>*Conservation Biology Group, Department of Zoology, University of Cambridge, Downing Street, Cambridge CB2 3EJ, UK*

<sup>2</sup>*Zoological Museum, University of Copenhagen, Universitetsparken 15, DK-2100, Copenhagen Ø, Denmark*

<sup>3</sup>*Biogeography and Conservation Laboratory, The Natural History Museum, Cromwell Road, London SW7 5BD, UK*

<sup>4</sup>*Center for Applied Biodiversity Science, Conservation International, 1919 M Street NW, Suite 600, Washington, DC 20036, USA*

<sup>5</sup>*Conservation Science Programme, World Wildlife Fund, 1250 24th Street NW, Washington, DC 20037-1193, USA*

<sup>6</sup>*Llanunwas, Upper Solva, near Haverfordwest, Pembrokeshire SA62 6UJ, UK*

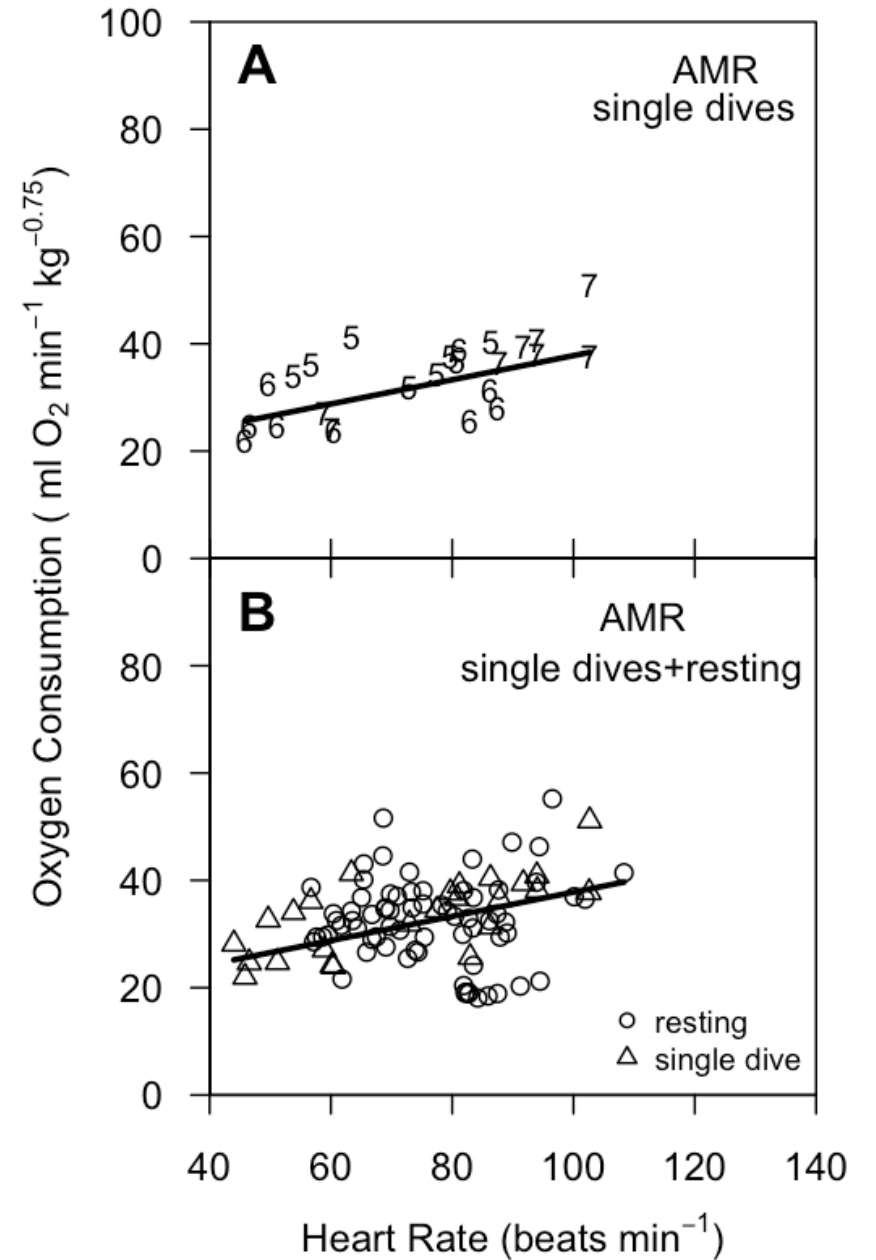
Anthropologists, biologists and linguists have all noted an apparent coincidence in species diversity and human cultural or linguistic diversity. We present, to our knowledge, one of the first quantitative descriptions of this coincidence and show that, for  $2^\circ \times 2^\circ$  grid cells across sub-Saharan Africa, cultural diversity and vertebrate species diversity exhibit marked similarities in their overall distribution. In addition, we show that 71% of the observed variation in species richness and 36% in language richness can be explained on the basis of environmental factors, suggesting that similar factors, especially those associated with rainfall and productivity, affect the distributions of both species and languages. Nevertheless, the form of the relationships between species richness and language richness and environmental factors differs, and it is unlikely that comparable mechanisms underpin the similar patterns of species and language richness. Moreover, the fact that the environmental factors considered here explain less than half of the variation in language richness indicates that other factors, many of which are likely to be historical or social, also influence the distribution of languages.

### 3. Represent magnitudes honestly

- *“A graphic does not distort if the visual representation of the data is consistent with the numerical representation” – Tufte (1983)*
- Bad examples:
  - Compare 2 plots with different Y-axis scales
  - Hide outliers
- Does the data illustrate what the model/analysis was actually doing?

# Example: Does the data show what the model was actually doing?

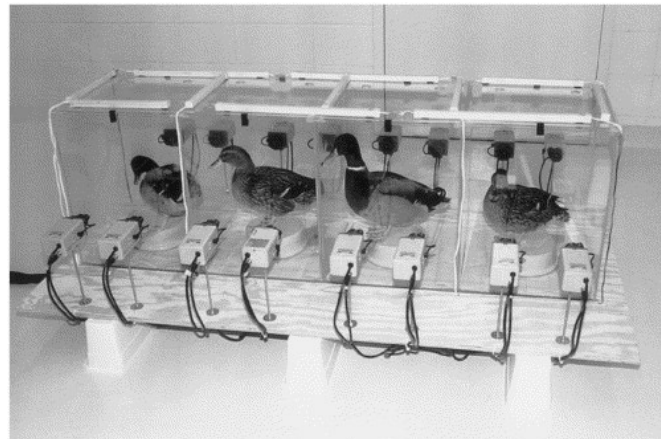
- **Question:** A) Can heart rate predict metabolic rate on single dives only?
- Linear-mixed effects model (LME) treats animal as a random effect, so **I don't plot symbols by animal, but rather treatment type**
- This was an appendix figure in Thesis, but not in publication



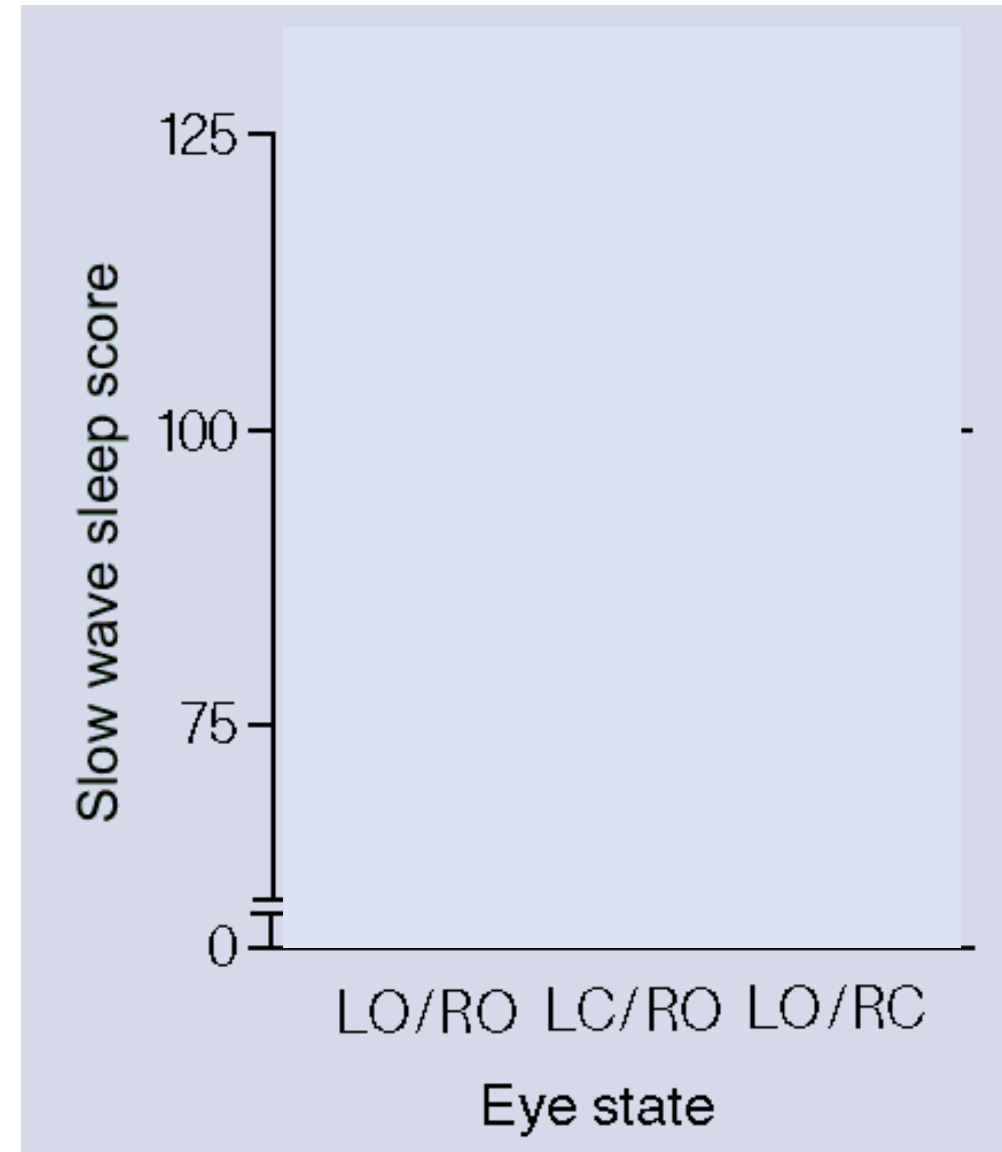
**Figure A3.5** The relationship between heart rate ( $fh$ ) averaged over a dive cycle and average metabolic rate ( $V\&O_2 = AMR$ ) for Steller sea lions over a single dive (A) was not significantly different than when animals were resting in water (B). See Table 3.1 for Animal ID to corresponding animal number.

# Example

- Birds sleep in risky situations by sleeping with 1 eye open and one hemisphere of the brain awake
  - Unihemispheric slow-wave sleep
- Some mammals do this too → whales, dolphins, seals, sea lions



Slow wave sleep in the brain hemispheres of mallard ducks sleeping with one eye open.  
From Rattenborg et al. (1999) *Nature*.

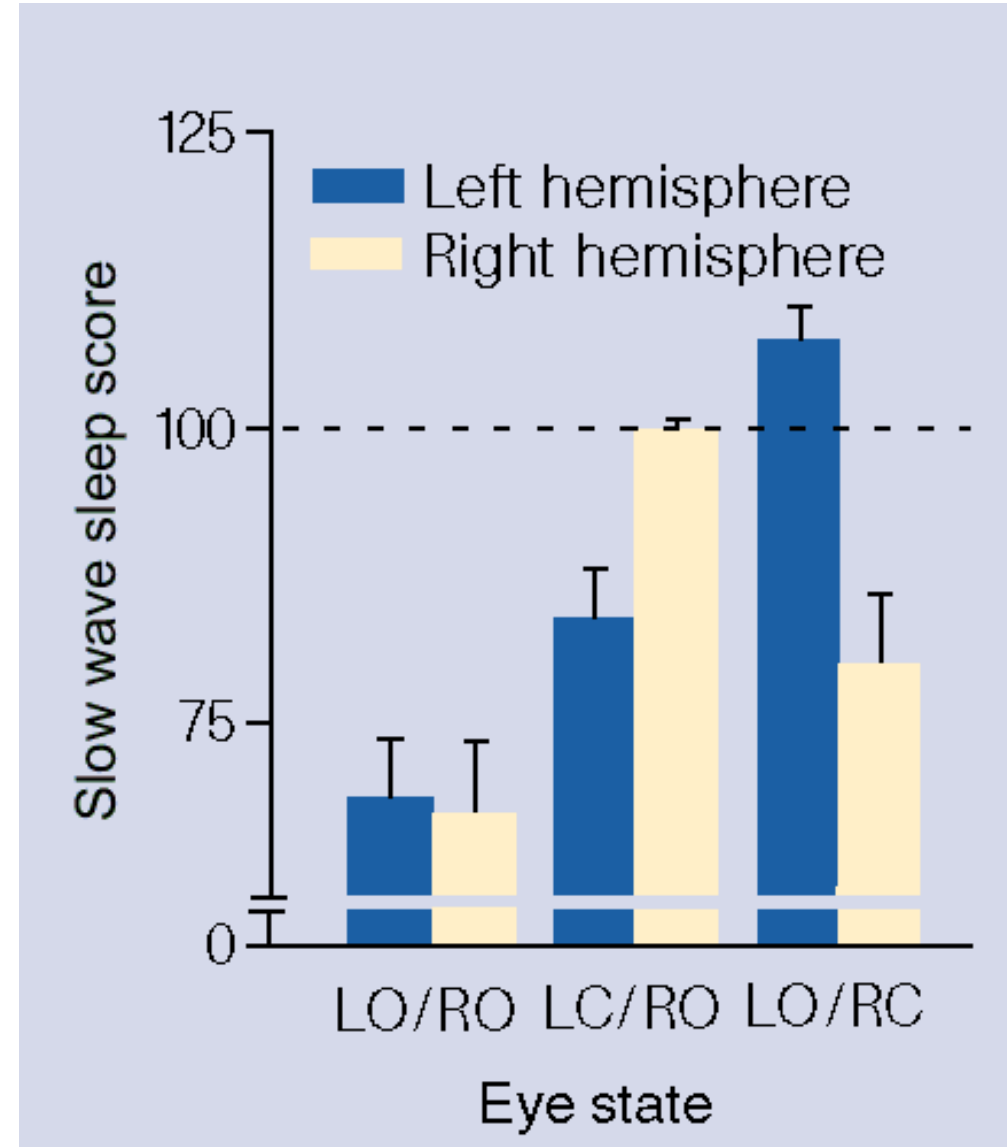


LO/RO=both eyes open  
LC/RO=Left closed/Right open  
LO/RC=Left open/Right closed

## 4. Represent magnitudes honestly

1. Are the bars “consistent with the numerical representation”?
2. Is 0 a reasonable baseline for evaluating sleep score?
3. Are there other issues with the graph?

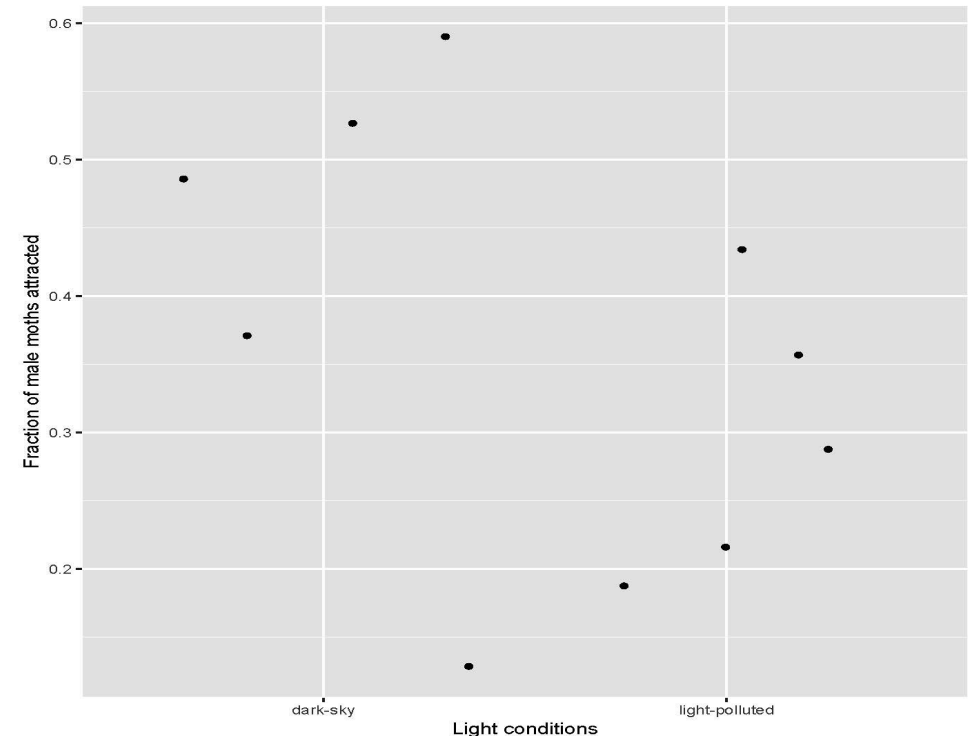
Slow wave sleep in the brain hemispheres of mallard ducks sleeping with one eye open.  
From Rattenborg et al. (1999) *Nature*.



## 4. Draw graphical elements clearly, minimize clutter

- *“Maximize the data-ink ratio, within reason”* – Tufte (1983)
- Originally journals charged for colour
- Modern take with online journals: How will this figure look printed black and white?
- Not a great use of negative space and solid grey background

Strip chart made with `ggplot()` using default graphical options.



Altermatt and Ebert (2016) measured light attraction of ermine moths (*Yponomeuta cagnagella*) from 10 different populations. Five of the populations were located in urban areas with plenty of human lights. The other 5 populations were located in pristine areas with no light pollution

**Default R plots are OK for exploration, but usually not polished enough to publish**

# 4 Principles to Increase Effectiveness of Graphs

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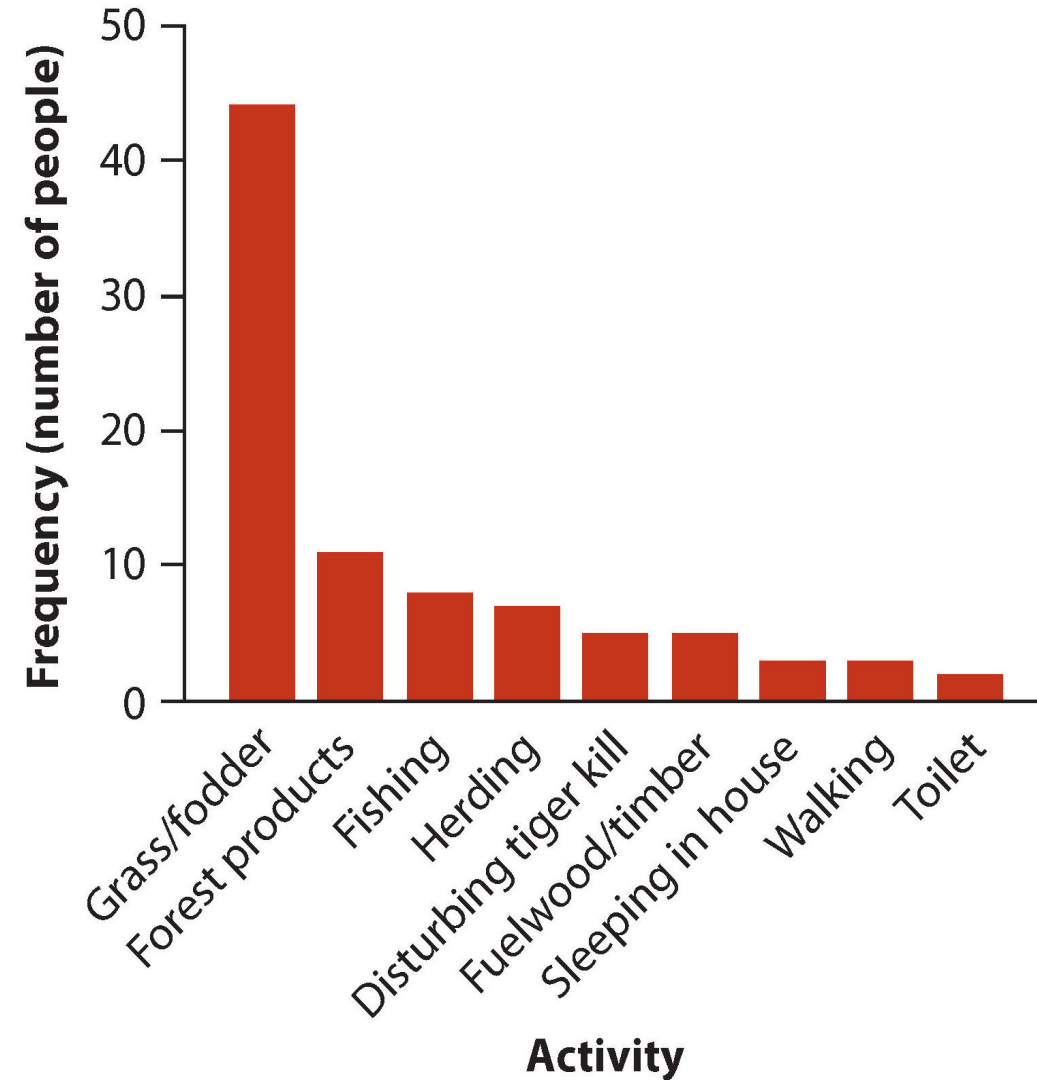


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# How to display category frequencies: Bar graph

- Height of bars displays frequency distribution of a categorical (grouping) variable
- Zero baseline
- Space between bars emphasize height
- Order of categories – most to least frequent is usually best
  - **R defaults order alphabetically**

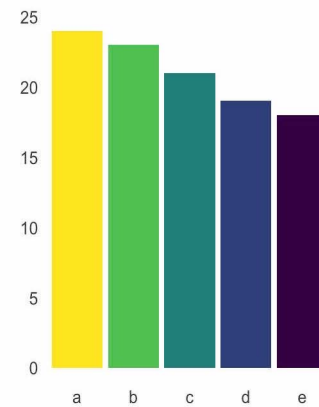
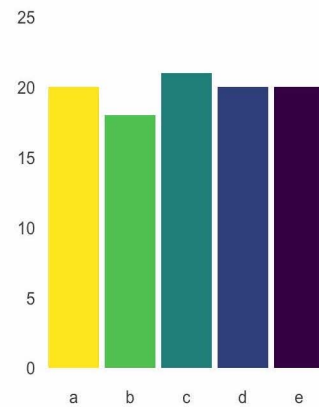
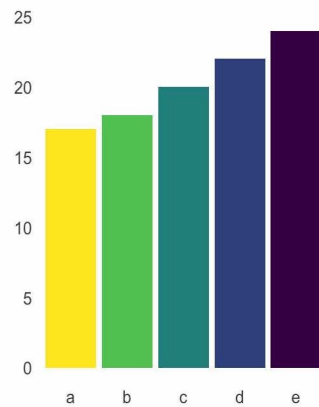
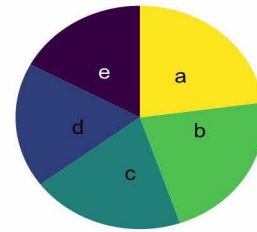
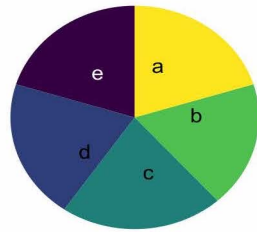
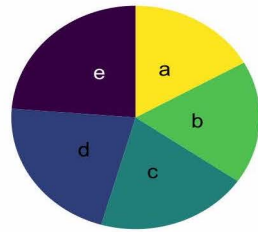


Activities of people at the time they were attacked and killed by tigers near Chitwan National Park, Nepal, between 1979 and 2006

# How to display category frequencies

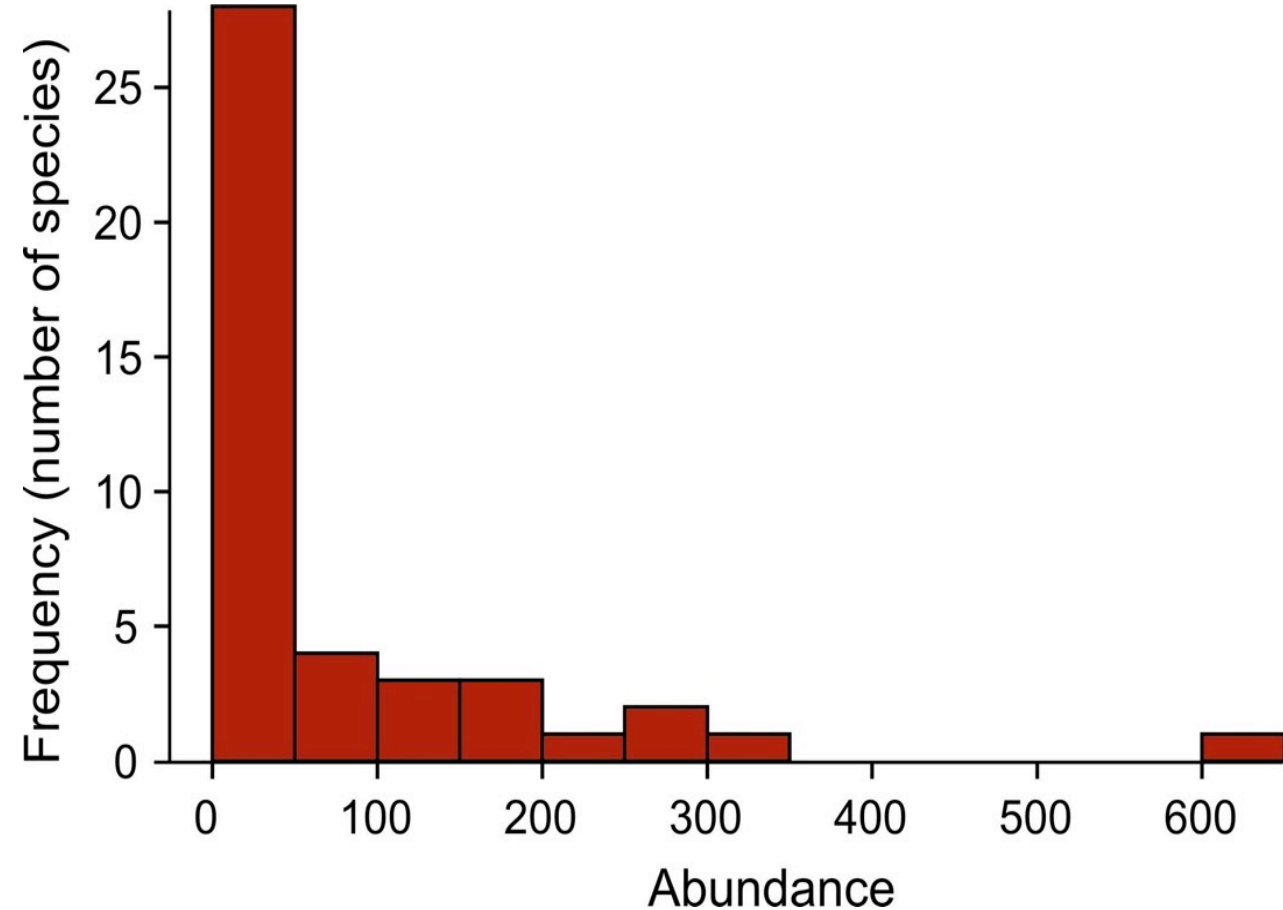
## Bar graph vs. pie chart: which is more successful?

- Humans are better at comparing relative areas in bar graphs than pie charts.



# How to display frequency distribution for numeric variable: Histogram

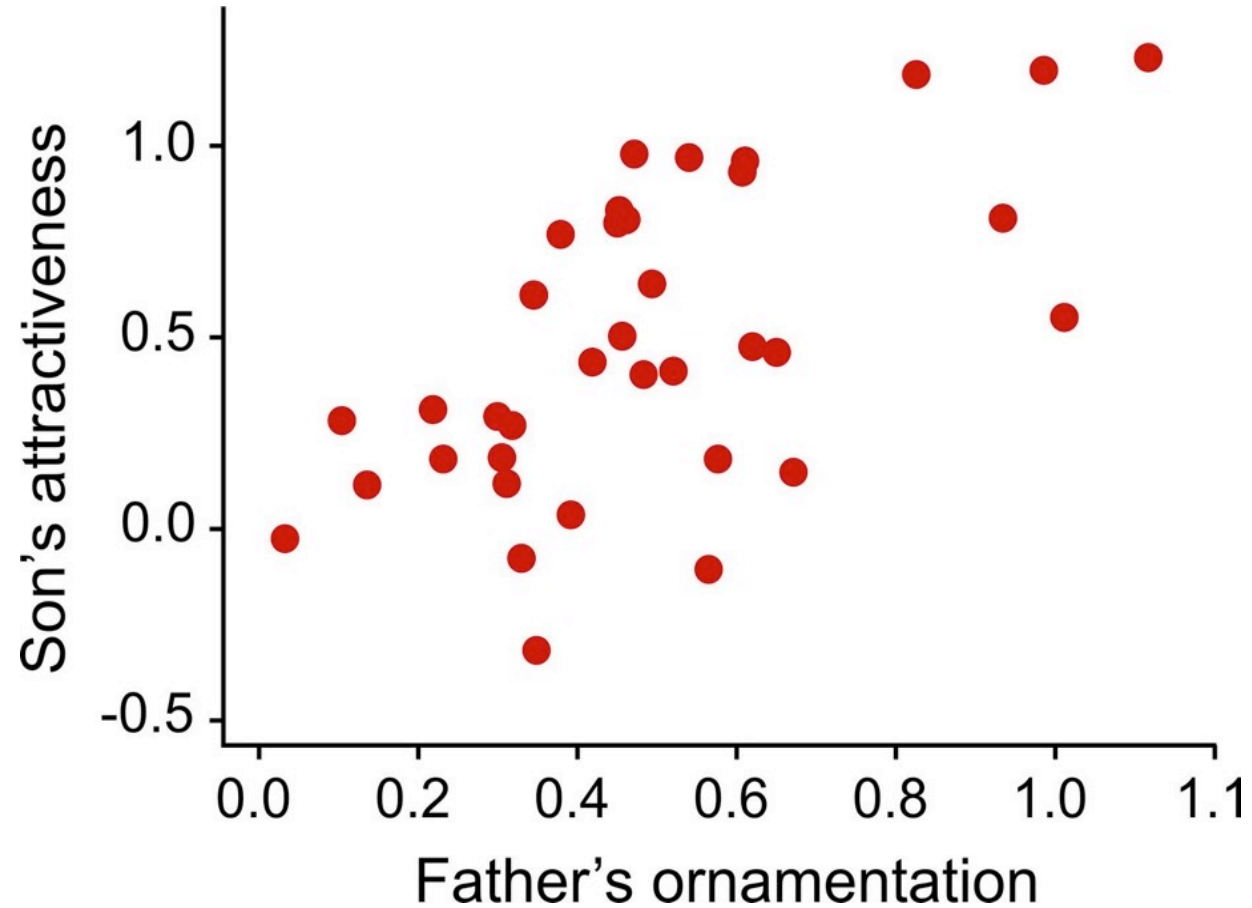
- Uses area of bars to display frequency distribution of a numerical variable
- Zero baseline
- No spaces between bars
- Choice of number of bins and bin width



# How to display association between 2 numerical variables: Scatterplot

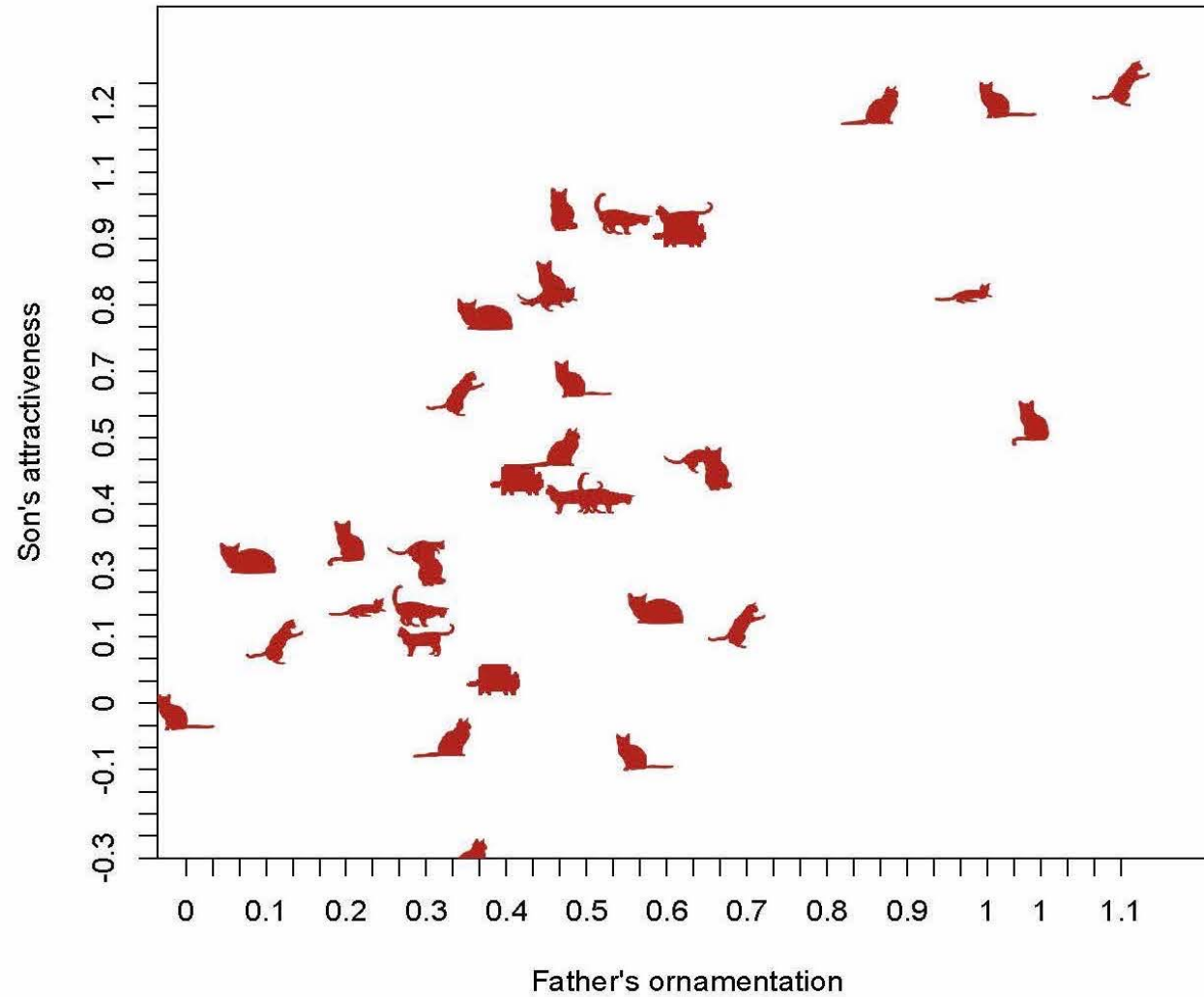
- Non-zero baseline often ok
- Goal is to show association, not height above 0
- Points fill the space available

**If you use base R plot(), R will just plot the data by it's best guess of plot type (based on the class of your variables)**



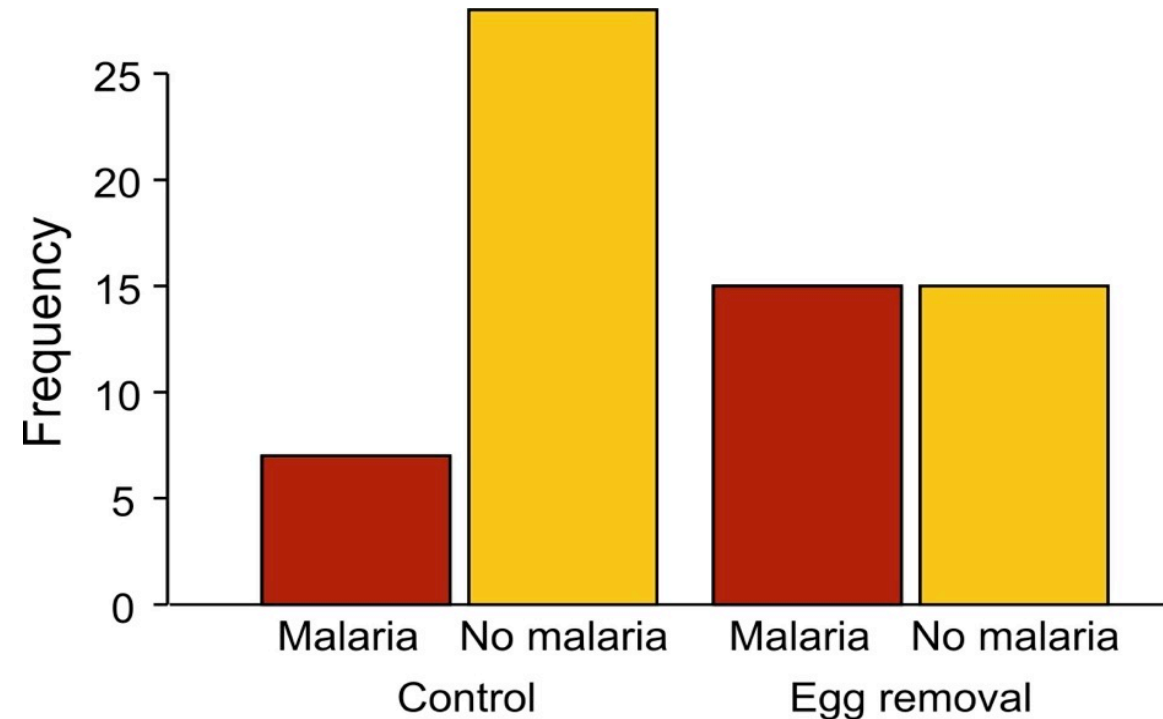
The relationship between the ornamentation of male guppies and the average attractiveness of their sons.  $n = 36$  families.

# A catterplot



# How to display association between categorical variables: **Grouped bar graph**

- Uses height of bars to display association between two (or more) categorical variables.
- Explanatory variable = outer groups; response variable = inner groups
- Zero baseline (so that height is proportional to frequency)
- Spacing between bars wider between outer groups



Incidence of malaria in female great tits in relation to experimental treatment.  $n = 65$  birds.

# How to display association between categorical variables: **Mosaic plot**

- Uses area of rectangles to display association between two (or more) categorical variables
- Explanatory variable along horizontal axis; response variable stacked
- Area proportional to frequency
- Like a graphical representation of a contingency table



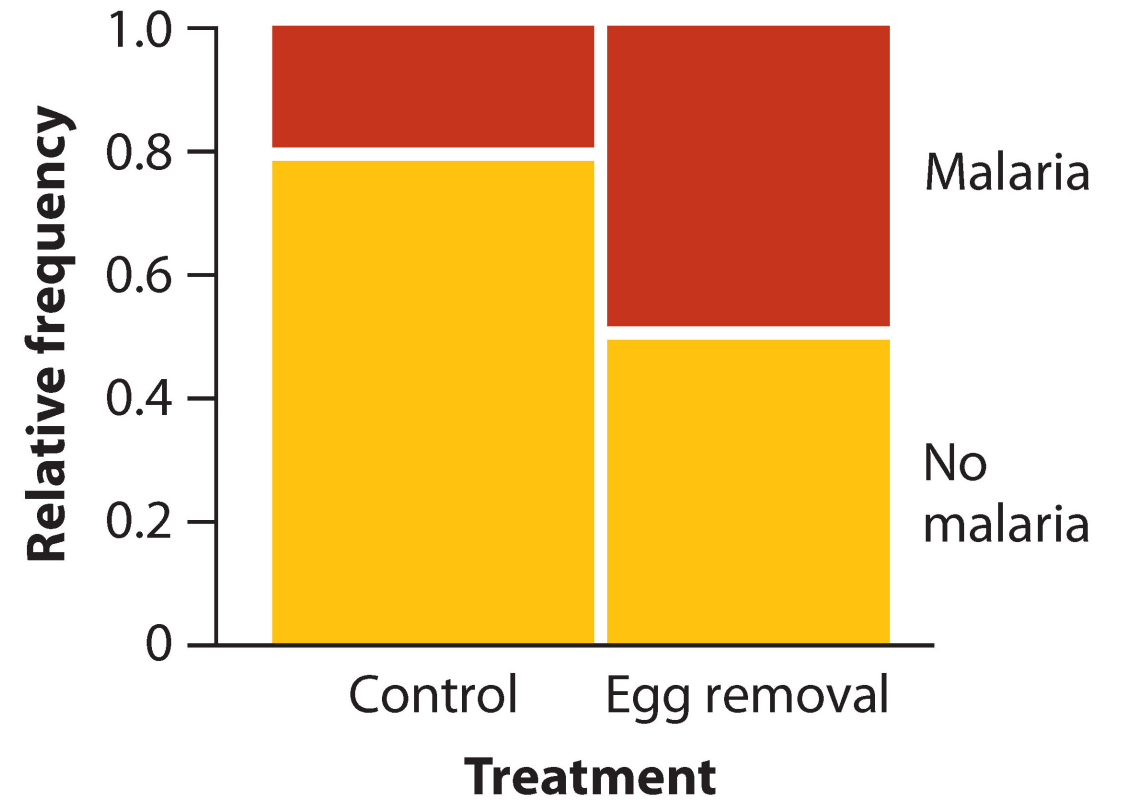
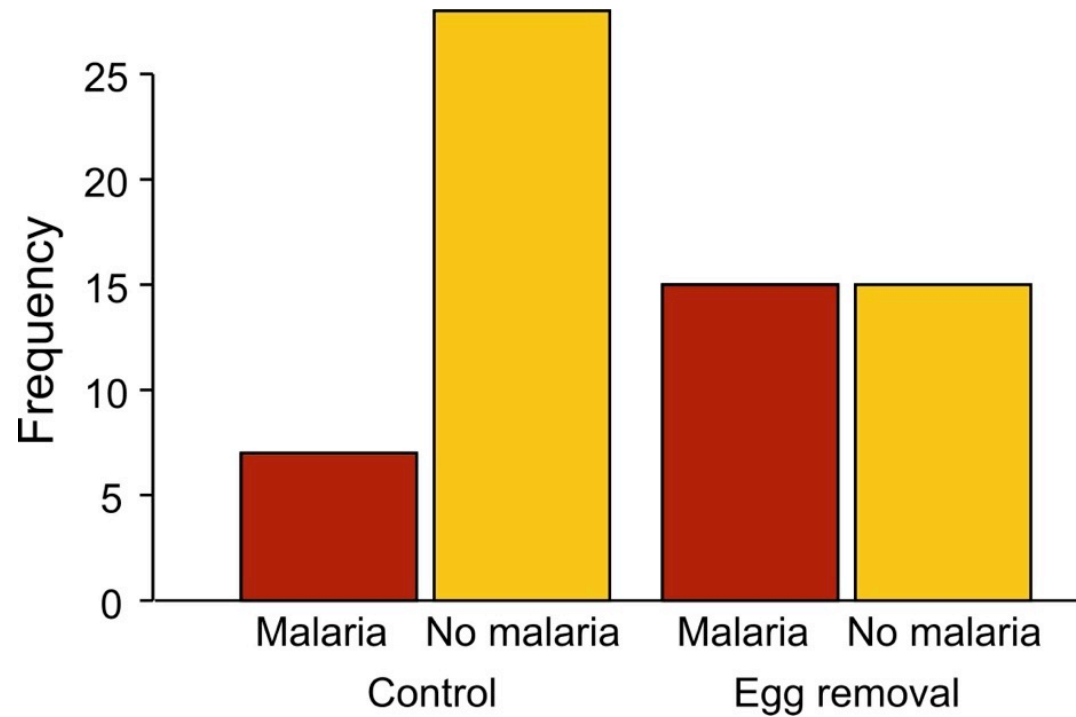
Incidence of malaria in female great tits in relation to experimental treatment.

$n = 65$  birds.



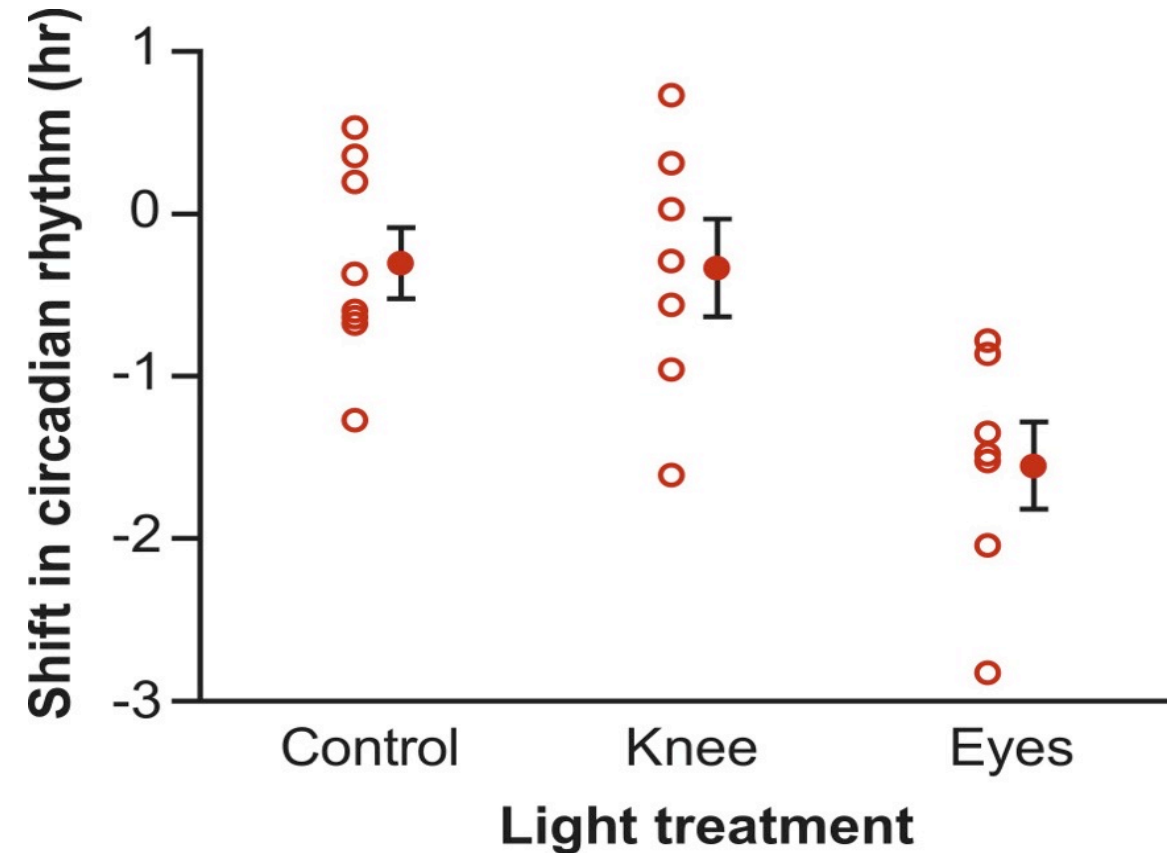
# Grouped bar graph vs. mosaic plot

## Which is more successful?



# How to display association between numerical and categorical variables? **Strip chart**

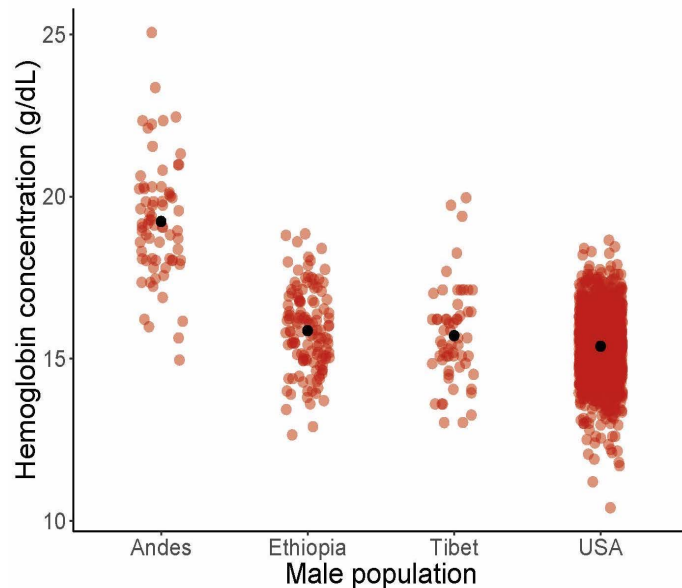
- Displays **differences between groups**
- Shows the data points
- Non-zero baseline often ok  
(goal is association not magnitude or frequency)
- Points fill the space available



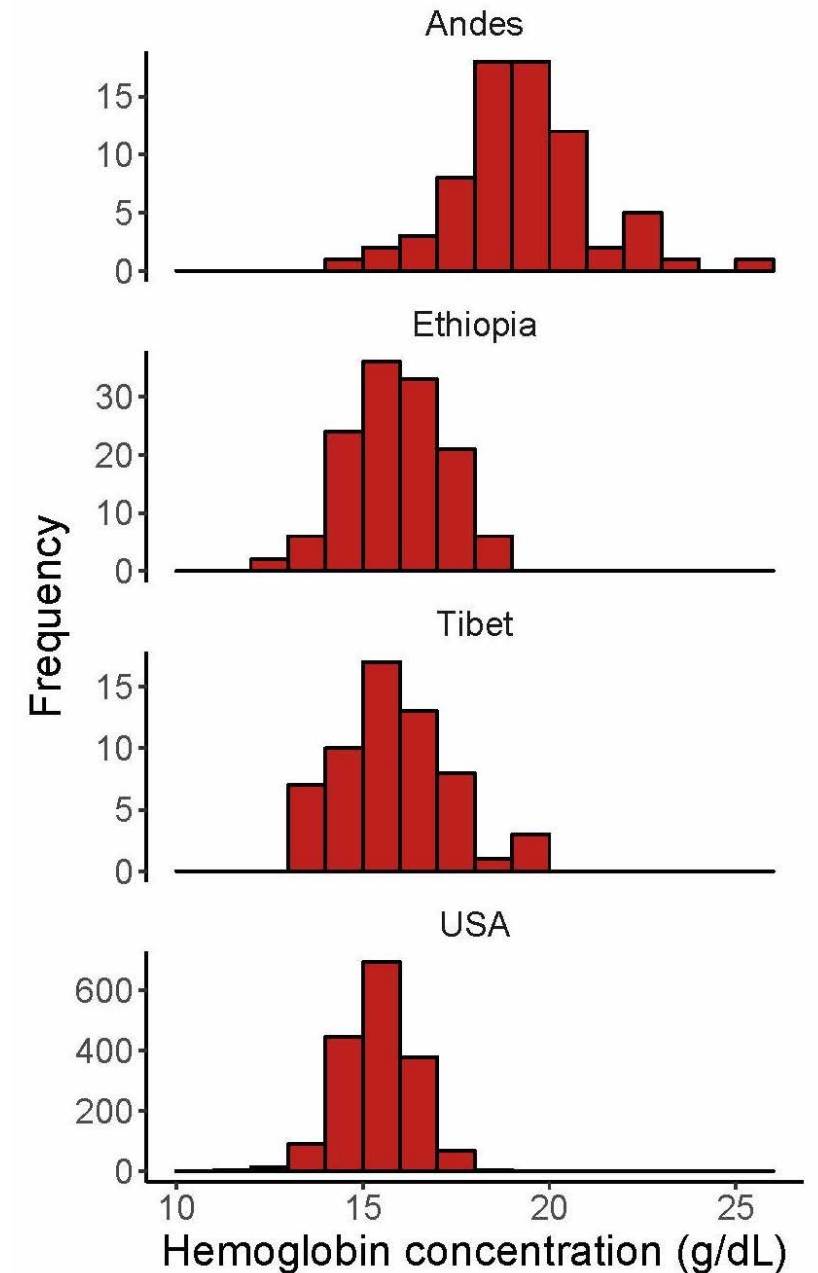
Phase shift in the circadian rhythm of melatonin production in 22 subjects given alternative light treatments (open circles). Group means  $\pm$  1 SE also shown.

# How to display association between numerical and categorical variables?

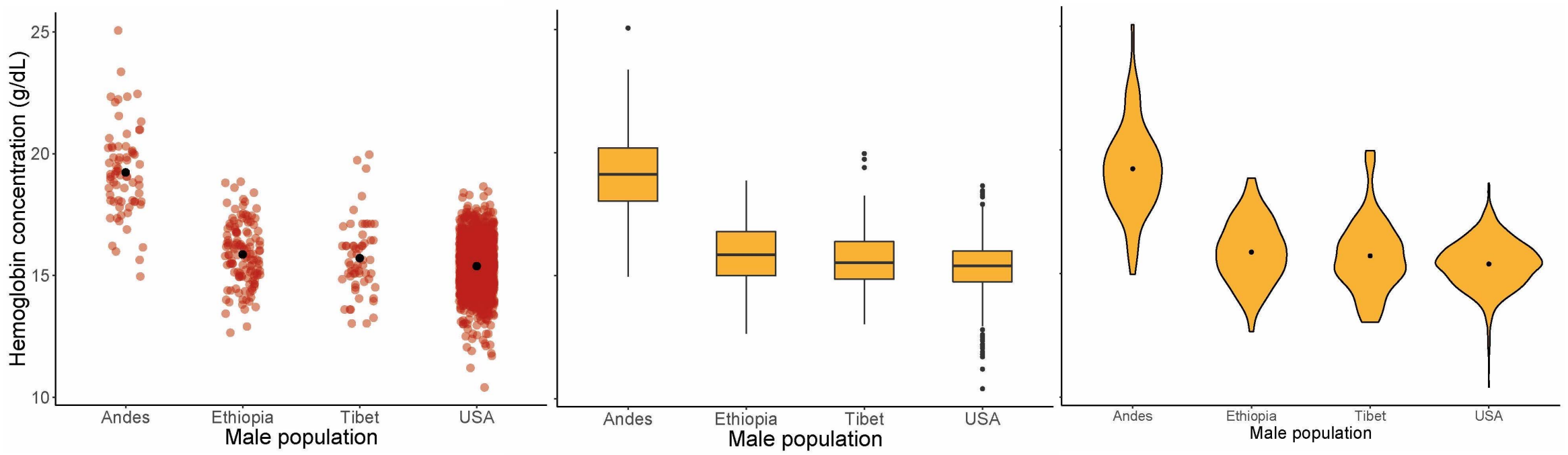
- Strip chart vs multiple histograms
- Too many data points for a strip chart.
- Stack histograms vertically to best compare



Hemoglobin concentration in blood of males living at high elevation compared to sea-level USA control



# Discussion: Which is more successful at conveying a pattern? Why?



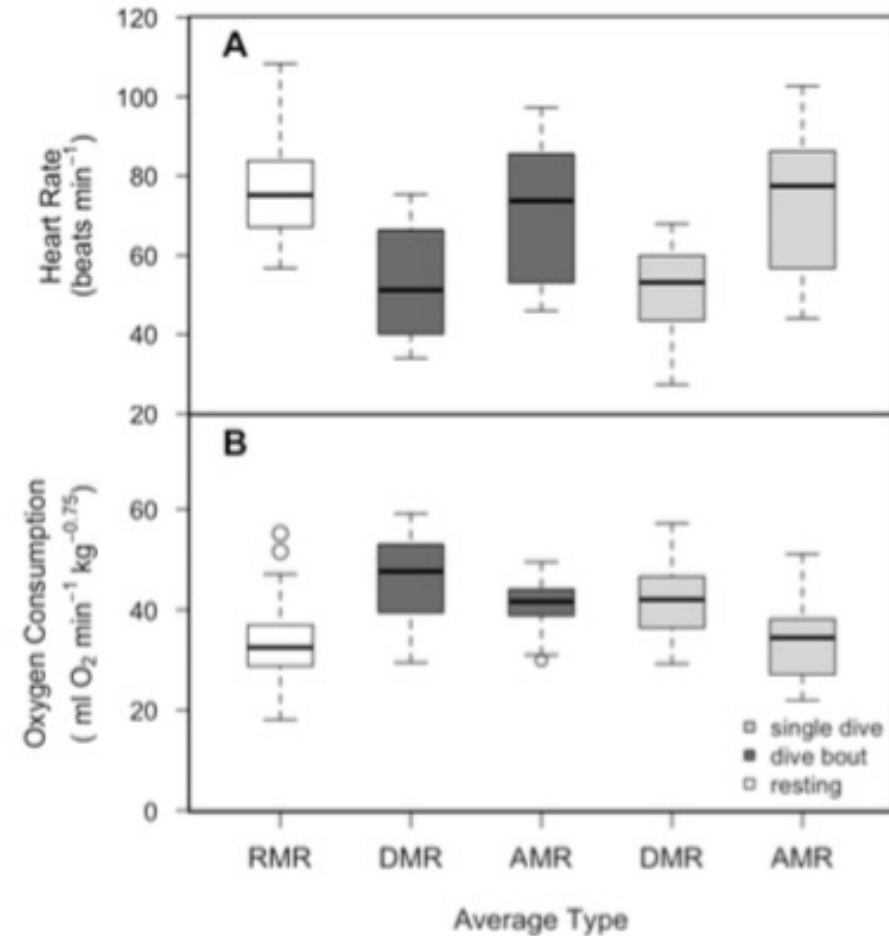
Strip chart

Box plot

Violin plot

# How to display association between numerical and categorical variables? **Box plot**

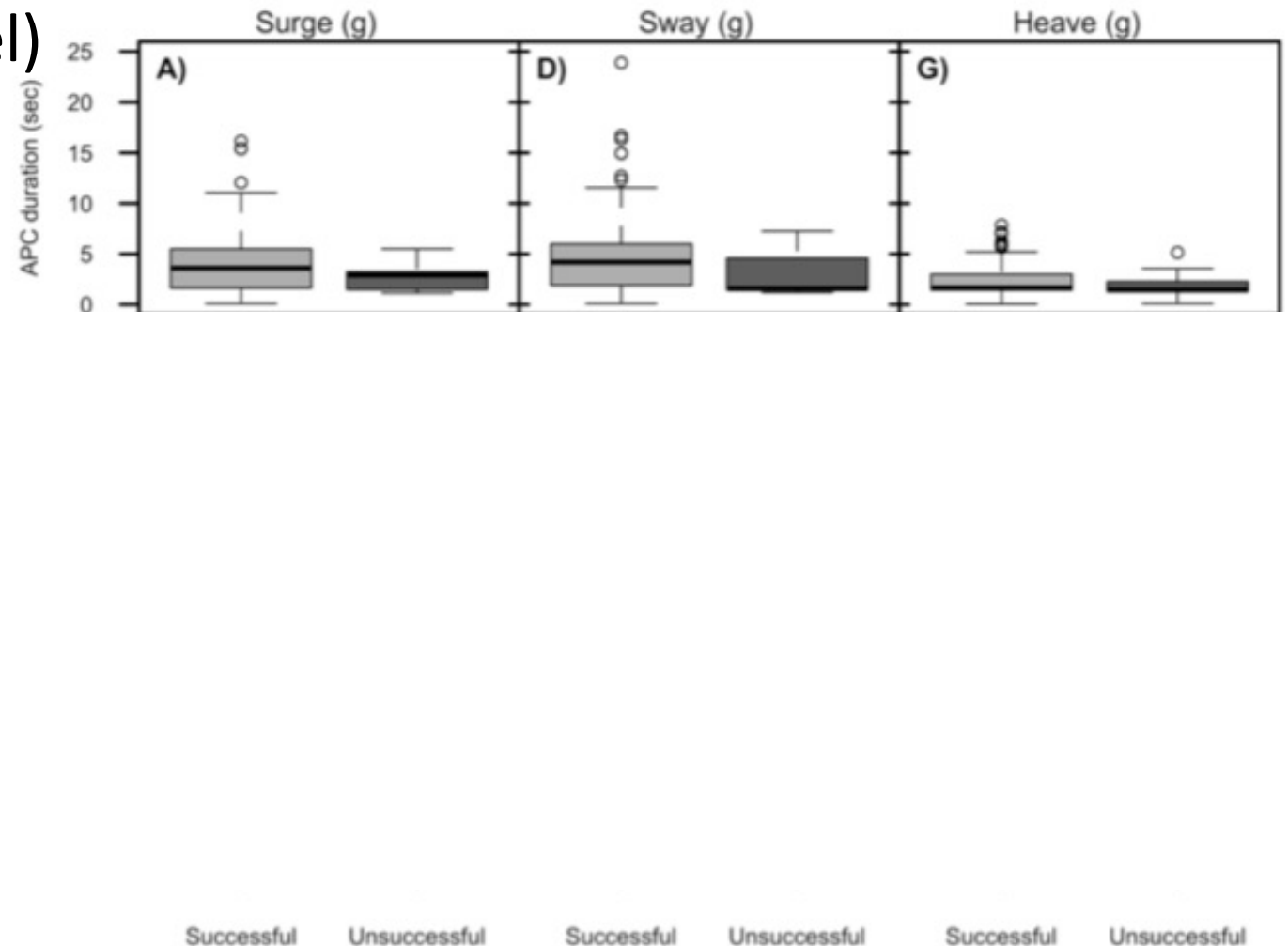
- I use these a lot with LME models that are not predictive
- You can change order of X variable (default is alphabetical)
- **Question:** Does Y variable differ among the X variable categories?
  - **NOT** predicting Y variable, just asking if different



## Boxplot example (with LME model)

Does the duration of an attempted prey capture (APC) differ between successful vs. unsuccessful attempts?

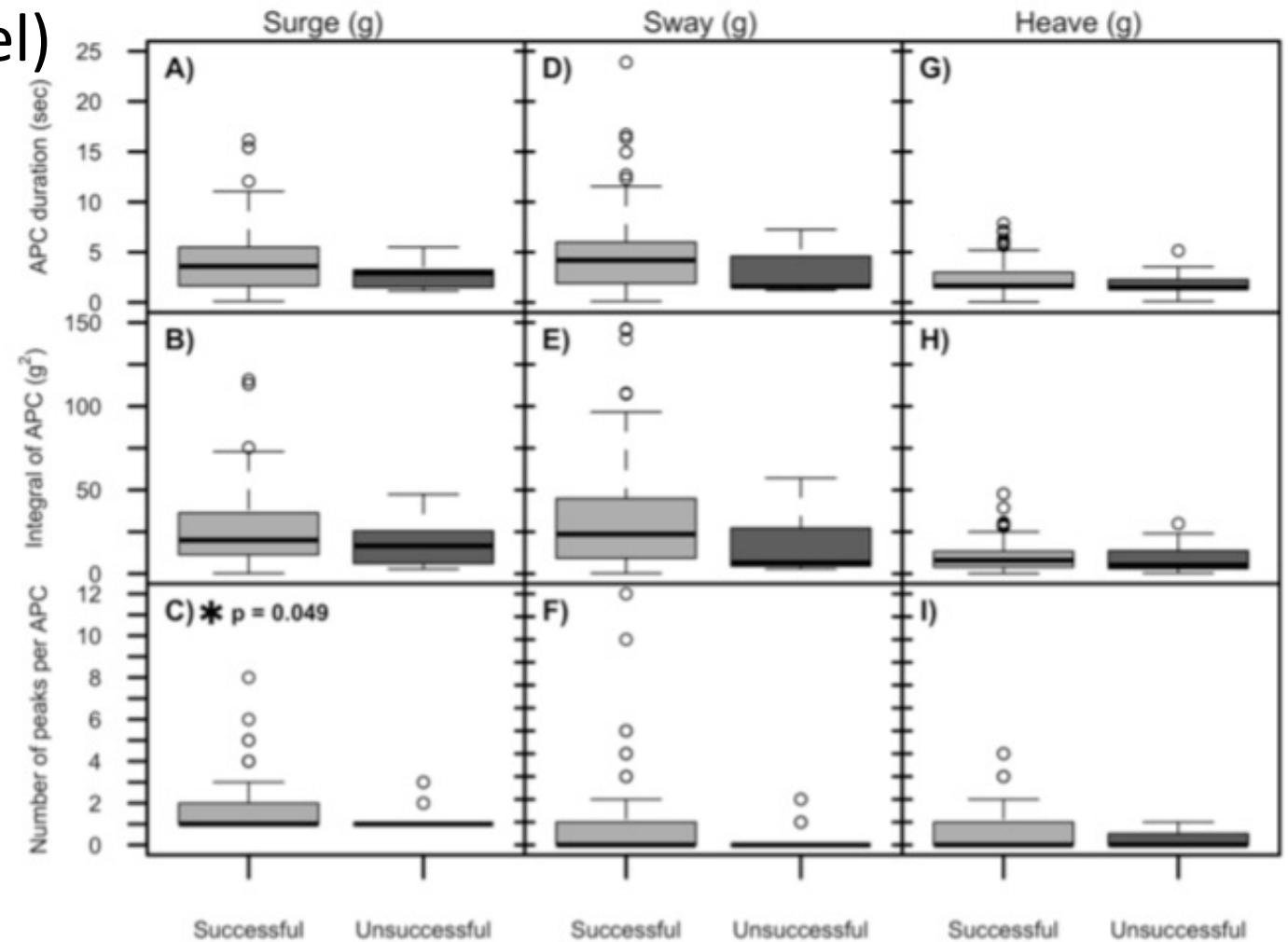
Is this different if I look at each type of acceleration??



**Fig 5. Analysis of successful prey captures by accelerometers.** The accelerometer was not able to distinguish between successful and unsuccessful attempted prey captures (APC) on any axis of acceleration. Successful events included video where prey was captured, and unsuccessful events included events where the fur seal chased and missed an attempted capture attempt of the prey on video. Neither mean event duration (A, D, G), or integral area under the APC (B, E, H) significantly varied between successful or unsuccessful APC. The number of peaks in surge marginally differed between types of APC (C), but not for sway (F) or heave (I). Results are from Function 1 optimized with generic parameters (0.1 variance and 5 s). Each data point represents an individual APC.

## Boxplot example (with LME model)

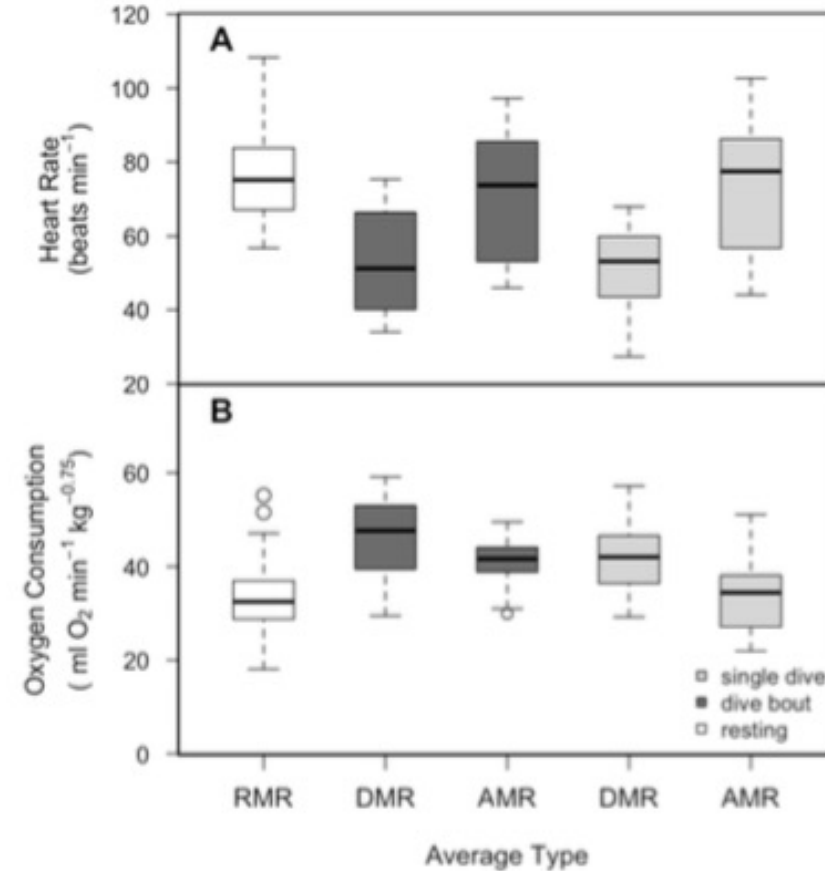
- Also looked at other acceleration metrics



**Fig 5. Analysis of successful prey captures by accelerometers.** The accelerometer was not able to distinguish between successful and unsuccessful attempted prey captures (APC) on any axis of acceleration. Successful events included video where prey was captured, and unsuccessful events included events where the fur seal chased and missed an attempted capture attempt of the prey on video. Neither mean event duration (A, D, G), or integral area under the APC (B, E, H) significantly varied between successful or unsuccessful APC. The number of peaks in surge marginally differed between types of APC (C), but not for sway (F) or heave (I). Results are from Function 1 optimized with generic parameters (0.1 variance and 5 s). Each data point represents an individual APC.

# Linear mixed-effects model: not predictive

- Numerical Y, **Categorical X**
- Animal random factor
- Fixed factor
- N=7 Steller sea lions
  
- Goal: **Is there a difference in metabolic rate among treatments (categorical X)**

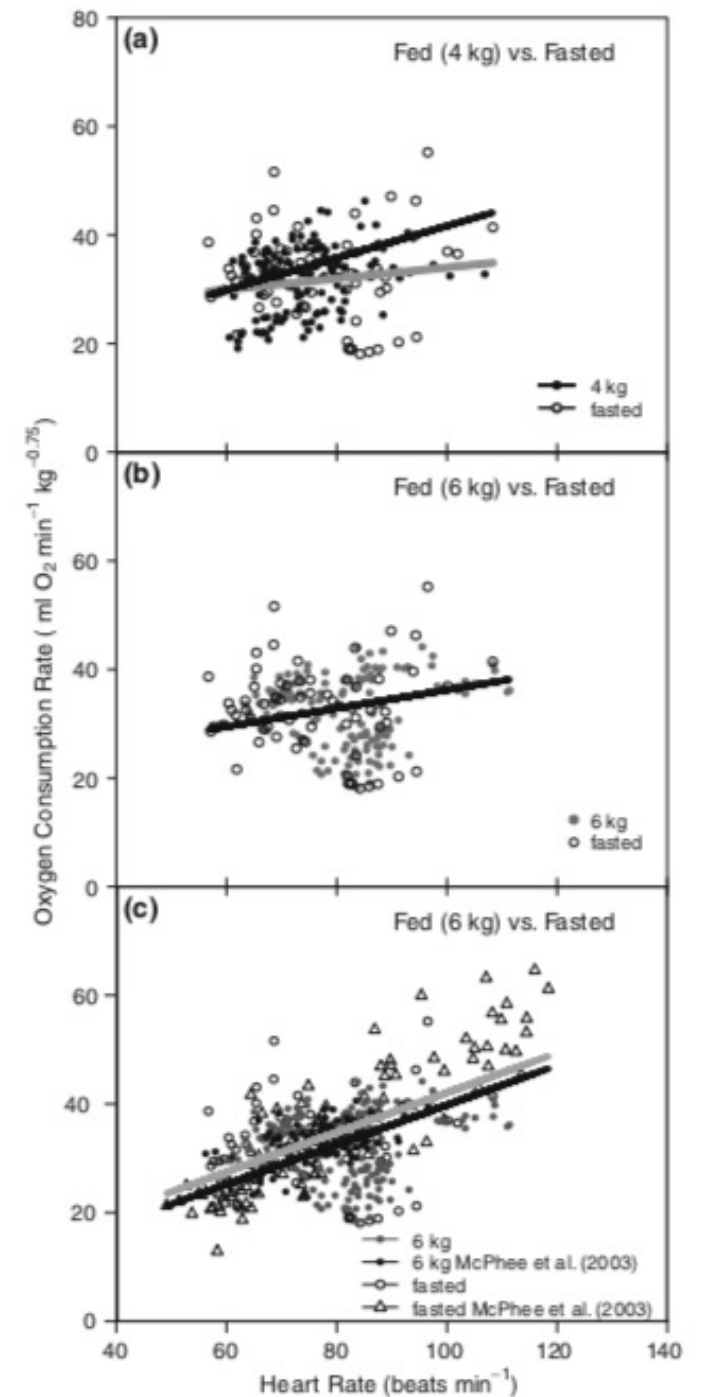


**Figure A3. 2** Box-and-whiskers plot showing minimum, median, and maximum heart rate ( $f_h$ ) and oxygen consumption  $\dot{V}_{O_2}$  for animals resting in water (white), averaged over dive bouts (dark grey), or averaged over single dives (light grey). Water composite includes data when animals were resting in a swim mill and resting at the surface of open water. Diving metabolic rate (DMR) includes data averaged over single dives or several dives in dive bout. Average metabolic rate (AMR) includes data averaged over a complete dive cycle (dive + surface interval) or entire dive bout (multiple dives and surface intervals).



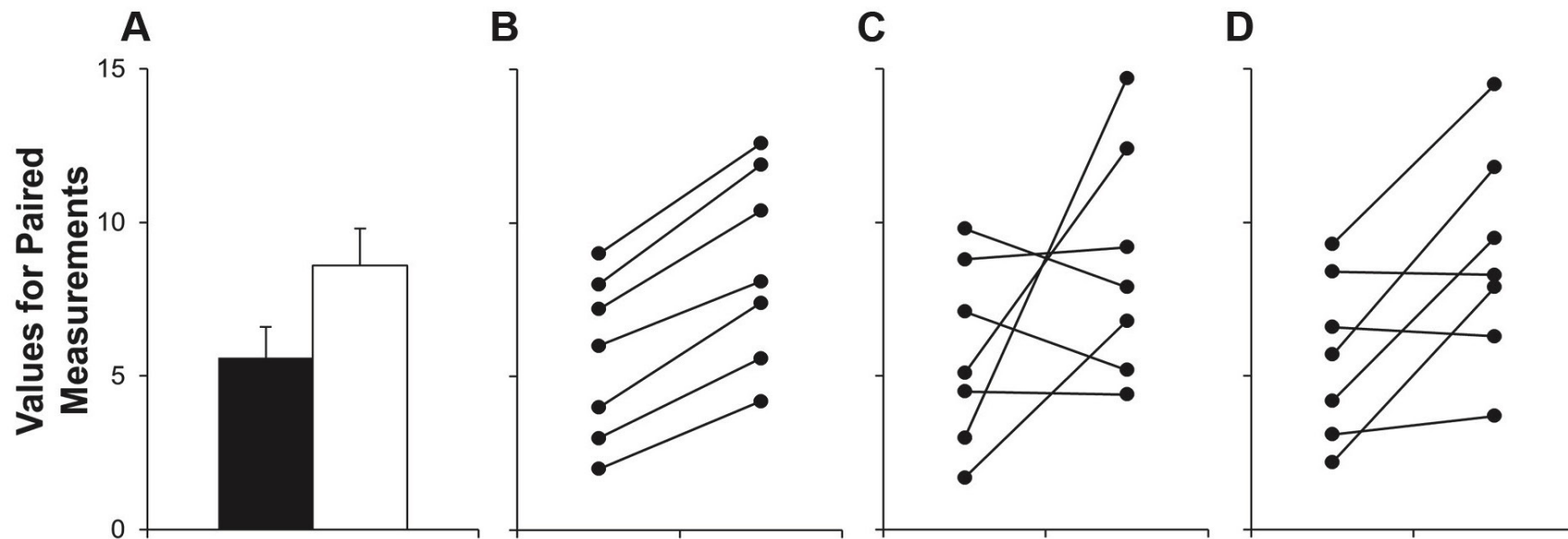
# Linear mixed-effects model: predictive

- Numerical X and Y
- Animal random factor
- Fixed factor –3 level treatment
  - Fasted
  - Fed 4 kg
  - Fed 6 kg
- N=7 Steller sea lions
- Goal: **predictive** equation for metabolic rate



# How to show repeated measures data

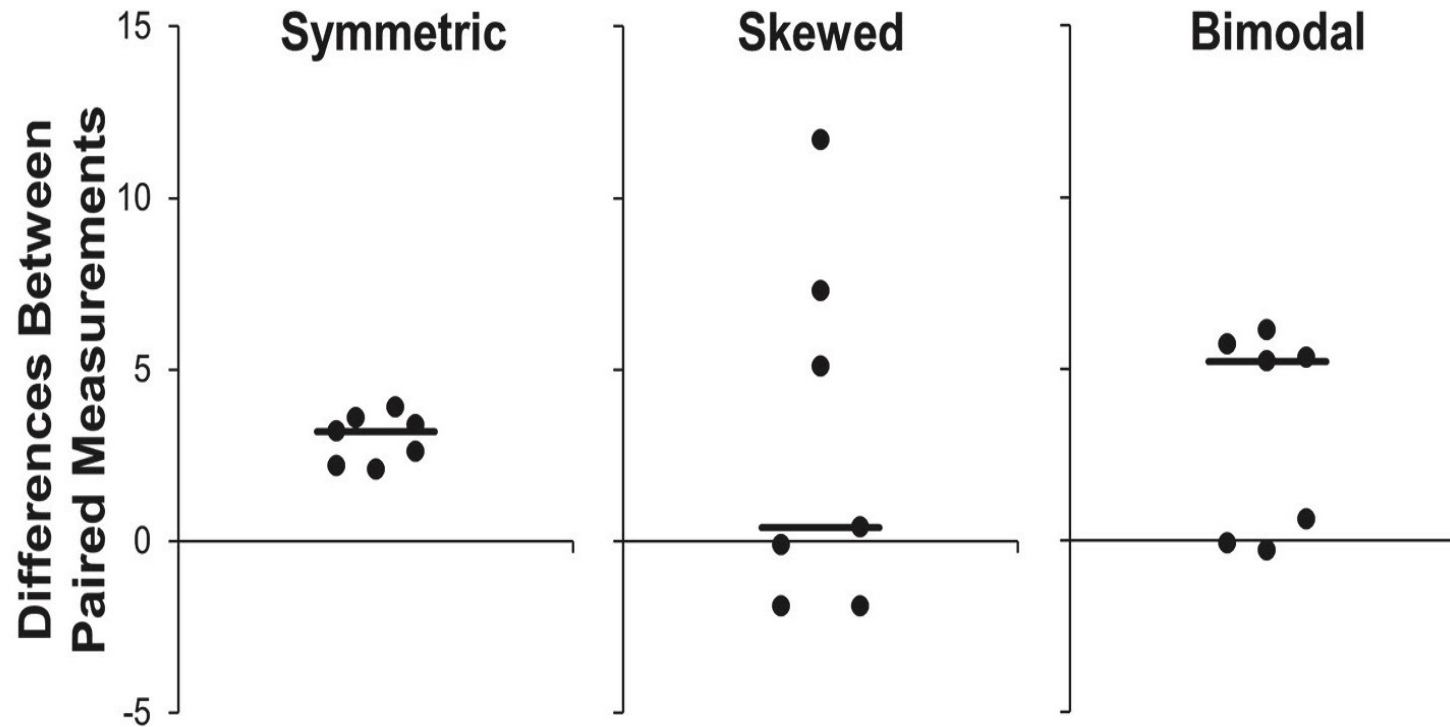
## Interaction plots



I use this more for exploratory, early LME plots than final plots in published papers

# How to show repeated measures data

## Strip charts of differences



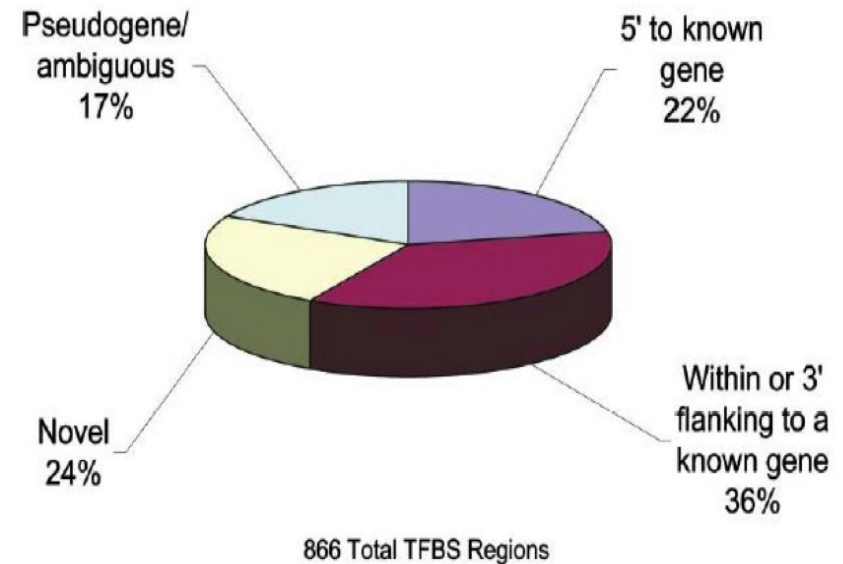
Weissgerber et al. (2015) beyond bar and line graphs: time for a new data presentation paradigm. *PLoS Biol.* DOI:10.1371/journal.pbio.1002128

# Avoid 3D graphs to display frequencies

- 3D makes it more difficult to see area, and hence the frequencies
- A graph that is meaningful only with numbers added is a failure to communicate clearly and effectively

**BUT, I often add numbers above bar plots, histograms etc when showing supervisor or presentations (when reader doesn't actually have paper in front of them)**

Distribution of All TFBS Regions



# Design graphs to show important trends not plot all of your raw data

**Instead:** Make use of supplemental material of thesis appendix

Bell ML, et al. (2007) Spatial and temporal variation in PM<sub>2.5</sub> chemical composition in the United States for health effects studies. Environmental Health Perspectives 115:989-995

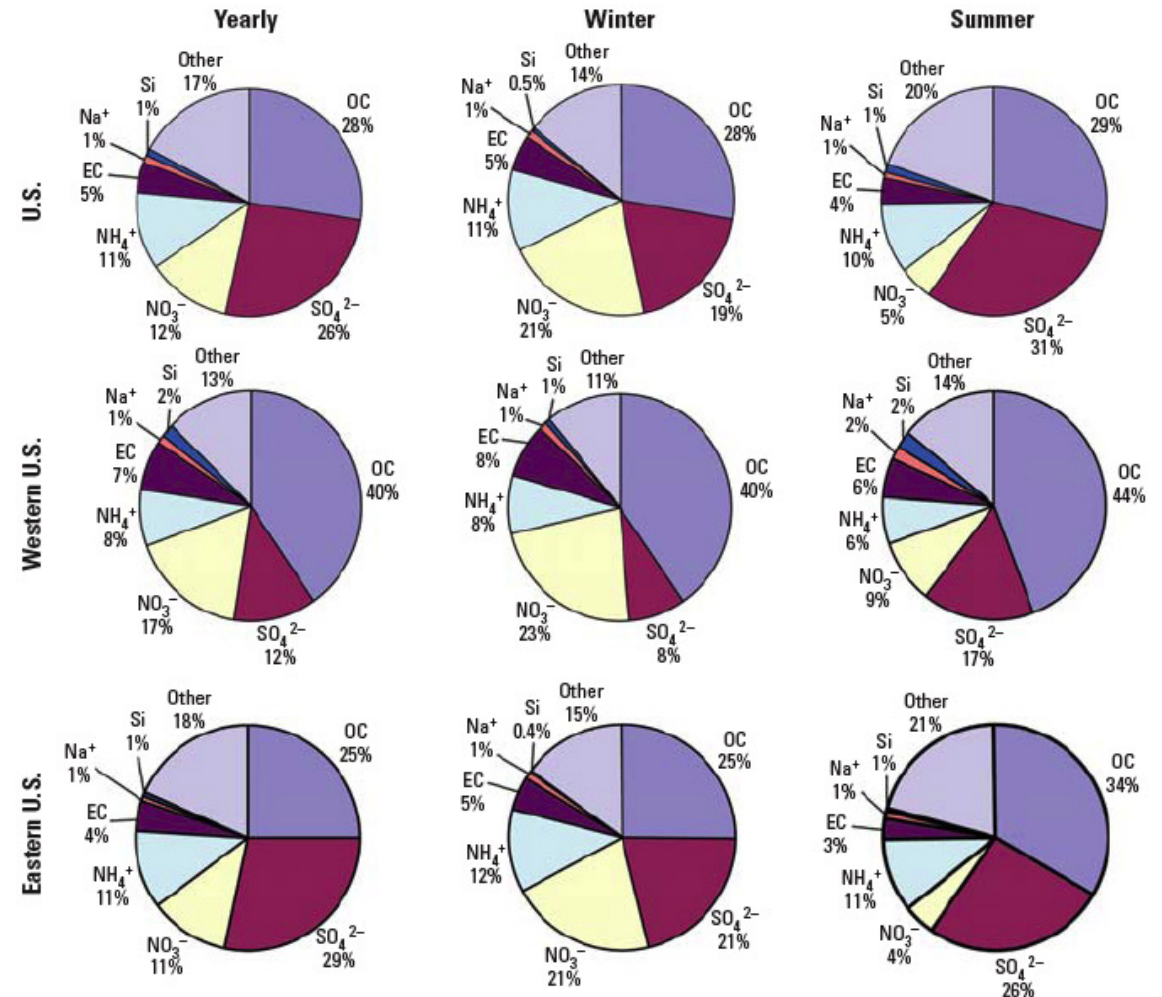


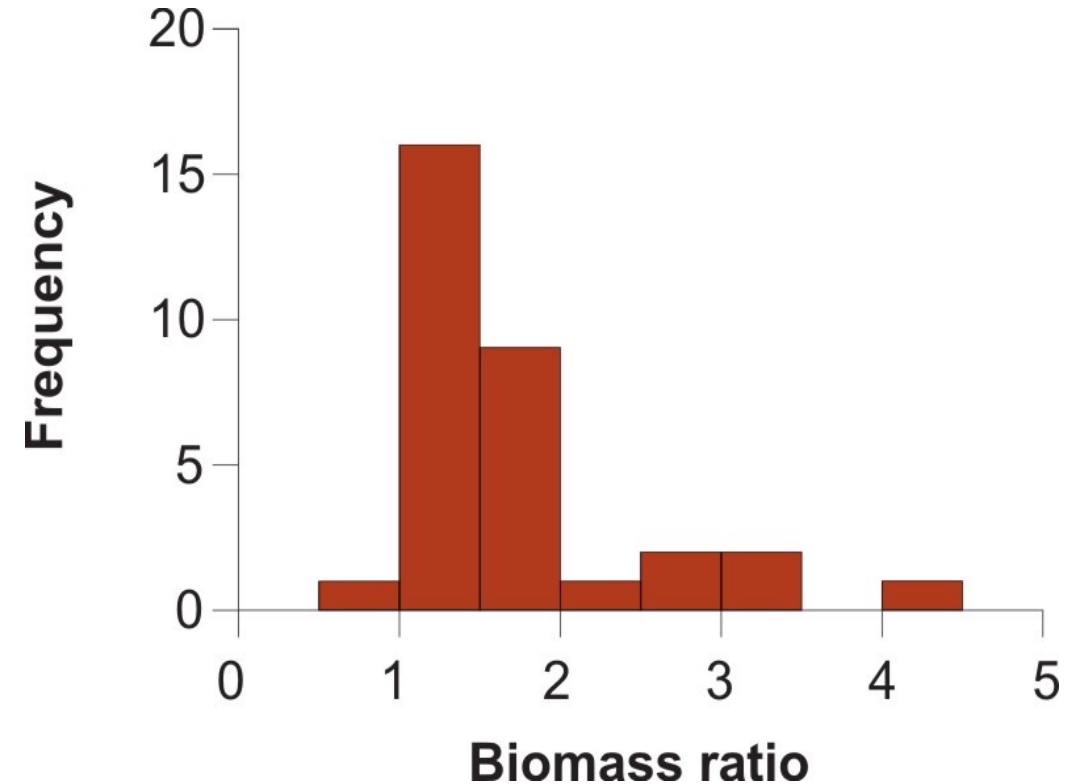
Figure 3. Percent of PM<sub>2.5</sub> composition by component for yearly, winter, and summer averages, by region.

# What was the point of 9 pie charts?

- The main patterns of interest were the change in  $\text{NO}_3$  and  $\text{SO}_4$  between winter and summer, and the consistency of change between geographic regions. This is **not** easy to see (“Huh?” not “Oh!”)
- **Try it later:** Design a graph to show the change from summer to winter in  $\text{NO}_3$  and  $\text{SO}_4$ , rather than try to display everything.

# Ratio Data (any proportion or %)

- **Problems:** Ratios less than 1 are sandwiched between 0 and 1, distorting magnitudes
- **Possible solutions:** Use log of ratio so that ratios above and below 1 have the same scale.



Biomass ratio is the total mass of all marine plants and animals per unit area of marine reserve divided by the same quantity in the unprotected control. N = 32 pairs (reserve and control). Data from Halpern (2003)

# Choosing colours for graphs

- Colour blindness (don't mix red and green)
- Publication costs for colour? (minimal online now)
- Most people still print articles in B&W so if use colour make sure it's readable in B/W greyscale
- **Keep colours and symbols for same treatment consistent throughout a manuscript**



# R Workshop Tips

## Point type cheat sheet (pch) for Base R

### R color cheatsheet

Finding a good color scheme for presenting data can be challenging. This color cheatsheet will help!

#### R uses hexadecimal to represent colors

Hexadecimal is a base-16 number system used to describe color. Red, green, and blue are each represented by two characters (#rrggbb). Each character has 16 possible symbols: 0,1,2,3,4,5,6,7,8,9,A,B,C,D,E,F:

"00" can be interpreted as 0.0 and "FF" as 1.0  
i.e., red= #FF0000, black=#000000, white = #FFFFFF

Two additional characters (with the same scale) can be added to the end to describe transparency (#rrggbbaa)

#### R has 657 built in color names

To see a list of names: `colors()`

Example:  
`peachpuff4`

These colors are displayed on P. 3.

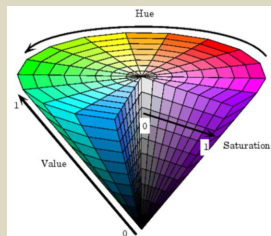
#### R translates various color models to hex, e.g.:

- RGB (red, green, blue): The default intensity scale in R ranges from 0-1; but another commonly used scale is 0-255. This is obtained in R using `maxColorValue=255`. `alpha` is an optional argument for transparency, with the same intensity scale.  
`rgb(r, g, b, maxColorValue=255, alpha=255)`
- HSV (hue, saturation, value): values range from 0-1, with optional alpha argument  
`hsv(h, s, v, alpha)`
- HCL (hue, chroma, luminance): hue describes the color and ranges from 0-360; 0 = red, 120 = green, blue = 240, etc. Range of chroma and luminance depend on hue and each other  
`hcl(h, c, l, alpha)`

#### A few notes on HSV/HCL

HSV is a better model for how humans perceive color. HCL can be thought of as a perceptually based version of the HSV model...blah blah blah...

Without delving into color theory: color schemes based on HSV/HCL models generally just look good.



#### R can translate colors to rgb (this is handy for matching colors in other programs)

`col2rgb(c("#FF0000", "blue"))`

### R Color Palettes

This is for all of you who don't know anything about color theory, and don't care but want some nice colors on your map or figure...NOW!

**TIP:** When it comes to selecting a color palette, **DO NOT** try to handpick individual colors! You will waste a lot of time and the result will probably not be all that great. R has some good packages for color palettes. Here are some of the options

#### Packages: grDevices and colorRamps

`grDevices` comes with the base installation and `colorRamps` must be installed. Each palette's function has an argument for the number of colors and transparency (`alpha`):

```
heat.colors(4, alpha=1)
> #FF0000FF "#FF8000FF" "#FFFF00FF" "#FFFFFF00FF"
```

For the `rainbow` palette you can also select start/end color (red = 0, yellow = 1/6, green = 2/6, cyan = 3/6, blue = 4/6 and magenta = 5/6) and saturation (`s`) and value (`v`):  
`rainbow(n, s=1, v=1, start=0, end=max(1, n-1)/n, alpha=1)`

#### Package: RcolorBrewer

This function has an argument for the number of colors and the color palette (see P. 4 for options).

```
brewer.pal(4, "Set3")
> "#8DD3C7" "#FFFFFFB3" "#BEBADA" "#FB8072"
```

To view colorbrewer palettes in R: `display.brewer.all(5)`  
There is also a very nice interactive viewer:  
<http://colorbrewer2.org/>

### ## My Recommendation ##

#### Package: colorspace

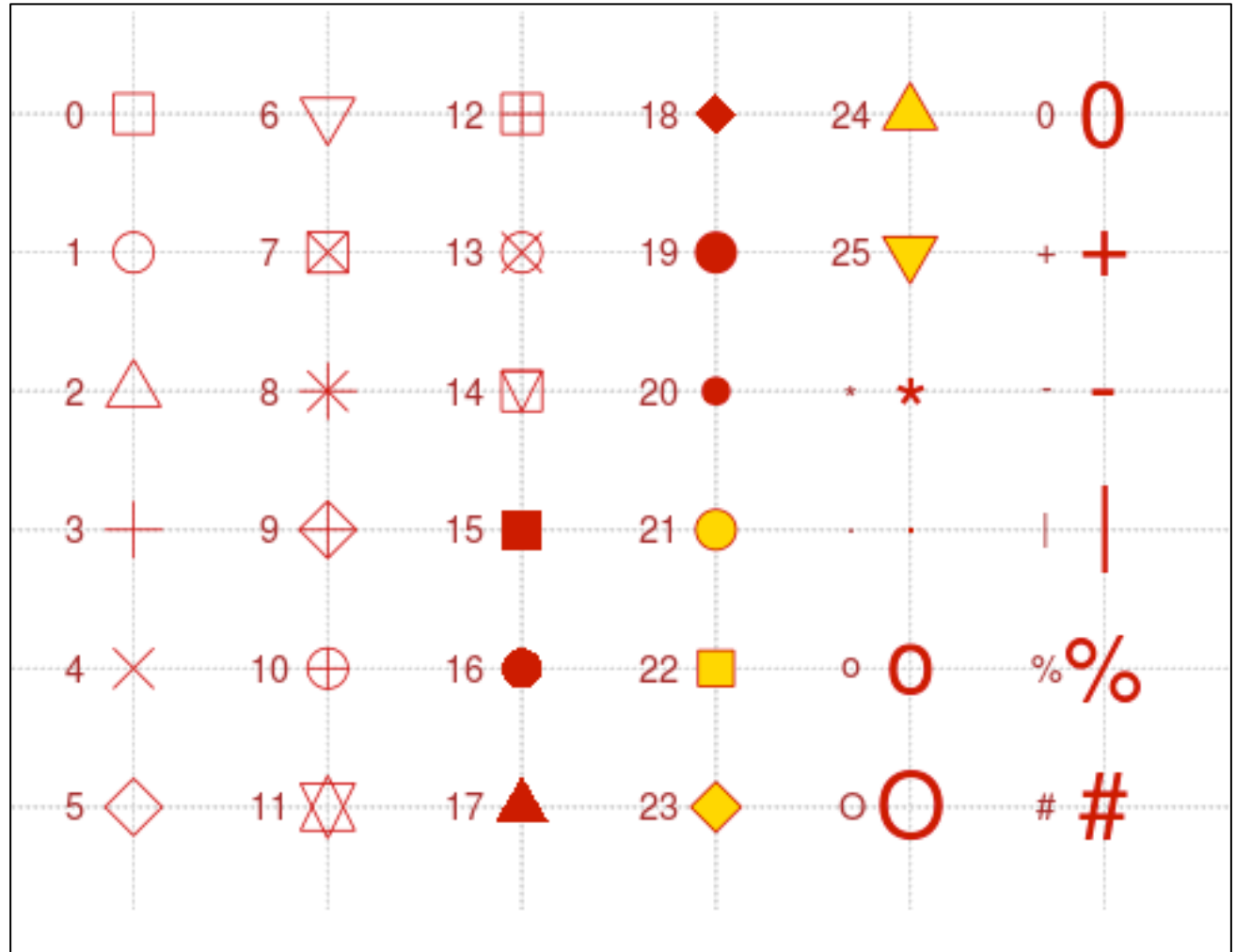
These color palettes are based on HCL and HSV color models. The results can be very aesthetically pleasing. There are some default palettes:

```
rainbow_hcl(4)
"#E495A5" "#ABB065" "#39BEB1" "#ACA4E2"
```

However, all palettes are fully customizable:  
`diverge_hcl(7, h = c(246, 40), c = 96, l = c(65, 90))`  
Choosing the values *would* be daunting. But there are some recommended palettes in the `colorspace` documentation. There is also an interactive tool that can be used to obtain a customized palette. To start the tool:  
`pal <- choose_palette()`

`grDevices`  
`palettes`  
`cm.colors`  
`topo.colors`  
`terrain.colors`  
`heat.colors`  
`rainbow`  
see P. 4 for options

`colorspace`  
`default palettes`  
`diverge_hcl`  
`diverge_hsl`  
`terrain_hcl`  
`sequential_hcl`  
`rainbow_hcl`



# Tables

- General rule of thumb is don't show the exact same data in **both** a table and Figure
- For some kinds of data, they may be the best way to communicate results to a wider audience.
- Use tables to illuminate patterns and show details of numbers you don't want to write fully in text of manuscript

# Tables

- Like graphs, tables are used to compare measurements between groups and expose relationships between variables.
- Put tables for storing numbers into online Appendix or Supplement
- Make your tables so that they cause the viewer to go “Oh!” and not “Huh?”

# Choosing a graph vs table

- Do you want readers to take away a picture of the overall trend or exact numerical values?
- Graphs show trends and patterns
- Tables show exact numbers → more difficult for our eyes to see patterns in tables

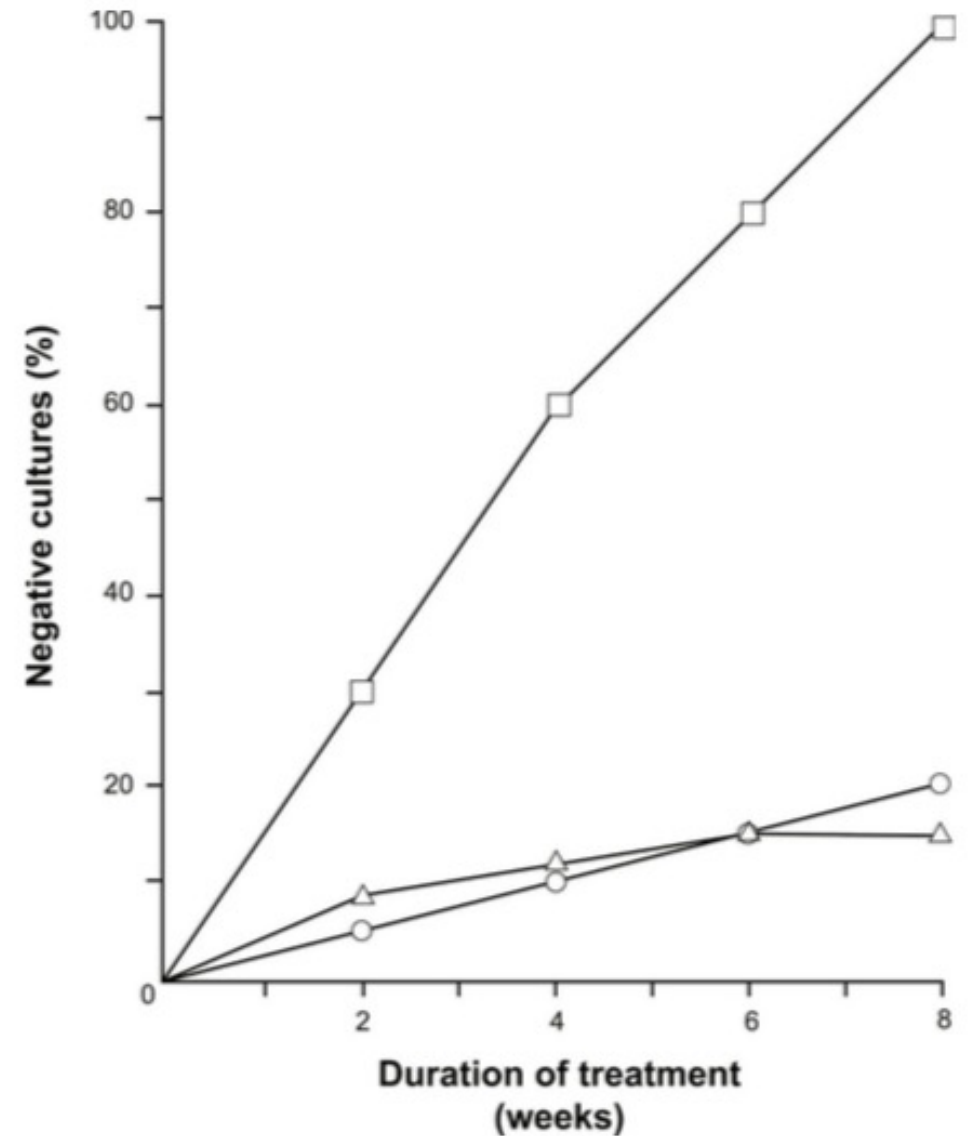
# Discussion: Which drug is more effective? Table or Graph best?

**Table 17.1.** Effect of streptomycin, isoniazid, and streptomycin plus isoniazid on *Mycobacterium tuberculosis*<sup>a</sup>

Treatment <sup>b</sup>	Percentage of negative cultures at:			
	2 wk	4 wk	6 wk	8 wk
Streptomycin	5	10	15	20
Isoniazid	8	12	15	15
Streptomycin + Isoniazid	30	60	80	100

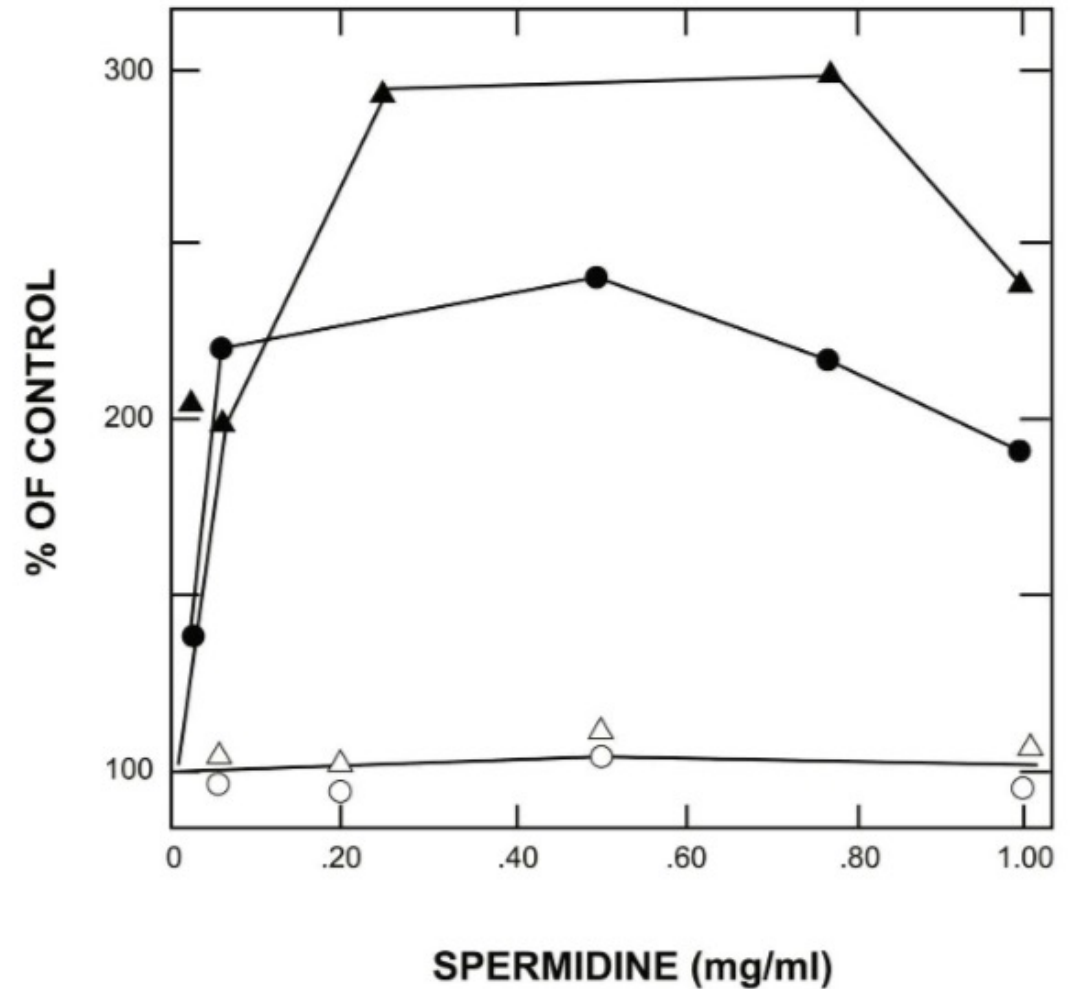
<sup>a</sup>The patient population, now somewhat less so, was described in a preceding paper (61).

<sup>b</sup>Highest quality available from our supplier (Town Pharmacy, Podunk, IA).



**Fig. 17.1.** Effect of streptomycin (  $\circ$  ), Isoniazid (  $\triangle$  ), and combination of both (  $\square$  ) on treating tuberculosis

What do you think of this graph?



**Figure 17.3.** Effect of spermidine on the transformation of *B. subtilis* BR 151. Competent cells were incubated for 40 min with spermidine prior to the addition of 5 µg of donor DNA per ml (●) or 0.5 µg of donor DNA per ml (▲). DNA samples of 5 µg (○) or 0.5 µg per ml (△) were incubated for 20 min prior to the addition of cells. (Redrawn from Fig. 1 in Clark PO and Leach FR. Stimulation of *Bacillus subtilis* transformation by spermidine. *Mol. Gen. Genet.* 178:21–25, 1980. © by Springer-Verlag 1980. With permission of Springer.)

# Improving Tables Examples

**Table 2.5-1** Inbreeding coefficient ( $F$ ) of Spanish Habsburg kings and queens and survival of their progeny.

King/Queen	$F$	Pregnancies	Miscarriages & stillbirths	Neonatal deaths	Later deaths	Survivors to age 10	Survival (total)	Survival (postnatal)
Ferdinand of Aragon								
Elizabeth of Castile	0.039	7	2	0	0	5	0.714	1.000
Philip I								
Joanna I	0.037	6	0	0	0	6	1.000	1.000
Charles I								
Isabella of Portugal	0.123	7	1	1	2	3	0.429	0.600
Philip II								
Elizabeth of Valois	0.008	4	1	1	0	2	0.500	1.000
Anna of Austria	0.218	6	1	0	4	1	0.167	0.200
Philip III								
Margaret of Austria	0.115	8	0	0	3	5	0.625	0.625
Philip IV								
Elizabeth of Bourbon	0.050	7	0	3	2	2	0.286	0.500
Mariana of Austria	0.254	6	0	1	3	2	0.333	0.400

*Source:* Data are from Alvarez et al. (2009).

# Critiquing my own table from PhD thesis

**Table 3. Summary of accelerometer error metrics relative to video data calculated on Australian fur seals.**

Acceleration	Function 1	Averaged over all animals (%)						Total
	Parameters	Detection ( $\pm$ SD)		FP rate ( $\pm$ SD)		Precision ( $\pm$ SD)		APC
X Surge	Generic	96.3	(3.2)	48.1	(15.3)	51.9	(15.3)	327
	Animal-specific	93.9	(3.0)	42.1	(15.0)	57.9	(15.0)	306
Y Sway	Generic	96.8	(3.2)	58.6	(5.8)	41.4	(5.8)	402
	Animal-specific	91.3	(7.1)	45.1	(18.5)	54.9	(18.5)	361
Z Heave	Generic	82.3	(10.8)	32.8	(13.9)	67.2	(13.9)	262
	Animal-specific	81.7	(9.4)	26.1	(16.1)	73.9	(16.1)	252

Error metrics included detection, false positive rate (FP rate), and precision as defined in the abbreviations. Animal-specific parameters were the parameters of minimum interval and variance threshold that yielded the greatest detection per animal per acceleration axis (Fig 4). Generic parameters (same for all animal) were set at 0.1 variance threshold with 5 second minimum interval for all animals and acceleration axis. Data are from the Random Testing subset (n = 97 dives).

doi:10.1371/journal.pone.0128789.t003

- Undefined abbreviation: APC
- Function 1 is not defined



# Outline

- Peer-feedback survey on Discussion Moderation and Presenting
- Why make graphs?
- Principles of effective display
- Types of graphs to achieve these principles
- Why some graphs fail, and what can be done
- Tables
- **Interactive Graphs**
- **Ggplot**
- **Motion (video data)**
- **Wrap-up**

# Interactive plots

- **Interactive plots**

- Graphs that allow the viewer to manipulate the image.
  - **Example:** Dive Move() for dive depth dataloggers on animals
- Line plot with basic “hover”
  - <https://rstudio.github.io/dygraphs/>
- Interactive scatter plot:
  - <https://cengel.github.io/R-data-viz/interactive-graphs.html#plotly>

# Data that moves: video data

Video graphs for displaying data are becoming more common

- <http://www.r-graph-gallery.com/3-r-animated-cube/>
- <https://ebird.org/science/status-and-trends/>

Measles outbreak simulation

- <https://www.theguardian.com/society/ng-interactive/2015/feb/05/-sp-watch-how-measles-outbreak-spreads-when-kids-get-vaccinated>

The Atlantic Slave Trade in Two Minutes

- [http://www.slate.com/articles/life/the\\_history\\_of\\_american\\_slavery/2015/06/animated\\_interactive\\_of\\_the\\_history\\_of\\_the\\_atlantic\\_slave\\_trade.html?wpsrc=sh\\_all\\_dt\\_fb\\_top](http://www.slate.com/articles/life/the_history_of_american_slavery/2015/06/animated_interactive_of_the_history_of_the_atlantic_slave_trade.html?wpsrc=sh_all_dt_fb_top)
- The fallen of WWII: <https://vimeo.com/128373915>

# Discussion paper next week

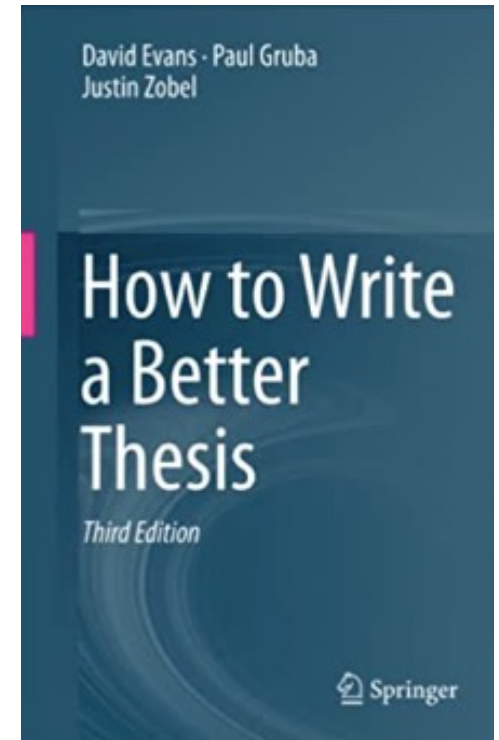
- Hurlbert, S. H. (1984). Pseudoreplication and the design of ecological field experiments. *Ecological Monographs* 54: 187–211
- Download from “**Handouts**” tab on course web site.
- Next Tuesday
  - Presenter: Leah Wessler
  - Moderator: Shawn Nirody
- **\*\*Please add your email to Google Sheets sign up\*\***
- Presenters and moderators are responsible for communicating with each other prior

# Assignment #1 Due 28 Jan @ 9pm

- Submit via Canvas only
- See header for Script
- See Rubric on Canvas
  
- Basic idea:
- Find a poor graph drawn from data and published by your thesis supervisor.
- Analyze it: why does it not succeed?
- Improve the graph using R.
- Analyze it: why does it succeed?
- Include R code **\*\*with meaningful annotations and R script header\***

# Optional Thesis Writing Resource

- Tips for writing the thesis as a whole
- Unique in describing thesis introductory & discussion chapter
- Ask me in workshop if you want an e-copy of scanned book (3<sup>rd</sup> ed)



Evans, David, Paul Gruba, and Justin Zobel. *How to write a better thesis*. Melbourne Univ. Publishing, 2011.

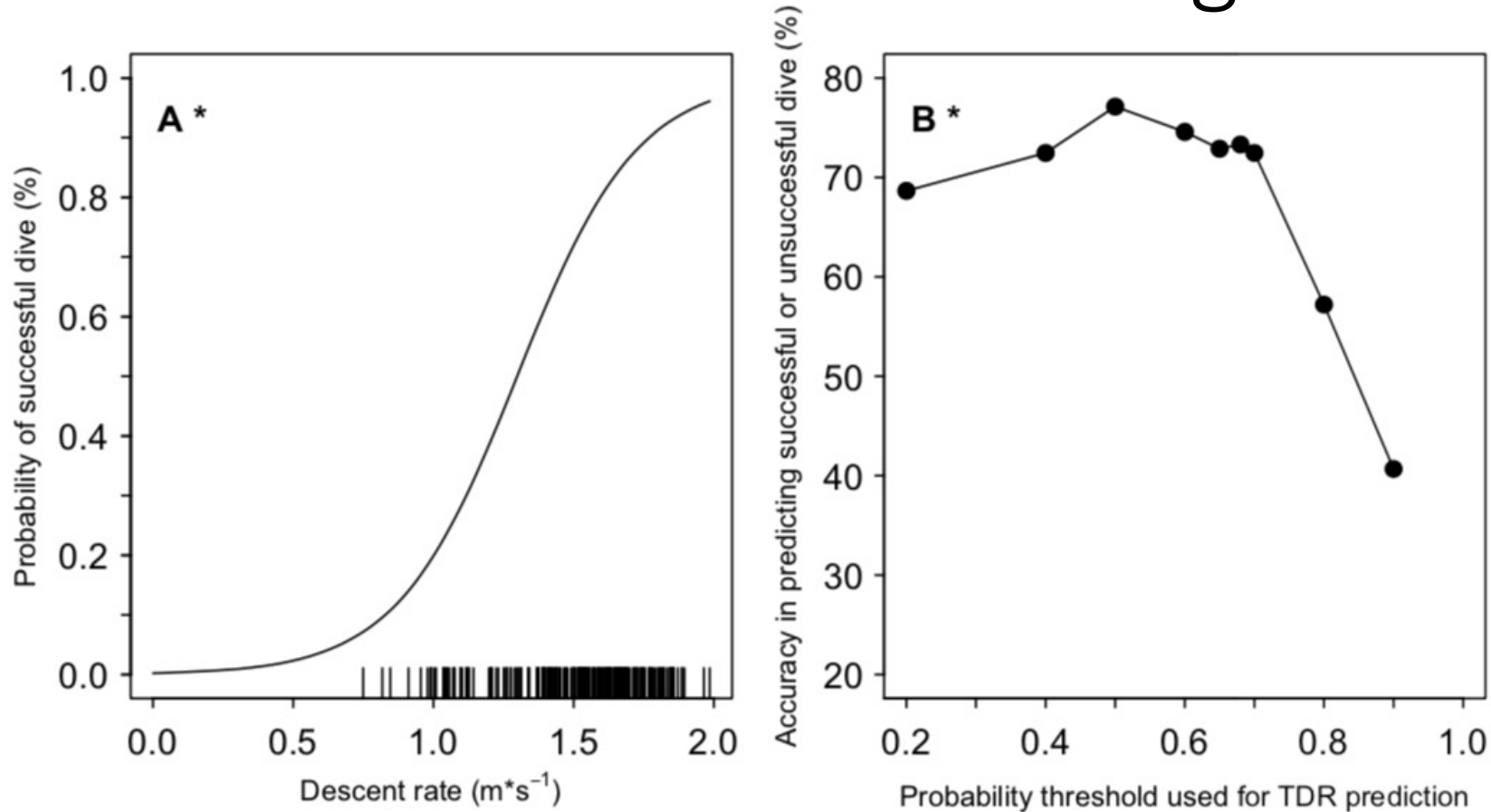
More example graphs

# Beth's examples of most used graphs

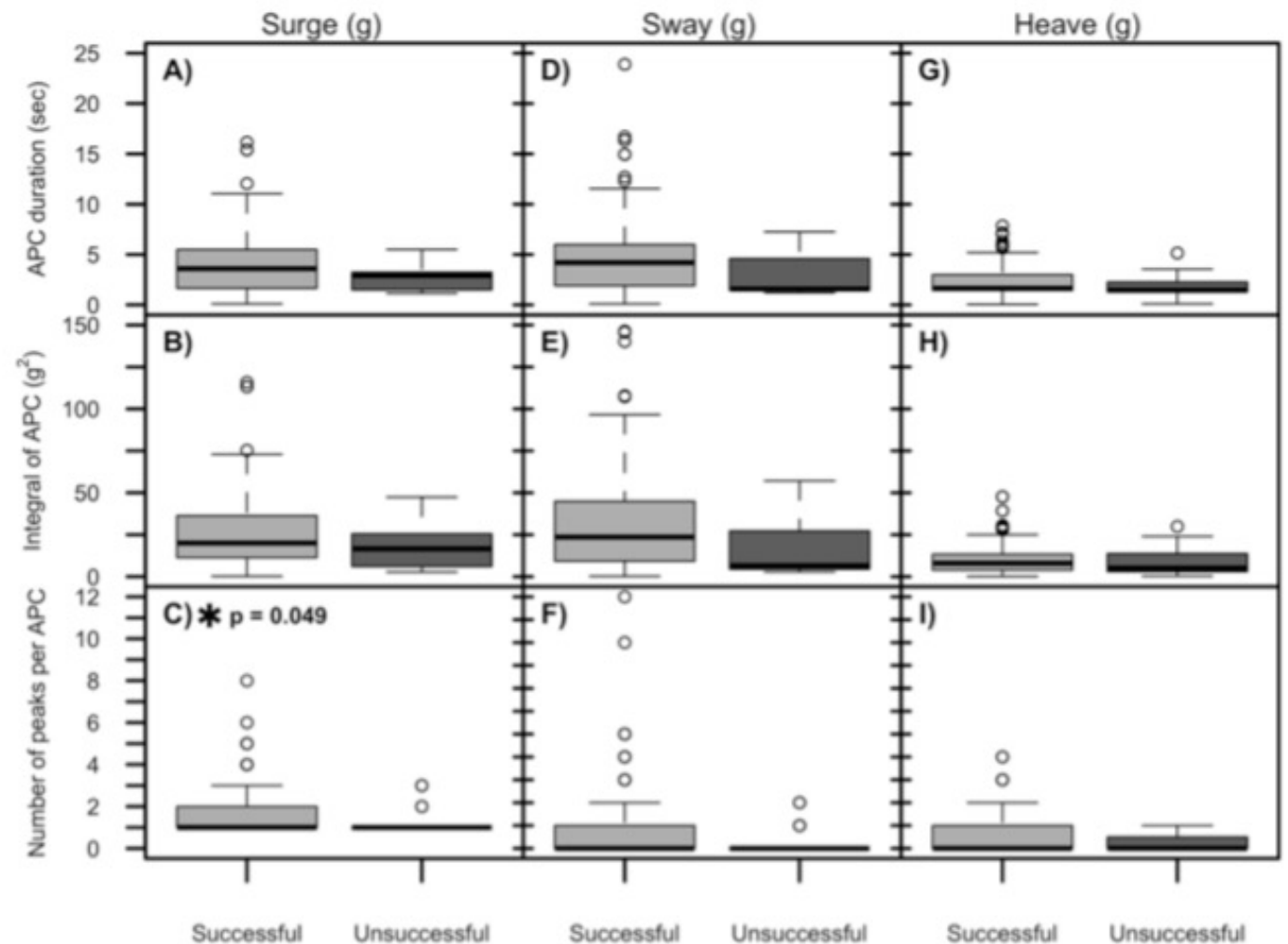
- Scatter plot with a model line (LME, both X and Y numerical)
- Box-n-whiskers (LME with categorical X variable, numerical Y)
- GLMM Graph example
- GAMM Graph example



# Generalized Linear Mixed Model Figure Example



**Fig. 2. Probability of a successful dive in response to dive characteristics and accuracy of the GLMM relative to animal-borne video.** (A) The most parsimonious model included descent rate as a predictor variable on the training subset (Table 2). Successful dives had at least one successful attempted prey capture (APC) per dive. Distributions are indicated with a rug plots. (B) Accuracy was calculated as the percent of dives correctly predicted as either successful or unsuccessful on the testing subset of dives (Table S2).



**Fig 5. Analysis of successful prey captures by accelerometers.** The accelerometer was not able to distinguish between successful and unsuccessful attempted prey captures (APC) on any axis of acceleration. Successful events included video where prey was captured, and unsuccessful events included events where the fur seal chased and missed an attempted capture attempt of the prey on video. Neither mean event duration (A, D, G), or integral area under the APC (B, E, H) significantly varied between successful or unsuccessful APC. The number of peaks in surge marginally differed between types of APC (C), but not for sway (F) or heave (I). Results are from Function 1 optimized with generic parameters (0.1 variance and 5 s). Each data point represents an individual APC.