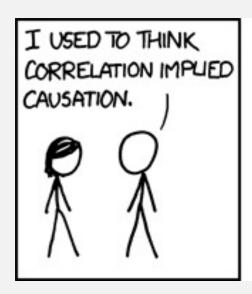
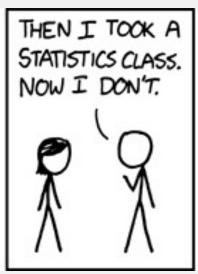
# Experimental Design

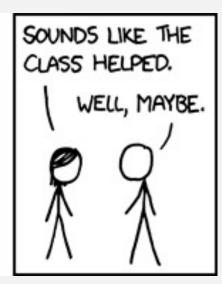
Biology 683

Lecture 1

Heath Blackmon







### Me

























I study the **evolution of traits**. Especially genomic traits. I use a variety of methods next-gen sequencing, experimental evolution, phylogenetic comparative methods, and theoretical approaches. I don't work with a single taxa; we have projects involving fish, mammals, reptiles, amphibians, insects, and bacteria all ongoing in my lab.



## You all

## Fill out index cards for me

## My Objectives

- Help you build an intuitive understanding of statistics
- Help you develop the confidence to think about the characteristics of the data that you will be collecting in your research and how you might analyze it.
- Get you comfortable with the idea of coding in R
- Help you develop the skills to handle datasets in R
- Help you develop the skills to build informative, honest, and intuitive data visualizations in R
- Make you a more productive and successful scientist!

## My View on Graduate Courses

- Graduate courses should open the door for you to become an expert in a field.
- However, you shouldn't need to become an expert to do well. Not every class is super central to your research.
- Therefore, I'm going to expose you to a lot of material and hope to really challenge you
  with some of the problems we solve. However, you can earn an A if you put forth an
  appropriate and reasonable effort.
- I hope that some of you embrace the skills that I am teaching you, as I believe they can be crucial to becoming an outstanding scientist.

## Pedagogy

Educational Evaluation and Policy Analysis Summer 1990, Vol. 12, No. 2, pp. 213-227

Class Size and Student Achievement: Research-Based Policy Alternatives

> Allan Odden University of Southern California





## What I will offer

 In class (MW 4-5:15) – lecture, discussion, questions and answers, live coding

- Monday Code Nights (MW 5:30-6:30) week before tests
- Saturday Code Days (9-12) donuts provided: Fill out form with weekends that you could attend.
- Lots of enthusiasm!

## Today

- Pedagological approach
- Syllabus / website / calendar
- Why I think you need this class
- R

# My opinions

Misuse of statistics is unethical as a scientist

## My opinions

Misuse or ignorance of statistics is unethical as a scientist

Poor training and maleficence are both responsible for failures

Statistical literacy in the general public is essential and lacking

Do your part: learn science of important topics and help friends and family understand them! This includes the statistical analysis and how we should let them inform our belief!

## Reproducibility crisis

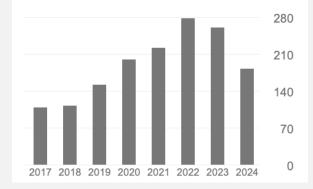
Started in the social sciences but some problems are widespread

possible causes of

f the reprodu

Citations	1634	1305	
h-index	20	20	SIS
i10-index	27	26	713

- small sample sizes
- p-hacking



unethical researchers/developers





Jonathan Pruitt Canada 150 Research Chair TED Talk 72.2 Million views

**Amy Cuddy** (2<sup>nd</sup> most popular TED Talk)

### Solutions

- Study preregistration (<u>COS</u>)
- Discuss What are possible
- solutions to the reproducibility
- · , crisis?

Systemic change - unlikely

# Why do biologists need statistics

- •We want to learn about the world often by testing hypotheses.
- •To test a hypothesis we have to design an experiment
- •Not all experiments have a traditional control and experimental treatment and this isn't always how we want to test a hypothesis
- •It is quite possible to design a study or collect data that cannot answer the questions that we have
- •This leads to poor manuscripts and can lead to bad practices like p-hacking

## Experimental Design

To design an experiment you need to understand how the data will be analyzed statistically.

- 1. How can you sample the population in which you are interested?
- 2. What tests are appropriate for your data?
- 3. What biases must be controlled for?
- 4. What sample size will be necessary?

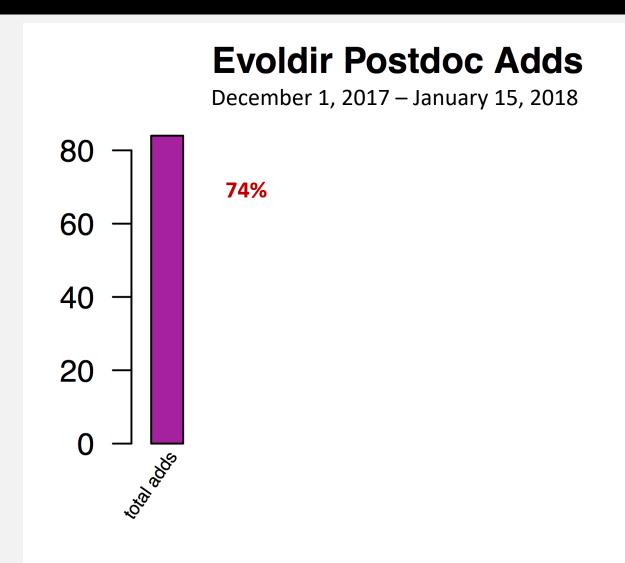
## Why not just collaborate with a statistician

- 1. In some cases this is a great option, but you have to understand enough to communicate.
- 2. If you publish a study you are responsible for its validity.
- 3. For most experiments simple methods suffice.
- 4. In many fields of biology there are sets of statistical tests that are expected for certain types of data.
- 5. For all of these reasons statistical analysis needs to involve people who understand the biological problem and the field of study.

## My stats philosophy

- Statistics is just another tool
- My responsibility as a scientists is to report the truth as accurately as possible and statistics help me in this regard
- We may NEED statistics to discern patterns in our data
- You need to understand where the signal that makes for a significant test comes from. Visualizing your data in the right way can do this!

# Why am I teaching this class?



### What is R

- R is an open and free statistical programming language that focuses on stats and graphics
- It works very similarly on all major operating systems
- It's also a full-fledged high level programming language (similar to Python)
- Very popular in industry so looks great on a CV.

## Why use R

1. Many statistical approaches have been implemented in the R environment.

- 2. Because it's open source, there are no proprietary secrets, as might be hiding in commercially available statistical packages.
- 3. Any program written in R will have access to all of R's tools for statistics and graphing.
- 4. New methods of analysis are being implemented in R by the scientists developing the methods.

## Why use R

- If you use R you can include a script with your manuscript <u>example</u>
   Reproducibility / Open science
   Reviewing
   Revising
- 6. Many methods (mixed models, quantitative genetics, etc.) are only available in R.
- 7. PLOTTING
- 8. Once you've learned one language you can learn others more easily.

## Downsides of R

- Learning curve
- Anyone can make a package so there is some junk out there
- Memory issues
- No language lasts forever and no language can do everything
  - Python
  - Awk
  - Julia

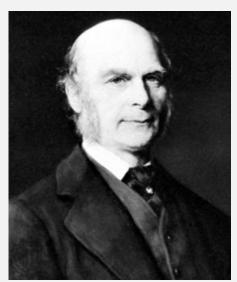
## The Origin of Statistics

### Much of modern statistics was an offshoot of genetics and evolution

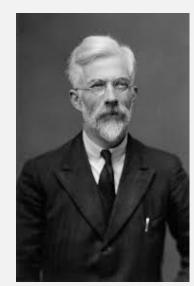
K. PEARSON 1857-1936 CORRELATION



F. GALTON 1822-1911 REGRESSION



R. FISHER 1890-1962 ANOVA



S. WRIGHT 1889-1988 PATH ANALYSIS



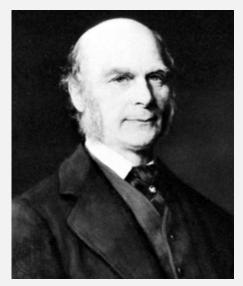
1900 rediscovery of Mendel's work was motivating problem.

# The Origin of Statistics

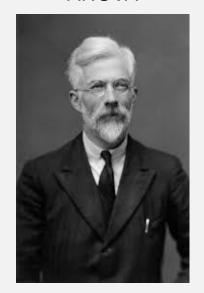
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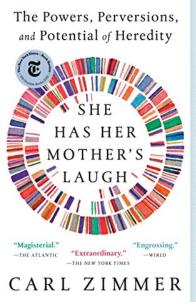


R. FISHER 1890-1962 ANOVA



# The disgraceful history of biostatistics

- Much of statistics was developed with the idea of showing that we could measure, scientifically analyze, and improve the "quality" of humans.
- The majority of geneticists and statisticians in the early 1900s were proponents of eugenics.
- What are the problems with this scientific/ethical



### For Future Classes

- 1. Install R and Rstudio on the computer you will use this semester
- 2. See me if you have problems installing it.

You can bring your laptop to class to follow along on coding that I do in front of you but this is not a requirement. Our room has insufficient plugs so charge ahead of time. I expect you to practice outside of class and come to practice sessions if you have problems.

Heath Blackmon
BSBW 425
Blackmon@tamu.edu

## Installing R and RStudio

#### **Installing R**

- 1. Go to the R homepage and click download R.
- Pick a mirror that is in Texas or at least in the United States.
- 3. Select the correct version for your system and follow the prompts.

#### **Installing Rstudio**

- 1. Go to the <u>RStudio homepage</u> and click on the download link below the free version of RStudio Desktop.
- 2. Select the correct version for your system and follow the prompts.

#### How you will be learning

1. I will code live in front of you, I will have some days times that we set aside extra time for help. HOMEWORK!

## Basics of R

#### 1. Demo R

#### **Data structures**

- vector
- matrix
- dataframe
- list

#### Data types

- numeric
- character
- logical
- factor

#### **Control elements**

- for
- if
- while

#### **Common functions**

- C
- matrix
- list
- sum
- mean
- sd
- sqrt
- abs
- paste
- rnorm
- rbinom
- rexp
- sample
- rep
- data
- Help
- which

#### **Basic base R plotting functions**

- hist
- plot
- density
- abline
- lines

#### **Operators**

- <
- ==
- >
- <
- %in%
- {
- •
- + \* / ^ %%

```
Practice

install.packages("swirl")

library("swirl")

swirl()

Complete two lessons of your choice
```