

# BIOSTATISTICS

... wait. What!?



Erik Kusch

[erik.kusch@au.dk](mailto:erik.kusch@au.dk)

Section for Ecoinformatics & Biodiversity  
Center for Biodiversity and Dynamics in a Changing World (BIOCHANGE)  
Aarhus University

28/10/2020

**1** Should you care?

**2** Biological Terminology

**3** Issues

**4** Me

**1** Should you care?

**2** Biological Terminology

**3** Issues

**4** Me

# The Big Question

Should you care about biostatistics?

# The Big Question

**YES!**

# The Big Question

**YES!**

Thank you for attending my TED talk.

1 Should you care?

2 Biological Terminology

3 Issues

4 Me

# Biological Terminology

**No**, biostatistics are **not just for math nerds**.

**Her: I'm a stats major**

**Me: [trying to think of something to impress her] yea I'm bad at math too**



Statisticians don't know important  
**biological background:**

- *Population vs. Sample*
- *Species, Family, Taxon, etc.*
- *Interpretation of results*

Biologists don't know important  
**statistical background:**

- *Unsupervised vs. Supervised Approaches*
- *Statistical Assumptions*
- *Parametric vs. Non-Parametric Tests*



# Biological Terminology

**No**, biostatistics are **not just for math nerds**.

**Her: I'm a stats major**

**Me:** [trying to think of something to impress her] **yea I'm bad at math too**



Statisticians don't know important  
**biological background:**

- *Population vs. Sample*
- *Species, Family, Taxon, etc.*
- *Interpretation of results*

Biologists don't know important  
**statistical background:**

- *Unsupervised vs. Supervised Approaches*
- *Statistical Assumptions*
- *Parametric vs. Non-Parametric Tests*

# Biological Terminology

**No**, biostatistics are **not just for math nerds**.

**Her: I'm a stats major**

**Me:** [trying to think of something to impress her] **yea I'm bad at math too**



Statisticians don't know important  
**biological background:**

- *Population vs. Sample*
- *Species, Family, Taxon, etc.*
- *Interpretation of results*

Biologists don't know important  
**statistical background:**

- *Unsupervised vs. Supervised Approaches*
- *Statistical Assumptions*
- *Parametric vs. Non-Parametric Tests*

# Basic Statistics

How often **do you** actually **check assumptions**?

## ■ *Assumptions:*

- Normality
- Independence
- Homogeneity of variances

→ Testing? Remedies?

## ■ *Scales and Distributions:*

- Continuous, Categorical
- Nominal, Binary, Ordinal, Interval, Relation/Ratio, Integer
- Gaussian Normal, Binomial, Poisson

→ Distinguish them?



# Basic Statistics

How often **do you** actually **check assumptions**?

## ■ Assumptions:

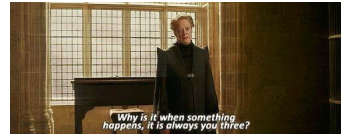
- Normality
- Independence
- Homogeneity of variances

→ Testing? Remedies?

## ■ Scales and Distributions:

- Continuous, Categorical
- Nominal, Binary, Ordinal, Interval, Relation/Ratio, Integer
- Gaussian Normal, Binomial, Poisson

→ Distinguish them?



# Basic Statistics

How often **do you** actually **check assumptions**?

## ■ Assumptions:

- Normality
- Independence
- Homogeneity of variances

→ Testing? Remedies?

## ■ Scales and Distributions:

- Continuous, Categorical
- Nominal, Binary, Ordinal, Interval, Relation/Ratio, Integer
- Gaussian Normal, Binomial, Poisson

→ Distinguish them?



# Basic Statistics

How often **do you** actually **check assumptions**?

## ■ Assumptions:

- Normality
- Independence
- Homogeneity of variances

→ Testing? Remedies?

## ■ Scales and Distributions:

- Continuous, Categorical
- Nominal, Binary, Ordinal, Interval, Relation/Ratio, Integer
- Gaussian Normal, Binomial, Poisson

→ Distinguish them?



# Basic Statistics

How often **do you** actually **check assumptions**?

## ■ Assumptions:

- Normality
- Independence
- Homogeneity of variances

→ Testing? Remedies?

## ■ Scales and Distributions:

- Continuous, Categorical
- Nominal, Binary, Ordinal, Interval, Relation/Ratio, Integer
- Gaussian Normal, Binomial, Poisson

→ Distinguish them?



# Correlations

Correlation is **not** necessarily **causation**.

Correlation tests yield two measurements:

- $r$  value (measure of correlation)
  - $r \approx 1$  (strong, positive correlation)
  - $r \approx 0$  (no correlation)
  - $r \approx -1$  (strong, negative correlation)
- $p$  value (measure of statistical significance)

**When you realize that all frequentist analyses are merely different versions of a correlation**



→ Get a feeling for it here <http://guessthecorrelation.com/>



# Correlations

Correlation is **not** necessarily **causation**.

Correlation tests yield two measurements:

- $r$  value (measure of correlation)
  - $r \approx 1$  (strong, positive correlation)
  - $r \approx 0$  (no correlation)
  - $r \approx -1$  (strong, negative correlation)
- $p$  value (measure of statistical significance)

**When you realize that all frequentist analyses are merely different versions of a correlation**



→ Get a feeling for it here <http://guessthecorrelation.com/>

# Correlations

Correlation is **not** necessarily **causation**.

Correlation tests yield two measurements:

- $r$  value (measure of correlation)
  - $r \approx 1$  (strong, positive correlation)
  - $r \approx 0$  (no correlation)
  - $r \approx -1$  (strong, negative correlation)
- $p$  value (measure of statistical significance)

**When you realize that all frequentist analyses are merely different versions of a correlation**



→ Get a feeling for it here <http://guessthecorrelation.com/>

# Advanced Statistics

What do you want to **analyse** and **predict**?

## ■ *Classifications:*

- K-Means
- Support-Vector Machines
- Hierarchies
- Networks

→ When to use which one?

## ■ *Regression:*

- Linear Models
- Least Squares vs. Maximum Likelihood
- Mixed Effect Models
- GLS/GLM, and GAM

→ How do you select the best model?

**Data not  
normal?**



**Want to  
appear more  
"computational"**



**Nonsignificant  
result?**



**Shoelace  
untied?**



# Advanced Statistics

What do you want to **analyse** and **predict**?

## ■ *Classifications:*

- K-Means
- Support-Vector Machines
- Hierarchies
- Networks

→ When to use which one?

## ■ *Regression:*

- Linear Models
- Least Squares vs. Maximum Likelihood
- Mixed Effect Models
- GLS/GLM, and GAM

→ How do you select the best model?

**Data not normal?**



**Want to appear more "computational"?**



**Nonsignificant result?**



**Shoelace untied?**



# Advanced Statistics

What do you want to **analyse** and **predict**?

■ *Classifications:*

- K-Means
- Support-Vector Machines
- Hierarchies
- Networks

→ When to use which one?

■ *Regression:*

- Linear Models
- Least Squares vs. Maximum Likelihood
- Mixed Effect Models
- GLS/GLM, and GAM

→ How do you select the best model?

**Data not normal?**



**Want to appear more "computational"**



**Nonsignificant result?**



**Shoelace untied?**



# Advanced Statistics

What do you want to **analyse** and **predict**?

■ *Classifications:*

- K-Means
- Support-Vector Machines
- Hierarchies
- Networks

→ When to use which one?

■ *Regression:*

- Linear Models
- Least Squares vs. Maximum Likelihood
- Mixed Effect Models
- GLS/GLM, and GAM

→ How do you select the best model?

**Data not normal?**



**Want to appear more "computational"**



**Nonsignificant result?**



**Shoelace untied?**



# Advanced Statistics

What do you want to **analyse** and **predict**?

■ *Classifications:*

- K-Means
- Support-Vector Machines
- Hierarchies
- Networks

→ When to use which one?

■ *Regression:*

- Linear Models
- Least Squares vs. Maximum Likelihood
- Mixed Effect Models
- GLS/GLM, and GAM

→ How do you select the best model?

**Data not normal?**



**Want to appear more "computational"**



**Nonsignificant result?**



**Shoelace untied?**



1 Should you care?

2 Biological Terminology

3 Issues

4 Me



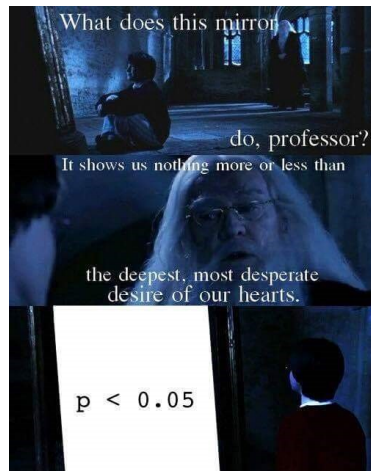
# Statistical Significance - the $p$ -value

## Misconceptions

- The  $p$ -value is not designed to tell us whether something is strictly true or false
- It is not the probability of the null hypothesis being true
- The size of  $p \neq$  strength of an observed effect

## Alternatives

- Effect Sizes
- Confidence Intervals
- Akaike Information Criterion (AIC)
- Bayes Factor
- Credible Intervals



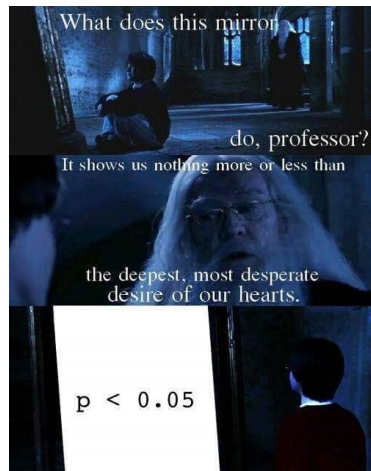
# Statistical Significance - the $p$ -value

## Misconceptions

- The  $p$ -value is not designed to tell us whether something is strictly true or false
- It is not the probability of the null hypothesis being true
- The size of  $p \neq$  strength of an observed effect

## Alternatives

- Effect Sizes
- Confidence Intervals
- Akaike Information Criterion (AIC)
- Bayes Factor
- Credible Intervals



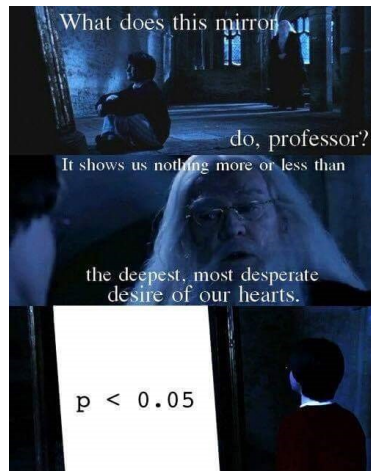
# Statistical Significance - the $p$ -value

## Misconceptions

- The  $p$ -value is not designed to tell us whether something is strictly true or false
- It is not the probability of the null hypothesis being true
- The size of  $p \neq$  strength of an observed effect

## Alternatives

- Effect Sizes
- Confidence Intervals
- Akaike Information Criterion (AIC)
- Bayes Factor
- Credible Intervals



# Coding Etiquette

## R Coding

- Object Modes
- Object Types
- Sub-setting
- Vectorisation
- Statements, Loops
- Functions, Packages

## Coding Schools

- Hard-coding vs. Soft-coding
- Base plot vs. ggplot2
- Base code vs. tidyverse



And what about **Git Hub**?

# Coding Etiquette

## R Coding

- Object Modes
- Object Types
- Sub-setting
- Vectorisation
- Statements, Loops
- Functions, Packages

## Coding Schools

- Hard-coding vs. Soft-coding
- `Base plot` vs. `ggplot2`
- `Base code` vs. `tidyverse`



And what about **Git Hub**?

# Coding Etiquette

## R Coding

- Object Modes
- Object Types
- Sub-setting
- Vectorisation
- Statements, Loops
- Functions, Packages

## Coding Schools

- Hard-coding vs. Soft-coding
- `Base plot` vs. `ggplot2`
- `Base code` vs. `tidyverse`



And what about **Git Hub**?

# Coding Etiquette

## R Coding

- Object Modes
- Object Types
- Sub-setting
- Vectorisation
- Statements, Loops
- Functions, Packages

## Coding Schools

- Hard-coding vs. Soft-coding
- Base `plot` vs. `ggplot2`
- Base `code` vs. `tidyverse`



And what about **Git Hub**?

# Manuscript Workflow

Using `Rmarkdown` for your research comes with a multitude of advantages:

- 1 Entire **workflow in one program** (`RStudio`)
- 2 **Research** and reports **reproducible** at the click of **one button**
- 3 **Combines** `R` functionality and  $\text{\LaTeX}$  formatting (if desired)
- 4 **Consistent formatting**
- 5 **Clear presentation of code**
- 6 **Dynamic documents** (you can generate various output document types)
- 7 Applicable for **almost all document types** you may desire as an output (e.g. manuscripts, presentations, posters, etc.)



1 Should you care?

2 Biological Terminology

3 Issues

4 Me

# Need Statistical Advice?

## Erik Kusch

### **Studies:**

PhD @ Aarhus University (currently enrolled)

M.Sc. @ University of Bergen

B.Sc. @ Technical University of Dresden

### **Experience:**

Biostatistics Tutor @ University of Leipzig

Biostatistics Research Assistant @ University of Leipzig

Biostatistics Research Assistant @ University of Kyoto

### **Research:**

- Dryland vegetation memory analyses
- Large-scale vegetation-climate modelling
- Remote sensing approaches in macroecology
- Biostatistical approaches in behavioural ecology
- Statistical downscaling of climate reanalysis data for use in biological analyses



Find me in room 318, building 1540 (Fridays, 09.00-12.00) or via [erik.kusch@bio.au.dk](mailto:erik.kusch@bio.au.dk).

# Need Statistical Advice?

## Erik Kusch

### **Studies:**

PhD @ Aarhus University (currently enrolled)

M.Sc. @ University of Bergen

B.Sc. @ Technical University of Dresden

### **Experience:**

Biostatistics Tutor @ University of Leipzig

Biostatistics Research Assistant @ University of Leipzig

Biostatistics Research Assistant @ University of Kyoto

### **Research:**

- Dryland vegetation memory analyses
- Large-scale vegetation-climate modelling
- Remote sensing approaches in macroecology
- Biostatistical approaches in behavioural ecology
- Statistical downscaling of climate reanalysis data for use in biological analyses



Find me in room 318, building 1540 (Fridays, 09.00-12.00) or via [erik.kusch@bio.au.dk](mailto:erik.kusch@bio.au.dk).