

# SIMPLE PARAMETRIC TESTS



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## 1 Background

## 2 Analyses

- t-Test (unpaired)
- t-Test (paired)
- Analysis of Variance (ANOVA)
- One-Way ANOVA
- Two-Way ANOVA
- ANCOVA

## 3 Our Data

- Choice Of Variables
- Research Questions

# Introduction

Parametric test are those statistical approaches which rely on **assumptions** about the parameters which define a population.

Prominent parametric tests include:

- Pearson correlation (Seminar 9 - Correlation Tests)
- **t-Test**
- **Analysis Of Variance** (ANOVA)
- Linear regression
- Multivariate extensions of parametric methods
- ...

# Terminology

A reminder about the distinction of parametric and non-parametric tests (taken from Seminar 6):

## Non-Parametric Tests

- Less *restrictive*
- Make *little to no assumptions*
- Often a **black box**
- Require *more data*

## Parametric Tests

- More *restrictive*
- Make *strict assumptions*
- **Easy to interpret**
- Require *less data*

→ Parametric tests are **numerous!**

# Purpose And Assumptions

## t-Test (unpaired)

`t.test(..., paired = FALSE)` in base R

*Purpose:*

To identify whether groups of variable values are different from one another.

$H_0$

*There is no difference in characteristics of the response variable values in dependence of the classes of the predictor variable.*

*Assumptions:*

- Predictor variable is binary
- Response variable is metric and **normal distributed** within their groups
- Variable values are **independent** (not paired)

→ Test whether variance of response variable values in groups are equal (`var.test()`) and adjust `t.test()` argument `var.equal` accordingly.

# Minimal Working Example

Let's feed data to our `t.test(..., paired = FALSE)` function that holds two groups with clearly differing means:

```
data <- c(rnorm(10, 5, 1), rnorm(10, 10, 1))
factors <- as.factor(rep(c("A", "B"), each = 10))
t.test(data ~ factors, paired = FALSE)

##
##  Welch Two Sample t-test
##
## data:  data by factors
## t = -10, df = 20, p-value = 3e-11
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##   -6.0 -4.4
## sample estimates:
## mean in group A mean in group B
##           5.1           10.3
```

The output above tells us that the means of our two groups are significantly different.

# Purpose And Assumptions

## t-Test (paired)

`t.test(..., paired = TRUE)` in base R

### *Purpose:*

To identify whether groups of variable values are different from one another.

### $H_0$

*There is no difference in characteristics of the response variable values in dependence of the classes of the predictor variable.*

### *Assumptions:*

- Predictor variable is binary
- Response variable is metric
- *Difference of response variable pairs* is **normal distributed**
- Variable values are **dependent** (paired)

# Minimal Working Example

Let's feed data to our `t.test(..., paired = TRUE)` function that holds two connected groups with clearly differing means:

```
data <- c(rnorm(10, 5, 1), rnorm(10, 10, 1))
factors <- as.factor(rep(c("A", "B"), each = 10))
t.test(data ~ factors, paired = TRUE)

##
## Paired t-test
##
## data: data by factors
## t = -10, df = 9, p-value = 5e-07
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -5.9 -4.1
## sample estimates:
## mean of the differences
## -5
```

The output above tells us that the means of our two connected groups are significantly different.



# Introduction to ANOVA

ANOVAs are used to test whether there is a difference between groups of variable values.

There are **multiple versions of ANOVAs**:

- One-way ANOVA (one predictor variable)
- Two-Way ANOVA (multiple predictor variables)
- MANOVA (multivariate ANOVA/multiple response variables)
- ANCOVA (categorical and continuous predictor variables)
- MANCOVA (multivariate ANCOVA)

# Data for ANOVA

We will use the `crabs` data set from the `MASS` package

```
library(MASS)
```

```
data(crabs)
```

```
head(crabs)
```

##	sp	sex	index	FL	RW	CL	CW	BD
## 1	B	M	1	8.1	6.7	16	19	7.0
## 2	B	M	2	8.8	7.7	18	21	7.4
## 3	B	M	3	9.2	7.8	19	22	7.7
## 4	B	M	4	9.6	7.9	20	23	8.2
## 5	B	M	5	9.8	8.0	20	23	8.2
## 6	B	M	6	10.8	9.0	23	26	9.8

# Purpose And Assumptions

## One-Way ANOVA

`anova()` in base R

*Purpose:* To explain the variance of a continuous response variable in relation to one predictor variables.

$H_0$  *Variance of response variable values is equal between levels of predictor variable.*

- Assumptions:*
- Predictor variable is categorical
  - Response variable is metric
  - *Response variable residuals* are **normal distributed**
  - Variance of populations/samples are equal (**homogeneity**)
  - Variable values are **independent** (not paired)

# Minimal Working Example

Let's test whether body depth (BD) of crabs are varying when grouped by sex:

```
OneWay <- with(crabs, lm(BD ~ sex))
anova(OneWay)

## Analysis of Variance Table
##
## Response: BD
##           Df Sum Sq Mean Sq F value Pr(>F)
## sex         1    19    18.8    1.61  0.21
## Residuals 198   2315    11.7
```

As we can see, sex does not make for a statistically significant predictor of crab body depth.

Take note that we do not deal with testing the assumptions here.

# Purpose And Assumptions

## Two-Way ANOVA

`anova()` in base R

*Purpose:* To explain the variance of a continuous response variable in relation to multiple predictor variables.

$H_0$  *Variance of response variable values is equal between levels of predictor variables.*

*Assumptions:*

- Predictor variables are categorical
- Response variable is metric
- *Response variable residuals* are **normal distributed**
- Variance of populations/samples are equal (**homogeneity**)
- Variable values are **independent** (not paired)

# Minimal Working Example

Let's test whether body depth (BD) of crabs are varying when grouped by sex and species as well as their interaction:

```
TwoWay <- with(crabs, lm(BD ~ sex * sp))
anova(TwoWay)

## Analysis of Variance Table
##
## Response: BD
##              Df Sum Sq Mean Sq F value    Pr(>F)
## sex             1      19      19      1.99    0.160
## sp              1     419     419    44.31 2.8e-10 ***
## sex:sp          1      42      42      4.48    0.035 *
## Residuals    196    1854         9
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The output above tells us that species and the interaction effect of sex and species are meaningful for understanding body depth of crabs.

# Purpose And Assumptions

## ANCOVA

`anova()` in base R

*Purpose:*

To explain the variance of a continuous response variable in relation to mixed (continuous and categorical) predictor variables.

$H_0$

*Adjusted variance and means of response variable values is equal between levels of predictor variables.*

*Assumptions:*

- Predictor variables are categorical or continuous
- Response variable is metric
- *Response variable residuals* are **normal distributed**
- Variance of populations/samples are equal (**homogeneity**)
- Variable values are **independent** (not paired)
- Relationship between the response and covariate is linear.

# Minimal Working Example

Let's test whether body depth (BD) of crabs are varying when grouped by species and the carapace width as a covariate:

```
Ancova <- with(crabs, lm(BD ~ sp * CW))
anova(Ancova)
```

```
## Analysis of Variance Table
##
## Response: BD
##              Df Sum Sq Mean Sq F value    Pr(>F)
## sp              1     419      419  2481.2 < 2e-16 ***
## CW              1    1880     1880 11130.5 < 2e-16 ***
## sp:CW           1         2         2    12.4  0.00054 ***
## Residuals    196         33         0
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The output above tells us that all of our model coefficients are significant.



# Variables We Can Use

## **Response variables** (metric)

- Weight
- Height
- Wing Chord
- Nesting Height
- Number of Eggs
- Egg Weight

## **Predictor variables** (categorical)

- Sex (binary)
- Climate (binary)
- Climate (3 levels - Continental, Semi-Coastal, Coastal)
- Home Range (3 levels - Small, Medium, Large)
- Site Index (11 levels)
- Predator Presence/Type (3 levels - Avian vs. Non-Avian vs. None)

# Research Questions And Hypotheses

So which of our major research questions (seminar 6) can we answer?

## unpaired t-Test

- *Climate Warming/Extremes*: Does sparrow morphology change depend on climate?
- *Sexual Dimorphism*: Does sparrow morphology change depend on Sex?

## paired t-Test (suppose a resettling program)

- *Climate Warming/Extremes*: Does sparrow morphology change depend on climate?

## One-Way ANOVA

- *Climate Warming/Extremes*: Does sparrow morphology depend on climate?
- *Predation*: Does nesting height depend on predator characteristics?
- *Site-wise variation*: Does sparrow morphology depend on sites?

## Two-Way ANOVA

- *Climate Warming/Extremes*: Does sparrow morphology depend on climate and sex?
- *Sexual Dimorphism*: Does sparrow morphology depend on population status and sex?

## ANCOVA

- *Climate Warming/Extremes*: Do sparrow characteristics depend on climate and latitude?

*Remember to **check assumptions**  
and find ways to circumvent  
violations of assumptions!*