STAT115: Introduction to Biostatistics

University of Otago Ōtākou Whakaihu Waka

Lecture 18: Paired Data

Outline

- Previous:
 - Started to look at relationships between variables
 - Frequency of brain waves (EEG) and sensory deprivation
 - ► Examples of relationship between one continuous and one categorical variable
 - Two groups are independent
- Today:
 - ► Look at paired data (two groups are not independent)
 - ► Start looking at relationships between two continuous variables

Motivating example

- Reaction time (ms) for 23 participants (press a button after stimulus)
 - University students
- There are two stimuli:
 - Auditory (a burst of white noise)
 - Visual (a circle flashing on a computer screen)
- Each participant exposed to both stimuli
 - ► Shouldn't use the approach from previous lecture
 - ► The two groups are not independent
 - We might expect someone with fast auditory reaction time to have a fast visual reaction
- Example of paired data
 - ▶ Each observation in group one has correspondence to an observation in group two

This is an exploratory study

Data

```
AV = read.csv('AV.csv')
head(AV)
```

```
## auditory visual
## 1 226.3 255.5

## 2 187.5 309.4

## 3 279.8 363.5

## 4 233.8 378.7

## 5 180.8 268.0

## 6 178.2 288.1
```

Paired: find the difference

• Look at the difference in the outcomes for each pair

```
AV$differ = AV$visual - AV$auditory
# this adds another variable (called differ) to the data frame AV
head(AV)
    auditory visual differ
##
## 1
       226.3 255.5 29.26
## 2
      187.5 309.4 121.91
## 3
      279.8 363.5 83.73
       233.8 378.7 144.83
## 4
## 5
      180.8 268.0 87.14
## 6
       178.2 288.1 109.87
```

Paired: back to the future

- Model the differences as if they were a single sample
 - lacktriangle The data are the differences and are given by y_d
 - ▶ The differences y_d are assumed to be normal with mean μ_d and variance σ_d^2
 - $lacktriangleq \mu_d$ is a parameter representing the mean difference in the population
- For our example:
 - $ightharpoonup y_d$ is the difference in reaction time (visual auditory)
 - \blacktriangleright μ_d is the population mean difference in reaction time (visual auditory)

In R

- For paired data: two ways to find confidence intervals and hypothesis tests in R
- Option 1: use t.test on the differenced values

```
t.test(AV$differ)
##
    One Sample t-test
##
## data: AV$differ
## t = 4.5, df = 22, p-value = 2e-04
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
   32.29 87.86
## sample estimates:
## mean of x
##
       60.08
```

In R

- For paired data: two ways to find confidence intervals and hypothesis tests in R
- Option 2: specify the two groups and include option paired = TRUE

```
t.test(AV$visual, AV$auditory, paired = TRUE)
##
    Paired t-test
##
## data: AV$visual and AV$auditory
## t = 4.5, df = 22, p-value = 2e-04
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
   32.29 87.86
## sample estimates:
## mean difference
##
             60.08
```

Output and interpretation

- Both approaches give identical confidence intervals
- Minor differences
 - ▶ Input differs: (1) input the differences; (2) input each group
 - Wording differences in output
 - 'One sample t-test' vs 'Paired t-test'
 - 'true mean' vs 'true mean difference'
 - 'mean of x' vs 'mean difference'
- Interpretation:
 - ▶ We are 95% confident that mean difference in the reaction times between visual and auditory stimuli is between (32.3, 87.9) ms

Hypothesis test

- Often with an exploratory study: use confidence interval
 - ► Calculate hypothesis test here as an example
- ullet The hypothesis test is in terms of μ_d
- Null hypothesis: assumption of no difference $(\mu_d=0)$
 - \vdash $\mathsf{H}_0: \mu_d = 0$
 - ▶ $H_A : \mu_d \neq 0$
- The *p*-value is 1.8498×10^{-4}
 - Evidence that data are incompatible with the null hypothesis
 - ▶ There is evidence (at the $\alpha=0.05$ level) that the data are incompatible with assumption of no difference

Extension

- Many applications may have more than two groups
 - Data from multiple independent groups
 - Multiple observations of each subject (repeated measures)
- There are statistical models for both cases
 - ► Independence: ANOVA (analysis of variance)
 - We will see this later in the course
 - ► Repeated measures: complex model
 - Outside the scope of this course

Relationship between continuous variables

- Previous examples: relationship between a continuous variable and a categorical variable
 - Continuous: reaction time; categorical: stimuli
 - ► Continuous: EEG frequency; categorical: sensory status (solitary/control)
- We are now going to consider relationships between two continuous variables

Motivating examples

- We are going to introduce three motivating examples
 - 1. The size of brushtail possums
 - Compare total length (mm) to head length (cm)
 - -n=104 observations
 - 2. Height of STAT 110 students
 - Compare father's height (cm) to son's height (cm)
 - -n=279 observations
 - 3. Squat weight of international power lifters
 - Comparing body weight (kg) to max squat weight (kg)
 - Photo from powerliftingtechnique.com
 - The athlete pictured (Kelly Branton) is in the dataset
 - -n = 9045 observations (athletes)
- All of these involve two continuous variables





Brushtail possums

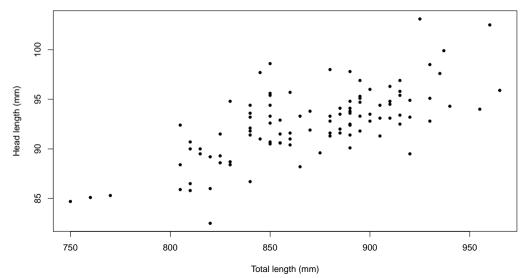
• Import the data

```
possum = read.csv('possum.csv')
```

• Have a look at the data:

```
head(possum)
##
     total_l head_l
## 1
         890
               94.1
               92.5
## 2
         915
## 3
         955
               94.0
               93.2
## 4
         920
## 5
         855
               91.5
## 6
         905
               93.1
```

Brushtail possums: scatterplot



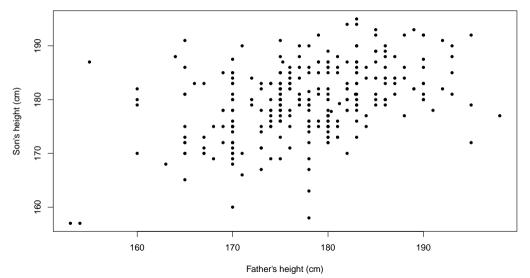
Father & son height

• Import the data

```
height = read.csv('height.csv')
```

• Have a look at the data:

Father & son height: scatterplot



Powerlifting

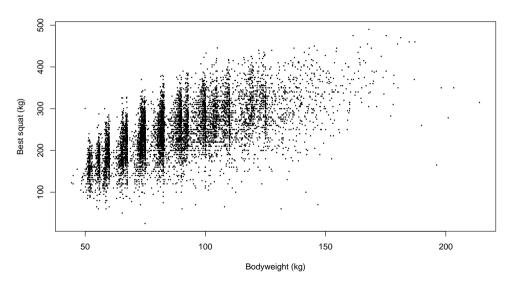
• Import the data

```
powerlift = read.csv('powerlift.csv')
```

• Have a look at the data:

```
head(powerlift)
##
     bodyweight bestsquat
## 1
           59.6
                    227.5
## 2
           67.2
                    255.0
## 3
           67.4
                    270.0
## 4
           59.9
                    260.0
## 5
           59.9
                    250.0
## 6
           56.0
                    210.0
```

Powerlift: scatterplot



Back to the beginning

- What was the first thing we did when we first encountered data in STAT115?
 - ▶ Found data summaries: sample mean and sample variance
- What summary describes the relationship between two continuous variables?

Correlation

- Correlation describes the strength of a linear relationship between two variables (let's call them x and y)
 - ▶ Always takes a value between -1 and 1
 - ▶ Population correlation represented by ρ (greek letter rho)
 - \triangleright Sample correlation represented by r
- With data $(x_1, y_1), (x_2, y_2), \ldots, (x_n, y_n)$, the correlation is given by

$$r = \frac{1}{n-1} \sum_{i=1}^{n} \frac{(x_i - \bar{x})}{s_x} \frac{(y_i - \bar{y})}{s_y}$$

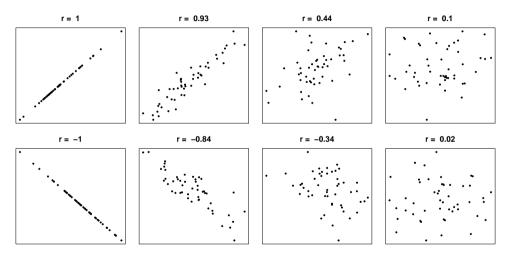
We will calculate the correlation using the R function cor

```
cor(possum$total_1, possum$head_1)
## [1] 0.6911
```

Understanding correlation

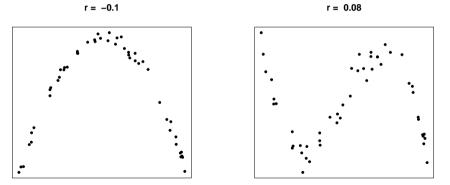
- Positive correlation:
 - \blacktriangleright If y is above its mean, then x is likely to be above it's mean (and vice versa)
- Negative correlation
 - \blacktriangleright If y is above its mean, then x is likely to be below it's mean (and vice versa)
- If the relationship is strong and positive
 - ightharpoonup r will be close to 1
- If the relationship is strong and negative
 - ightharpoonup r will be close to -1
- If there is no apparent (linear) relationship between x and y
 - ightharpoonup r will be close to 0

Understanding correlation: graphically I



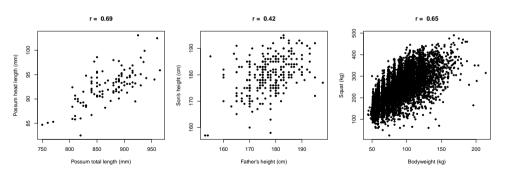
Understanding correlation: graphically II

- r measures the strength of the linear relationship
 - ightharpoonup Strong non-linear relationships can produce r values that do not reflect the strength of the relationship



Data

```
rposs = cor(possum$total_1, possum$head_1)
rheight = cor(height$son, height$father)
rpower = cor(powerlift$bodyweight, powerlift$bestsquat)
```



Test yourself at https://www.guessthecorrelation.com/

Limitations

- The correlation r is a useful summary
 - We may want to learn how precise it is: confidence interval
 - ▶ Such intervals can be found: cor.test in R
 - We will not consider them in STAT115
- The correlation as a summary is limited
- What might we want to know?
 - 1. Possum data: predict head length from a measurement of total length
 - 2. Height data: understanding and quantifying heritability of height as a trait
 - 3. Powerlifting: compare the squat weight of an athlete to their peers of a similar weight
- Correlation does not help us for 1 and 3
 - ▶ Limited for 2: quantifies the linear relationship, but does not describe it
 - What is the expected difference in height between a son with father who is 170 cm tall, and a son with father who is 180 cm tall?

Summary

- Looked at paired data
 - Model the difference between the two groups
 - Confidence intervals
 - Hypothesis test
- Looked at relationships between two continuous variables
- Explored a data summary: correlation
 - ► Gives the strength of a linear relationship between two variables
 - ▶ Always between -1 and +1
 - ► Easy to calculate in R