# Statistics for Linguists 08 July 2022

10:00	Workshop introduction
10:15	Loading and exploring datasets
10:45	Data transformation and coding
11:15	Practical exercise
12:15	Review of practical
12:30 - 13:30	LUNCH BREAK
13:30	lmer and glmer
14:30	Post-hoc analysis and model visualization
15:00	Practical exercise
16:00	Review of practical
16:15	Model building
17:00	End of workshop

# Statistics for Linguists

Post-hoc analysis and model visualization

# Learning objectives

- You will learn to load/import data
- Explore a dataset and create descriptive statistics
- Transform a dataset (if needed)
- Code your factors
- Build a mixed model
- Perform post-hoc statistics
- Visualize your data and your model

# What does this mean?

> m5.lmer <- lmer(log(ReadingTime) ~ capitalization \* determiner + (1 | participant), data = psycholinguistics\_data)

> summary(m5.lmer)

```
Groups
            Name
                       Variance Std.Dev.
participant (Intercept) 0.03543 0.1882
Residual
                       0.31941 0.5652
Number of obs: 2039, groups: participant, 30
Fixed effects:
                                                        df t value Pr(>|t|)
                             Estimate Std. Error
(Intercept)
                                                  28.85073 158.753
                             5.81156
                                        0.03661
                                                                    <2e-16 ***
capitalization1
                             0.01593 0.01258 2006.66649 1.266
                                                                     0.205
determiner1
                            -0.19560
                                       0.01260 2009.09620 -15.520
                                                                    <2e-16 ***
                            -0.01616 0.01257 2006.50155 -1.285
capitalization1:determiner1
                                                                     0.199
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## What does this mean?

```
> m6.lmer <- lmer(log(ReadingTime) ~ condition + (1 | participant),
data = psycholinguistics data)
> summary(m6.lmer)
                   Name
                        Variance Std.Dev.
         Groups
         participant (Intercept) 0.03543 0.1882
         Residual
                              0.31941 0.5652
        Number of obs: 2039, groups: participant, 30
        Fixed effects:
                       Estimate Std. Error df t value Pr(>|t|)
        (Intercept) 5.97508 0.04188 49.36933 142.665 <2e-16 ***
        condition+C/-D -0.35888 0.03593 2008.31218 -9.988 <2e-16 ***
        condition-C/+D 0.06418 0.03402 2006.24503 1.887 0.0594 .
        condition-C/-D -0.35935 0.03511 2007.16928 -10.235 <2e-16 ***
        Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# Post-hoc analyses

 Post-hoc analyses might be necessary because Imers don't provide a comparison of every level against each other level of a factor

 For continuous variables, a post-hoc analysis may involve correlation testing

• For factors, we can use special packages such as emmeans

### emmeans

- Computes estimated marginal means (EMMs) for specified factors or factor combinations in a linear model; and optionally, comparisons or contrasts among them
- > m5.lmer <- lmer(log(ReadingTime) ~ capitalization \* determiner + (1 | participant), data = psycholinguistics\_data)
- > emmeans(m5.lmer, pairwise ~ capitalization \* determiner, adjust = "Tukey")

### emmeans

> emmeans(m5.lmer, pairwise ~ capitalization \* determiner, adjust =

"Tukey")

```
$emmeans
capitalization determiner emmean SE df lower.CL upper.CL
```

посар	nouet	3.02 0.0429	34.7	5.55	3.70
cap	nodet	5.62 0.0436	58.1	5.53	5.70
nocap	det	6.04 0.0420	50.4	5.95	6.12

E 62 0 0420 E4 7

cap det 5.98 0.0419 49.7 5.89 6.06

Degrees-of-freedom method: kenward-roger

Results are given on the log (not the response) scale.

Confidence level used: 0.95

#### \$contrasts

```
contrast estimate SE df t.ratio p.value nocap nodet - cap nodet -0.000471 0.0371 2007 -0.013 1.0000 nocap nodet - nocap det -0.423531 0.0353 2007 -12.006 <.0001 nocap nodet - cap det -0.359354 0.0351 2007 -10.235 <.0001 cap nodet - nocap det -0.423061 0.0361 2009 -11.719 <.0001 cap nodet - cap det -0.358884 0.0359 2008 -9.987 <.0001 nocap det - cap det 0.064177 0.0340 2006 1.887 0.2341
```

### emmeans

> emmeans(m6.lmer, pairwise ~ condition, adjust = "Tukey")

#### \$emmeans

```
        condition emmean
        SE
        df lower.CL upper.CL

        +C/+D
        5.98 0.0419 49.7
        5.89 6.06

        +C/-D
        5.62 0.0436 58.1
        5.53 5.70

        -C/+D
        6.04 0.0420 50.4
        5.95 6.12

        -C/-D
        5.62 0.0429 54.7
        5.53 5.70
```

Degrees-of-freedom method: kenward-roger Results are given on the log (not the response) scale. Confidence level used: 0.95

#### \$contrasts

```
contrast estimate SE df t.ratio p.value (+C/+D) - (+C/-D) 0.358884 0.0359 2008 9.987 <.0001 (+C/+D) - (-C/+D) -0.064177 0.0340 2006 -1.887 0.2341 (+C/+D) - (-C/-D) 0.359354 0.0351 2007 10.235 <.0001 (+C/-D) - (-C/+D) -0.423061 0.0361 2009 -11.719 <.0001 (+C/-D) - (-C/-D) 0.000471 0.0371 2007 0.013 1.0000 (-C/+D) - (-C/-D) 0.423531 0.0353 2007 12.006 <.0001
```

# Post-hoc analyses

- For continuous variables, this makes little sense.
- > emmeans(mn.lmer, pairwise ~ sequential\_trial, adjust="Tukey")
- How could we further investigate the relation between sequential\_trial and reading time?
  - Group the independent variable (often done in cognitive domains, e.g. high vs. low working memory)
  - Correlations (with different subsets of the data, when needed)
  - Visually presenting the data or model output

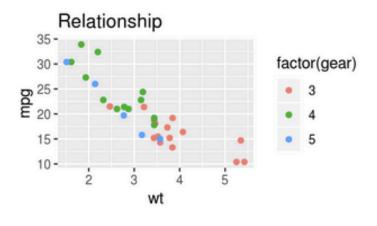
# Post-hoc analyses

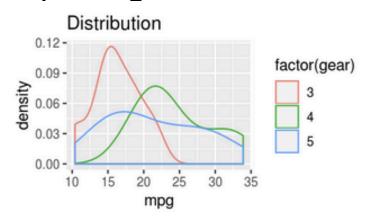
- Correlation
- > cor.test(x,y)
- Multiple types possible (pearson's, spearman's,...)
- However, this again ignores the issue of variability between participants and items

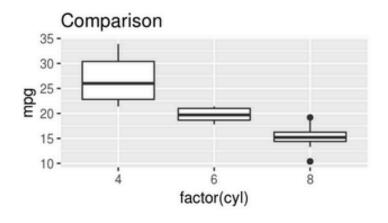
Pearson's product-moment correlation

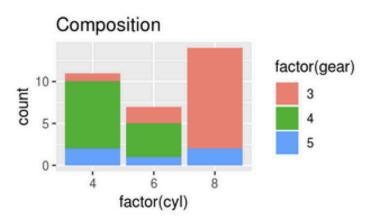
```
data: psycholinguistics_data2$sequential_trial and psycholinguistics_data2$ReadingTime
t = 0.4664, df = 2037, p-value = 0.641
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
   -0.03309113    0.05371890
sample estimates:
        cor
0.01033335
```

Data plots are easiest with the ggplot package





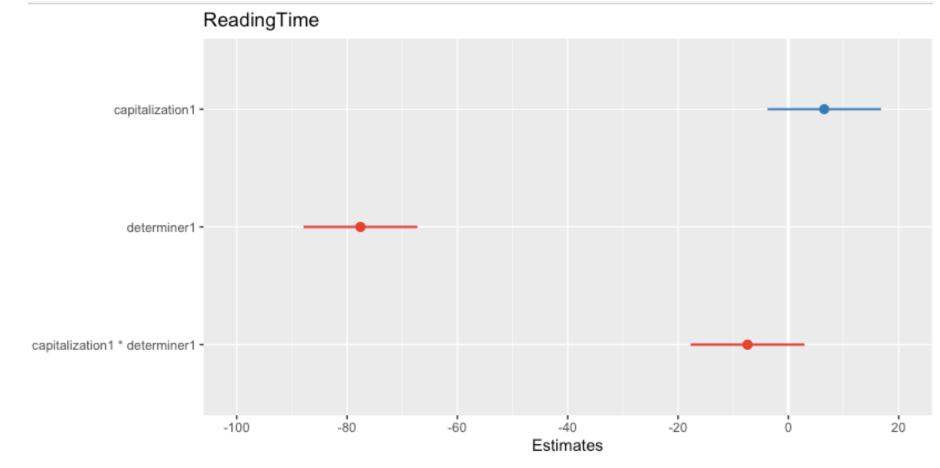




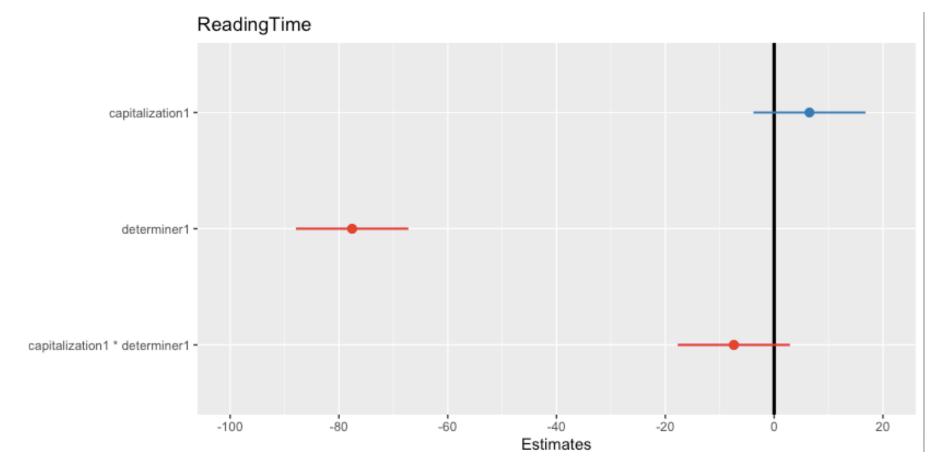
- However, model plots display the model effects, not the original data
- The can be used for presentation (in articles etc.) or for better understanding your results

> plot\_model(m5.lmer) # from sjPlot package

> plot\_model(m5.lmer) # from sjPlot package

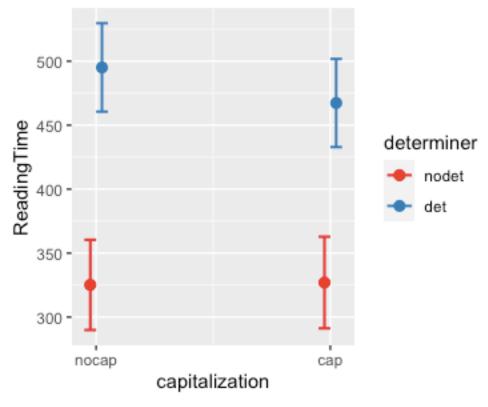


> plot\_model(m5.lmer) # from sjPlot package



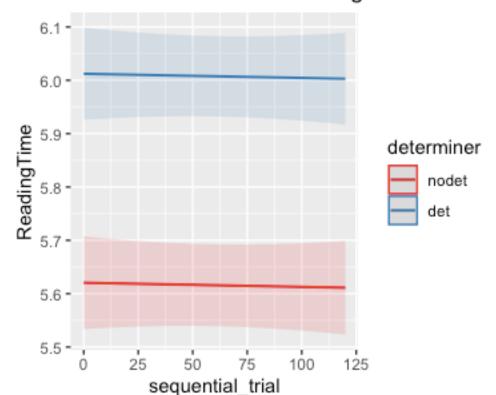
• > plot\_model(m5.lmer, type = "eff", terms = c("capitalization", "determiner"))

#### Predicted values of ReadingTime



Also with one continuous variable

• > plot\_model(m7.lmer, type = "eff", terms = c("sequential\_trial", "determiner")) Predicted values of ReadingTime



# Effect sizes

- Effect size is the size of the effect
- In our data, the effect of capitalization was 430-419 = 11 ms (dependent on outlier removal technique)
- Whether 11 ms is a lot or not depends on the variation in the data
- This is reflected in a relative or standardized effect size
- Popular measure of standardized effect size for comparisons between two means: Cohen's d

# Effect sizes

 Popular measure of standardized effect size for comparisons between two means: Cohen's d

> lme.dscore(m5.lmer, psycholinguistics\_data, type="lme4")

```
t df d
capitalization1 1.5909857 2025.474 0.07070221
determiner1 -14.3795260 2027.232 -0.63873819
capitalization1:determiner1 -0.8180792 2025.402 -0.03635547
```

- This value will then have to be interpreted
  - Typically d = 0.2 represents a 'small' effect size, 0.5 represents a 'medium' effect size and 0.8 a 'large' effect size

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