

Course outline

- Session 1. Simple Regression; Individual Difference; Intro to Linear mixed-effect models (LMMs)
- Session 2. LMMs (lmer); Introduction to Generalised LMMs (glmer)
- Session 3. Ggeneralised LMMs (glmer);
 Model assumptions and diagnostics



Session 3 Roadmap (today)

Model Comparison & Selection

- Generalised LMMs
- Model Assumptions
- Exercise and Q&A





Model Comparison and Selection

Recall: Structure of LMMS

• R code:

```
Imer (y \sim x1 * x2 +
(1 + x1 * x2 | Grouping_1) +
(1 + x1 | Grouping_2),
data = datafilename)
```

Recall: Model Fit

- No *R*²
- Use Maximum likelihood Ratio test to compare models
- R code: anova(model1, modle2)

Model Comparison & Selection

In what order should you build your models?

- Which model should you select as final model?
 - > Barr, Levy, Scheepers, & Tily, (2013): keep it maximal
 - Matuschek, Kliegl, Vasishth, Baayen & Bates (2017): make it parsimonious to balance Type1 error and power

Common Issues

Convergence

```
Warning message:
In checkConv(attr(opt, "derivs"), opt$par, ctrl = control$checkConv, :
Model failed to converge with max|grad| = 0.0206805 (tol = 0.002, component 1)
```

Overfitting

```
boundary (singular) fit: see ?isSingular
```

Deal with Convergence Issue

```
Warning message:
In checkConv(attr(opt, "derivs"), opt$par, ctrl = control$checkConv, :

Model failed to converge with max|grad| = 0.0206805 (tol = 0.002, component 1)
```

Solutions:

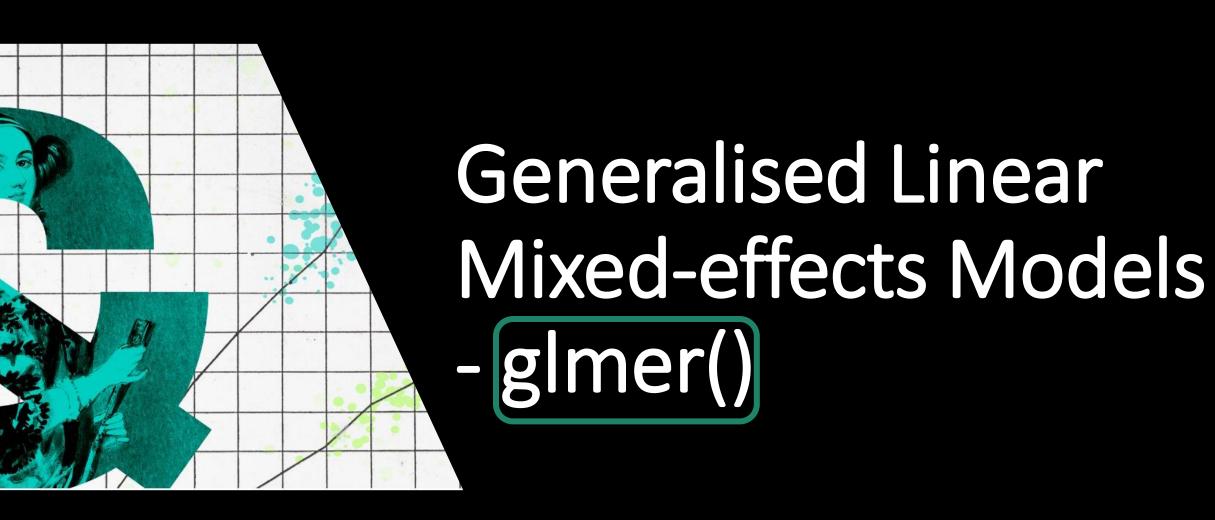
- Adjust stopping (convergence) tolerances for the nonlinear optimizer, using the optCtrl() argument to ImerControl.
- Centre and standardise continuous predictor variables
 - the scale() function

Deal with Overfitting Issue

boundary (singular) fit: see ?isSingular

Solutions:

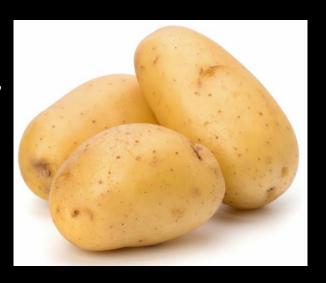
- Remove the most complex part of the random effects structure (i.e. random slopes)
- Maybe acceptable to remove a specific random effect term when its variance estimates are very low



Binary Outcome

- Pass or Fail
- Invest or Not
- Vote 'Yes' or 'No'
- 'Correct' or 'incorrect' answer
- Fixation on the target image or not
- Align with partner or not
 - Lexical choice
 - Grammatical choice

'potato' or 'tattie'





- (a) Gromit gave Wallace some cheese.
- (b) Gromit gave some cheese to Wallace.

Generalised Linear Mixed-effects Models

• glmer () for binary outcome

• *R* code:

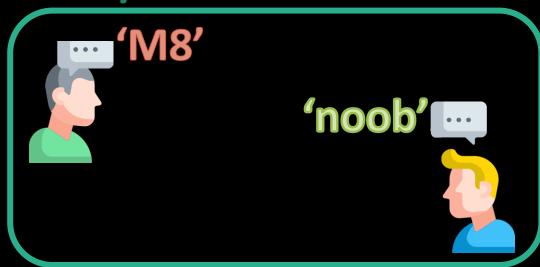
```
glmer (y \sim x1 * x2 +
	(1 + x1 * x2 | Grouping_1) +
	(1 + x1 | Grouping_2),
	data = datafilename,
	family = 'binomial'
```

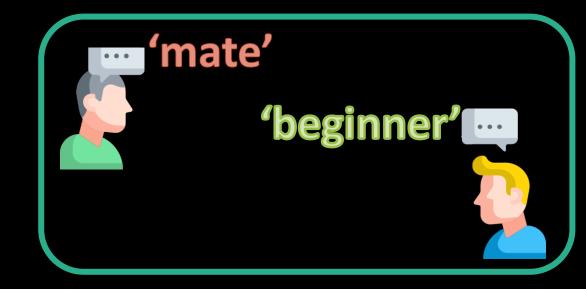
Example: Lexical Choice Study

'Noob' or 'Beginner'?

Internet slang data (simulated data)

• Hypothesis: People were more likely to use internet slang words (as relative to alternative expressions) after seeing their conversational partner uses a slang rather than a stand word.





Example: Lexical Choice Study

Repeated measure (36 items/participant)

One predictor (3 conditions):
 'prime' = slang / standard words / other

• Two sources of random variance

•	Participant †	Item [‡]	List [‡]	Trial [‡]	Prime [‡]	Response [‡]
15	1	11	1	81	standard	0
16	1	11	1	109	other	0
17	1	12	1	8	slang	0
18	1	13	1	106	standard	0
19	1	14	1	25	other	0
20	1	15	1	1	slang	1
21	1	16	1	103	standard	0
22	1	17	1	38	slang	1
23	1	18	1	49	standard	0
24	1	18	1	70	other	0
25	1	19	1	42	other	0
26	1	21	1	46	slang	1
27	1	22	1	87	standard	0
28	1	23	1	18	other	0
29	1	24	1	120	slang	0
30	1	25	1	116	standard	0
24		20	4	70	-1	
Showing 15 to 32 of 5,434 entries, 6 total columns						

Build a Glmer Model

Interpret Coefficients

```
Generalized linear mixed model fit by maximum likelihood (Laplace Approximation) ['glmerMod']
 Family: binomial (logit)
Formula: Response ~ Prime + (1 | Participant) + (1 | Item)
   Data: data ISD
Control: glmerControl(optimizer = "bobyga". optCtrl = list(maxfun = 2e+05))
                   logLik deviance df.resid
           5658.3 -2807.7
                            5615.4
Scaled residuals:
             10 Median
-4.9925 -0.5758 -0.2008 0.6024 4.8213
Random effects:
                         Variance Std.Dev.
 Groups
 Participant (Intercept) 2.4345 1.5603
             (Intercept) 0.1262 0.3552
Number of obs: 5386, groups: Participant, 160; Item, 30
Fixed effects:
              Estimate Std, Error z value Pr(>|z|)
             -0.27907 0.15230 1.832
Primestandar -0.93888
                        0.08594 -10.925
              1.12819
                         0.08583 13.145
Primesland
                                           <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Correlation of Fixed Effects:
            (Intr) Prmstn
Primestndrd -0.266
Primeslang -0.277 0.441
```

β 0 Probability of the intercept:

$$\frac{e^{-0.28}}{1+e^{-0.28}}=0.43$$

β 1 Probability of slope 1:

$$\frac{e^{(-0.28+(-0.94))}}{1+e^{(-0.28+(-0.94))}}=0.23$$

β 2 Probability of slope 2:

$$\frac{e^{(-0.28+1.13)}}{1+e^{(-0.28+1.13)}} = 0.70$$

- Coefficients are in logit units (log-odds)
- Can be transformed to probabilities: $prob(x) = \frac{e^x}{1+e^x}$



Model Assumptions

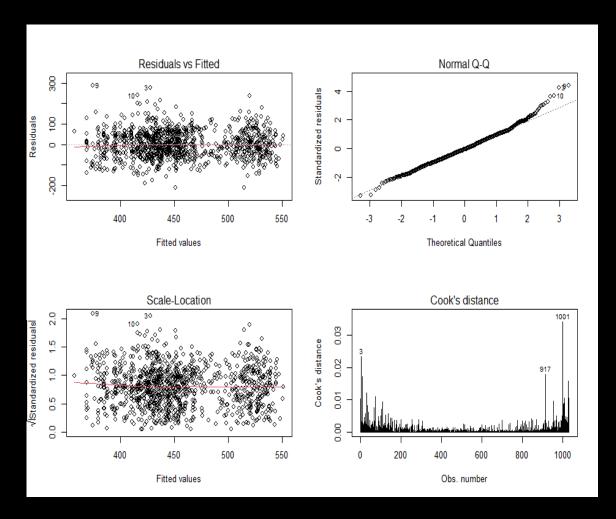
Model Assumptions: Linear regression

- Nature of the model
 - the relation between predictor and outcome has to be linear

- Nature of the errors (i.e., residuals)
 - normal and independent of each other

Model Assumptions: Simple regression Im()

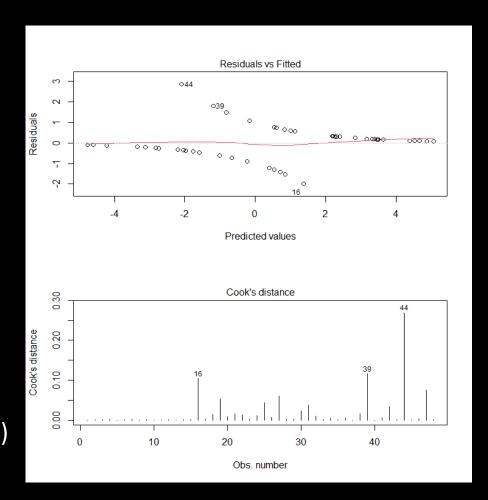
- Required(LINE):
 - Linearity of relationships
 - Independence of residuals
 - Normality of residuals
 - Equal variances for residuals
- Desirable:
 - uncorrelated predictors (no collinearity)
 - no outliers



Model Assumptions: Simple regression glm()

For binomial DVs, (logistic regression)

- Required:
 - LINEAR relationships between IVs and log-odds
 - Normality of residuals
 - homogeneity of variance
 - Independence of residuals
- Desirable:
 - uncorrelated predictors (no collinearity)
 - no "bad" (overly influential) observations
 - large samples (due to maximum likelihood fitting)



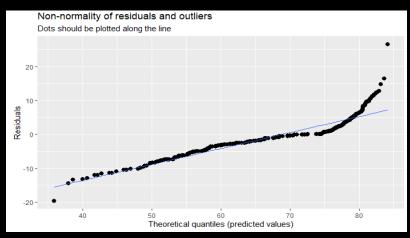
Model Assumptions: Mixed-effects Models

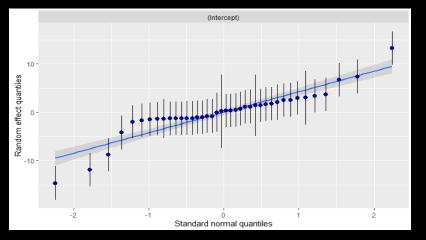
Similar to simple linear regressions model

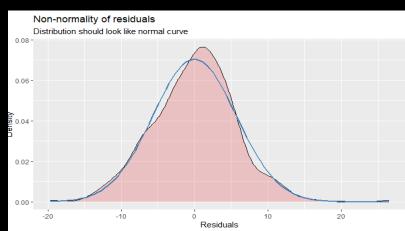
- Error is random
- Residuals at multiple levels
 - Level1 residuals: mean = 0, variance constant (R code: residual())
 - Level2+ residuals: mean = 0, variance constant (R code: ranef())

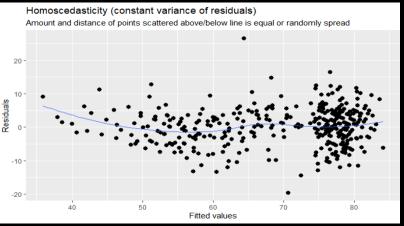
Model Assumptions: Mixed Models – Imer()

plot_model(mMixed_reduced , type = "diag")



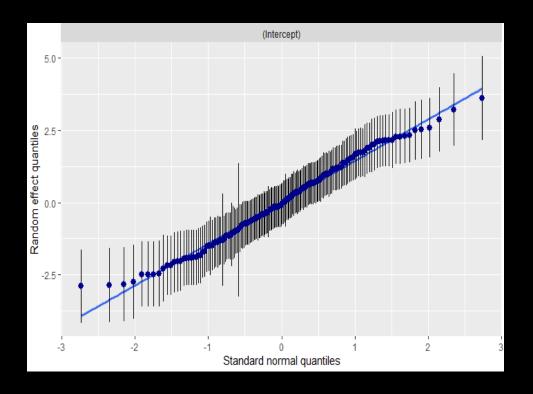


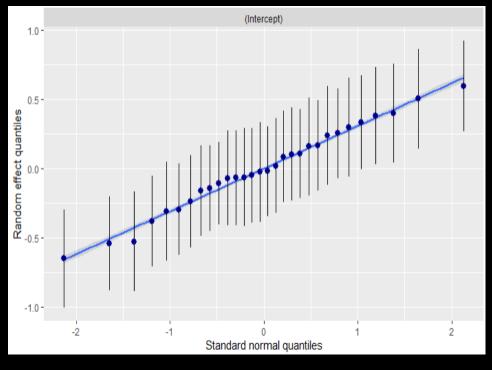




Model Assumptions: Mixed Models – glmer()

plot_model(m3_dataISD, type = "diag")





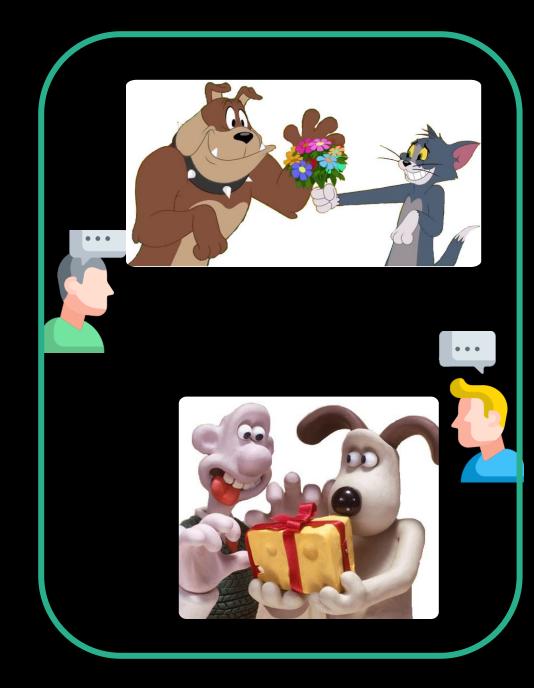


Exercise & Q&A

Exercise

The 'cheese data': Simulated data based on real psycholinguistic findings on structural priming.

- Two Predictors :
 - 2-level factor "Prime"
 - (a): Tom gave Spike some flowers.
 - (b): Tom gave some flowers to Spike.
 - 2-level factor "communication" (video- vs audio-call)
- Binary outcome: (a) or (b)?
 - (a) Gromit gave ... (Wallace some cheese)
 - (b) Gromit gave ... (some cheese to Wallace).



Exercise



Can you investigate the priming effect in the 'cheese data'?

• R code:

```
glmer (y \sim x1 * x2 +
	(1 + x1 * x2 | Grouping_1) +
	(1 + x1 | Grouping_2),
	data = datafilename,
	family = 'binomial'
)
```

Further Reading

Paper

Brown, VA. (2021). An Introduction to Linear Mixed-Effects Modeling in R. *Advances in Methods and Practices in Psychological Science*. 4(1). doi:10.1177/2515245920960351

Advances in Methods and Practices in Psychological Science

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An Introduction to Linear Mixed-Effects Modeling in R

Violet A. Brown View all authors and affiliations

All Articles | https://doi.org/10.1177/2515245920960351

E-Book

https://vasishth.github.io/Freq CogSci/

Shravan Vasishth, Daniel Schad, Audrey Bürki, Reinhold Kliegl

Linear Mixed Models in Linguistics and Psychology: A Comprehensive Introduction

THANK YOU

