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# Mixed-effects regression

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# This lecture

- Introduction
- Recap: multiple regression
- Mixed-effects regression analysis: explanation
- Case-study: [Influence of alcohol on L1 & L2](#)
- Conclusion

# Introduction

- Consider the following situation (taken from [Clark, 1973](#)):
  - Mr. A and Mrs. B study reading latencies of verbs and nouns
  - Each randomly selects 20 words and tests 50 subjects
  - Mr. A finds (using a sign test) **verbs** to have faster responses
  - Mrs. B finds **nouns** to have faster responses
- How is this possible?

# The language-as-fixed-effect fallacy

- The problem is that Mr. A and Mrs. B disregard the (huge) variability in the words
  - Mr. A included a difficult noun, but Mrs. B included a difficult verb
  - Their set of words does not constitute the complete population of nouns and verbs, therefore their results are limited to **their words**
- This is known as the language-as-fixed-effect fallacy (LAFEF)
  - **Fixed-effect factors** have repeatable and a small number of levels
  - Word is a **random-effect** factor (a non-repeatable random sample from a larger population)

# Why linguists are not always good statisticians

- LAFEF occurs frequently in linguistic research until the 1970's
  - Many reported significant results are wrong (the method is anti-conservative)!
- [Clark \(1973\)](#) combined a by-subject ( $F_1$ ) analysis and by-item ( $F_2$ ) analysis in measure *min F'*
  - Results are significant and generalizable across subjects and items when *min F'* is significant
  - Unfortunately many researchers (>50%!) incorrectly interpreted this study and may report wrong results ([Raaijmakers et al., 1999](#))
  - E.g., they only use  $F_1$  and  $F_2$  and not *min F'* or they use  $F_2$  while unnecessary (e.g., counterbalanced design)

# Our problems solved...

- Apparently, analyzing this type of data is difficult...
- Fortunately, using mixed-effects regression models solves these problems!
  - The method is easier than using the approach of [Clark \(1973\)](#)
  - Results can be generalized across subjects and items
  - Mixed-effects models are robust to missing data ([Baayen, 2008](#), p. 266)
  - We can easily test if it is necessary to treat item as a random effect
  - No balanced design necessary (as in repeated-measures ANOVA)!
- But first some words about regression...

# Recap: multiple regression (1)

- Multiple regression: predict one numerical variable on the basis of other independent variables (numerical or categorical)
- We can write a regression formula as  $y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \epsilon$ 
  - $y$ : (value of the) dependent variable
  - $x_i$ : (value of the) predictor
  - $\beta_0$ : **intercept**, value of  $y$  when all  $x_i = 0$
  - $\beta_i$ : **slope**, change in  $y$  when the value of  $x_i$  increases with 1
  - $\epsilon$ : residuals, difference between observed values and predicted (fitted) values

# Recap: multiple regression (2)

- Factorial predictors are (automatically) represented by binary-valued predictors:  $x_i = 0$  (reference level) or  $x_i = 1$  (alternative level)
  - Factor with  $n$  levels:  $n - 1$  binary predictors
  - Interpretation of factorial  $\beta_i$ : change from reference to alternative level
- Example of regression formula:
  - Predict the reaction time of a subject on the basis of word frequency, word length, and subject age:  $RT = 200 - 5WF + 3WL + 10SA$

# Mixed-effects regression modeling: introduction

- Mixed-effects regression distinguishes **fixed effects** and **random-effect** factors
- Fixed effects:
  - All numerical predictors
  - Factorial predictors with a repeatable and small number of levels (e.g., Sex)
- Random-effect **factors**:
  - Only factorial predictors!
  - Levels are a non-repeatable **random sample** from a larger population
  - Generally a large number of levels (e.g., Subject, Item)

# What are random-effects factors?

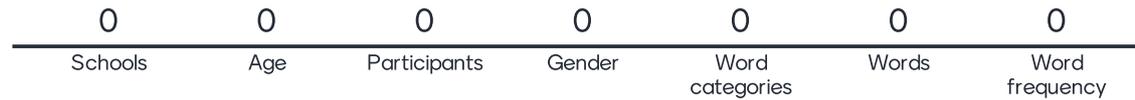
- Random-effect factors are factors which are likely to introduce systematic variation (here: **subject** and **item**)
  - Some **subjects** have a slow response (RT), while others are fast  
= **Random Intercept for Subject** (i.e.  $\beta_0$  varies per subject)
  - Some **items** are easy to recognize, others hard  
= **Random Intercept for Item** (i.e.  $\beta_0$  varies per item)
  - The effect of item frequency on RT might be higher for one **subject** than another: e.g., non-native participants might benefit more from frequent words than native participants  
= **Random Slope for Item Frequency per Subject** (i.e.  $\beta_{WF}$  varies per subject)
  - The effect of subject age on RT might be different for one **item** than another: e.g., modern words might be recognized faster by younger participants  
= **Random Slope for Subject Age per Item** (i.e.  $\beta_{SA}$  varies per item)
- Note that it is **essential** to test for random slopes!

# Question 1

Go to [www.menti.com/ebdd7c](https://www.menti.com/ebdd7c)

## What are examples of random effect factors?

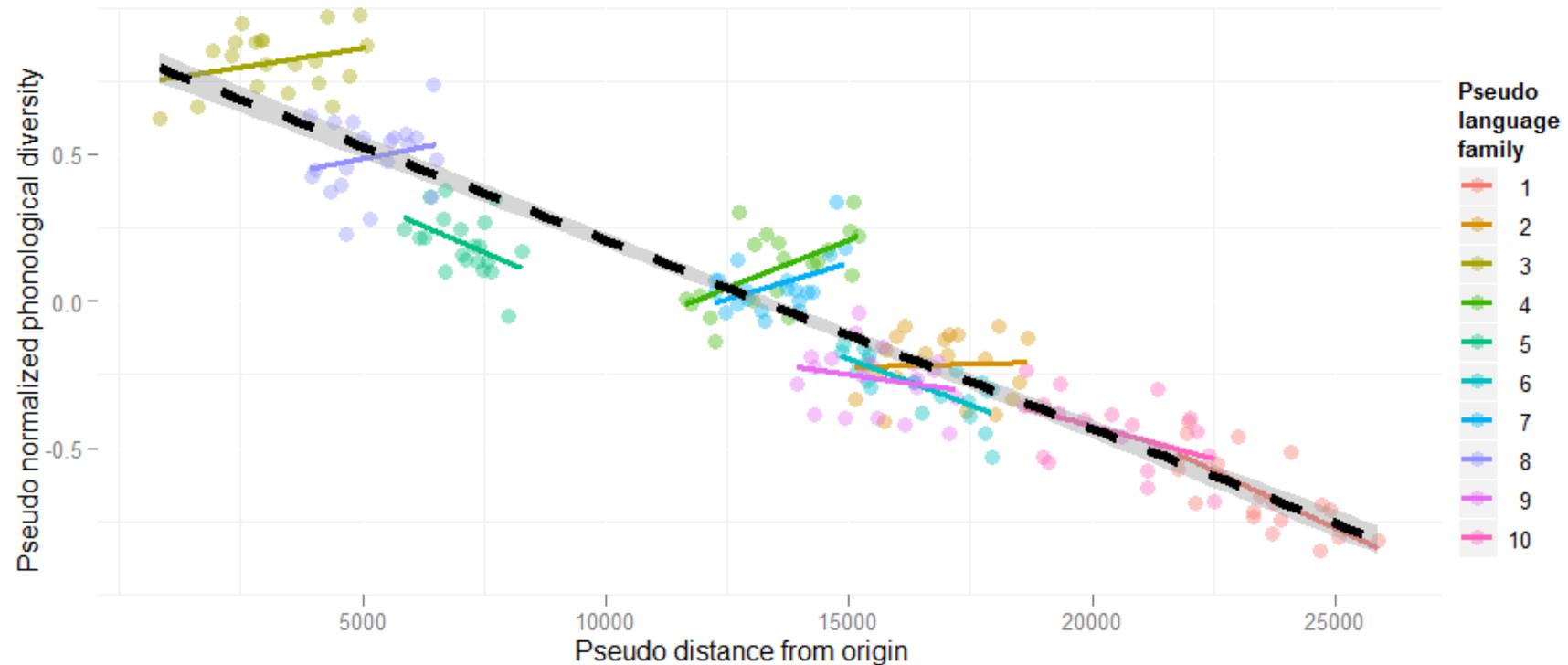
 Mentimeter



Press **ENTER** to show correct



# Random slopes are necessary!



		Estimate	Std. Error	t value	Pr(> t )
Linear regression	DistOrigin	-6.418e-05	1.808e-06	-35.49	<2e-16
+ Random intercepts	DistOrigin	-2.224e-05	6.863e-06	-3.240	<0.001
+ Random slopes	DistOrigin	-1.478e-05	1.519e-05	-0.973	n.s.

(This example is explained at the [HLP/Jaeger lab blog](#))

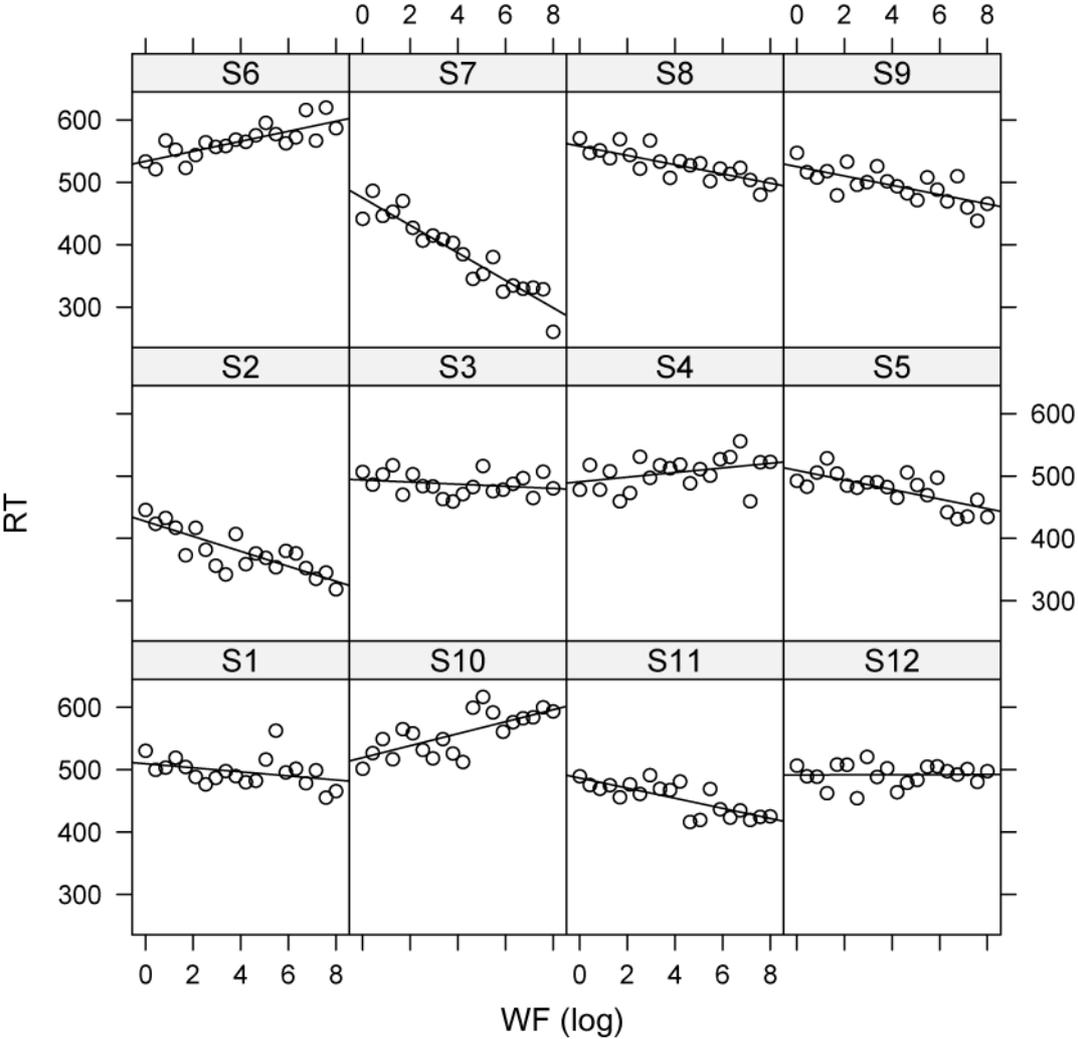
# Modeling the variance structure

- Mixed-effects regression allow us to use random intercepts and slopes (i.e. adjustments to the population intercept and slopes) to include the variance structure in our data
  - Parsimony: a single parameter (standard deviation) models this variation for every random slope or intercept (a normal distribution with mean 0 is assumed)
  - The slope and intercept adjustments are **B**est **L**inear **U**nbiased **P**redictors
  - Model comparison determines the inclusion of random intercepts and slopes
- Mixed-effects regression is only required when each level of the random-effect factor has multiple observations (e.g., participants respond to multiple items)

# Specific models for every observation

- $RT = 200 - 5WF + 3WL + 10SA$  (general model)
  - The intercepts and slopes may vary (according to the estimated variation for each parameter) and this influences the word- and subject-specific values
- $RT = 400 - 5WF + 3WL - 2SA$  (word: scythe)
- $RT = 300 - 5WF + 3WL + 15SA$  (word: twitter)
- $RT = 300 - 7WF + 3WL + 10SA$  (subject: non-native)
- $RT = 150 - 5WF + 3WL + 10SA$  (subject: fast)
- And it is not hard to use!
  - `lmer( RT ~ WF + WL + SA + (1+SA|Wrd) + (1+WF|Subj) )`
  - (`lmer` automatically discovers random-effects structure: nested/crossed)

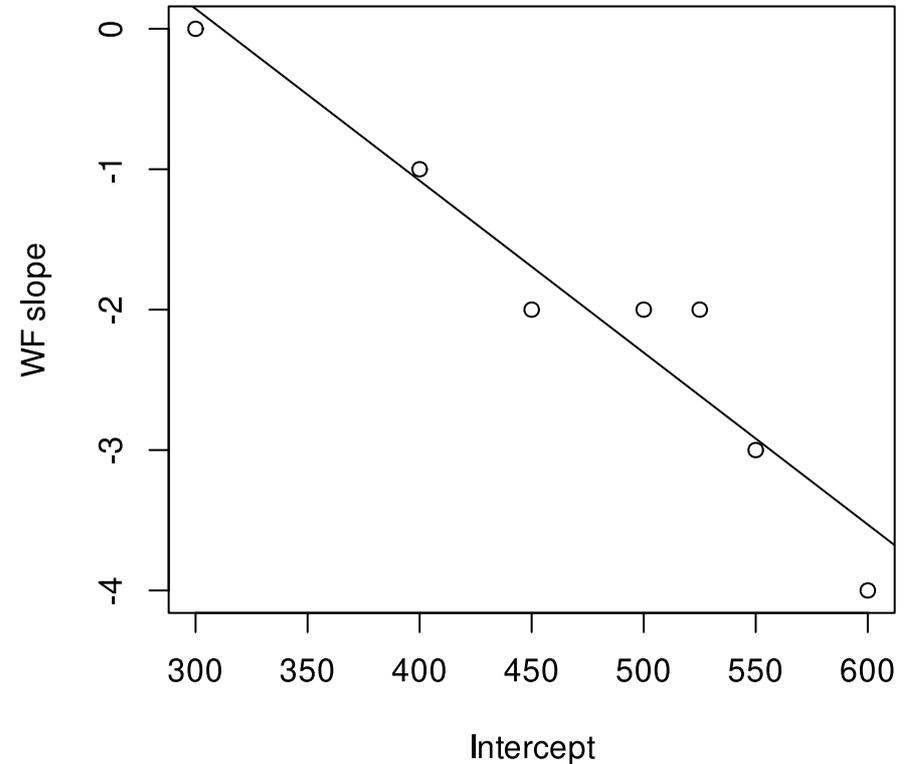
# Variability in intercept and slope



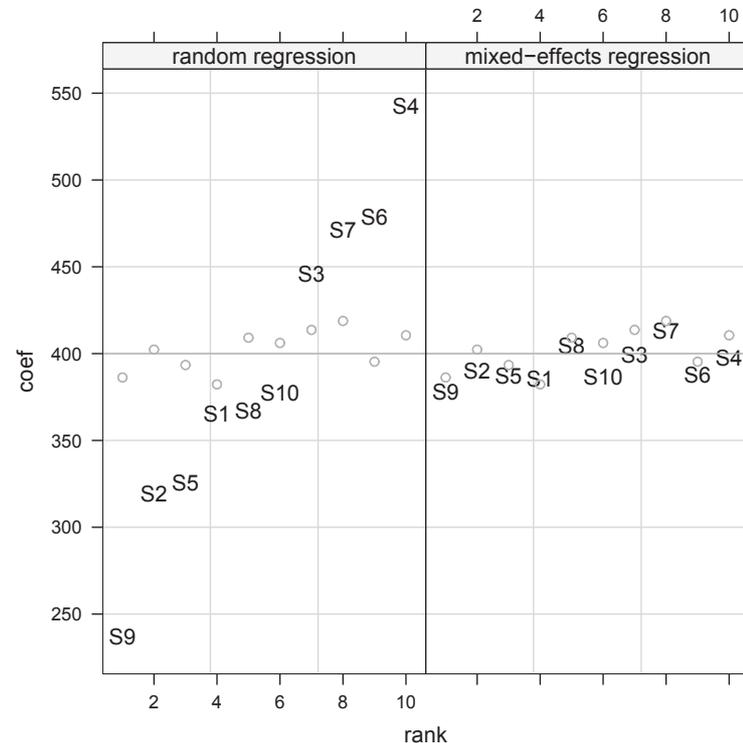
# Random slopes and intercepts may be (cor)related

- For example:

SUBJECT	INTERCEPT	WF SLOPE
S1	525	-2
S2	400	-1
S3	500	-2
S4	550	-3
S5	450	-2
S6	600	-4
S7	300	0



# BLUPs of `lmer` do not suffer from shrinkage



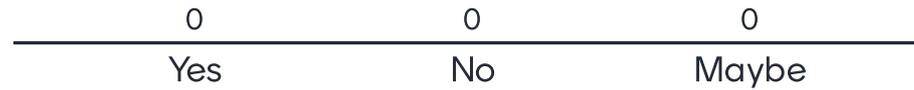
- The BLUPS (i.e. adjustment to the model estimates per item/participant) are close to the real adjustments, as `lmer` takes into account regression towards the mean (fast subjects will be slower next time, and slow subjects will be faster) thereby avoiding overfitting and improving prediction - see [Efron & Morris \(1977\)](#)

# Question 2

Go to [www.menti.com/ebdd7c](https://www.menti.com/ebdd7c)

Do you include a random slope which improves the model, if your primary effect becomes n.s.?

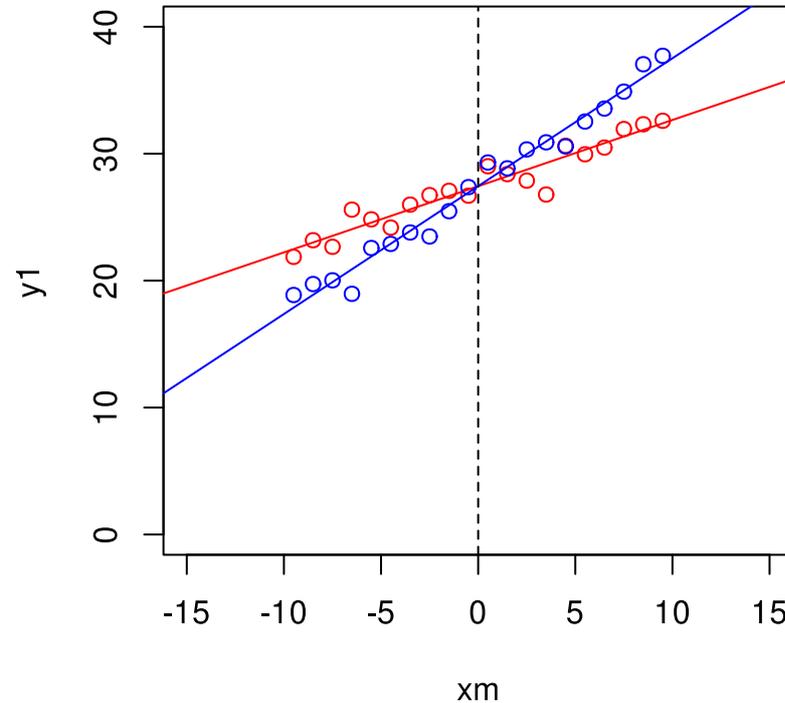
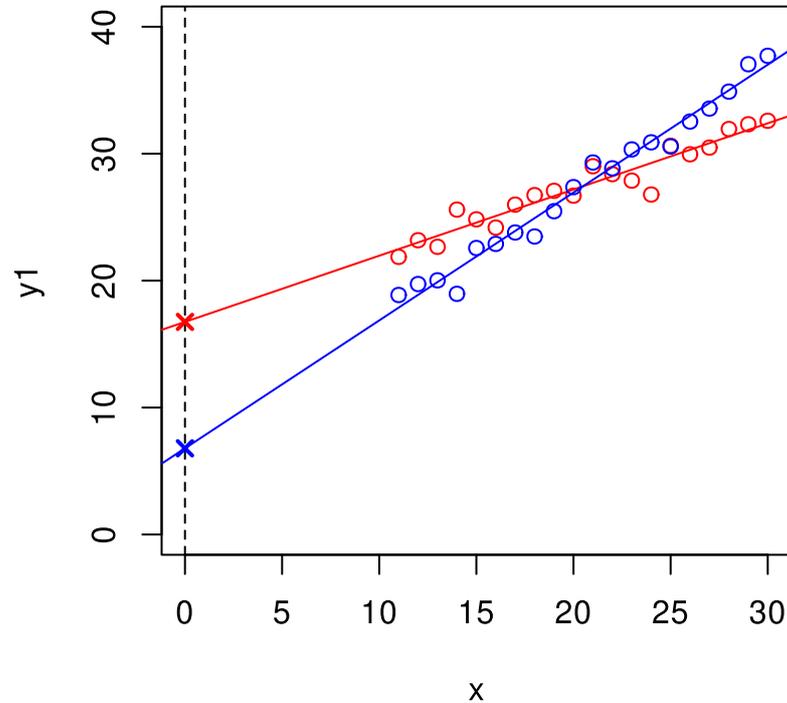
 Mentimeter



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# Center your variables (i.e. subtract the mean)!



- Otherwise random slopes and intercepts may show a spurious correlation
- Also helps the interpretation of interactions (see [this lecture](#))

# Mixed-effects regression assumptions

- Independent observations within each level of the random-effect factor
- Relation between dependent and independent variables linear
- No strong multicollinearity
- Residuals are not autocorrelated
- Homoscedasticity of variance in residuals
- Residuals are normally distributed
- (Similar assumptions as for [regression](#))

# Question 3

Go to [www.menti.com/ebdd7c](https://www.menti.com/ebdd7c)

Does the dependent variable have to be normally distributed?

 Mentimeter

0	0	0	0
Yes	No	It depends	?



Press **ENTER** to show correct



# Model criticism

- Check the distribution of residuals: if not normally distributed and/or heteroscedastic residuals then transform dependent variable or use generalized linear mixed-effects regression modeling
- Check outlier characteristics and refit the model when large outliers are excluded to verify that your effects are not 'carried' by these outliers
  - See [regression lecture](#)

# Model selection I

- “The data analyst knows more than the computer.” ([Henderson & Velleman, 1981](#))
  - Get to know your data!
- There is no adequate *automatic* procedure to find the best model
- Fitting the most complex random effects structure ([Barr et al., 2013](#)) is mostly **not** a good idea
  - The model may not converge, or is uninterpretable (see [Baayen et al., 2017](#))
  - The power of your method is reduced ([Matuschek et al., 2017](#))

# Model selection II

- My stepwise variable-selection procedure (for **exploratory analysis**):
  - Include random intercepts
  - Add other potential explanatory variables one-by-one
  - Insignificant predictors are dropped
  - Test predictors for inclusion which were excluded at an earlier stage
  - Test possible interactions (don't make it too complex)
  - Try to **break** the model by including significant predictors as random slopes
  - Only choose a more complex model if supported by model comparison

# Model selection III

- For a hypothesis-driven analysis, stepwise selection is **problematic**
  - Confidence intervals too narrow:  $p$ -values too low (multiple comparisons)
  - See, e.g., [Burnham & Anderson \(2002\)](#)
- Solutions:
  - Careful specification of potential *a priori* models lining up with the hypotheses (including optimal random-effects structure) and evaluating only these models
    - This may be followed by an exploratory procedure
  - Validating a stepwise procedure via cross validation (e.g., bootstrap analysis)

# Case study: influence of alcohol on L1 and L2



# Case study: influence of alcohol on L1 and L2

- Reported in [Wieling et al. \(2019\)](#) and [Offrede et al. \(2020\)](#)
- Assess influence of **alcohol** (BAC) on L1 (clarity) vs. L2 (nativeness) ratings
- Prediction: higher BAC has negative effect on L1, but positive effect on L2 nativeness (based on [Renner et al., 2018](#))
- ~80 recordings from native Dutch speakers included (all BACs < 0.8, no drugs)
- Dutch ratings were given by >100 native (sober) Dutch speakers (at Lowlands)
- English ratings were given by >100 native American English speakers (online)
- Dependent variable is  $z$ -transformed rating (5-point Likert scale)
- Numerical variables are centered (not  $z$ -transformed)

# Dataset

```
load("lls.rda")  
head(lls)
```

```
#      SID  BAC Sex BirthYear L2cnt SelfEN LivedEN L2anxiety  Edu      LID Lang Rating  
# 1 S0045188-17 0.392  F      -5.66 -1.04  0.463      Y      0.357 0.817 L0637009  NL  0.946  
# 2 S0045188-17 0.392  F      -5.66 -1.04  0.463      Y      0.357 0.817  L196  EN  0.330  
# 3 S0045188-17 0.392  F      -5.66 -1.04  0.463      Y      0.357 0.817  L86  EN -0.298  
# 4 S0045188-17 0.392  F      -5.66 -1.04  0.463      Y      0.357 0.817 L0614758  NL  0.325  
# 5 S0045188-17 0.392  F      -5.66 -1.04  0.463      Y      0.357 0.817  L220  EN -0.435  
# 6 S0045188-17 0.392  F      -5.66 -1.04  0.463      Y      0.357 0.817  L225  EN -0.239
```

# Fitting our first model

(fitted in R version 4.2.2 Patched (2022-11-10 r83330), **lme4** version 1.1.31)

```
library(lme4)
m <- lmer(Rating ~ BAC + (1 | SID) + (1 | LID), data = lls) # does not converge
```

```
# boundary (singular) fit: see help('isSingular')
```

```
summary(m)$coef # show fixed effects
```

```
#           Estimate Std. Error t value
# (Intercept)    0.144    0.0638   2.252
# BAC            -0.221    0.3262  -0.677
```

```
summary(m)$varcor # show random-effects part only: no variability per listener (due to z-scaling)
```

```
# Groups   Name          Std.Dev.
# LID      (Intercept) 0.000
# SID      (Intercept) 0.560
# Residual                    0.738
```

# Evaluating our hypothesis (1)

(note: random intercept for LID excluded)

```
m1 <- lmer(Rating ~ Lang * BAC + (1 | SID), data = lls)
```

- This model represents our **hypothesis test** (but likely without the correct random-effects structure)

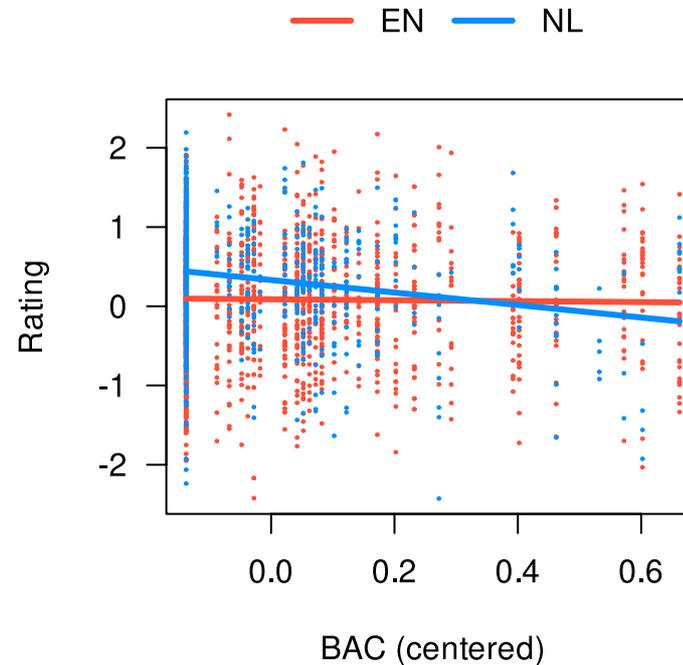
# Results (likely overestimating significance)

```
summary(ml, cor = F) # suppress expected correlation of regression coefficients; Note: |t| > 2 => p < .05 (N > 100)
```

```
# Linear mixed model fit by REML ['lmerMod']
# Formula: Rating ~ Lang * BAC + (1 | SID)
#   Data: lls
#
# REML criterion at convergence: 5853
#
# Scaled residuals:
#   Min      1Q  Median      3Q      Max
# -3.668 -0.647  0.032  0.707  3.187
#
# Random effects:
#   Groups   Name                Variance Std.Dev.
#   SID      (Intercept)  0.305    0.552
#   Residual                    0.533    0.730
# Number of obs: 2541, groups:  SID, 82
#
# Fixed effects:
#              Estimate Std. Error t value
# (Intercept)   0.0883    0.0635   1.39
# LangNL        0.2430    0.0358   6.78
# BAC           -0.0592    0.3255  -0.18
# LangNL:BAC   -0.7286    0.1938  -3.76
```

# Visualization helps interpretation

```
library(visreg)
visreg(m1, "BAC", by = "Lang", overlay = T, xlab = "BAC (centered)", ylab = "Rating")
```



- Interpretation: a higher BAC has a negative effect on L1, but not L2

# Is the added interaction an improvement?

```
m0 <- lmer(Rating ~ Lang + (1 | SID), data = lls) # comparison: best simpler model (BAC n.s.)
anova(m0, m1) # interaction necessary
```

```
# Data: lls
# Models:
# m0: Rating ~ Lang + (1 | SID)
# m1: Rating ~ Lang * BAC + (1 | SID)
#   npar  AIC  BIC logLik deviance Chisq Df Pr(>Chisq)
# m0    4 5866 5889 -2929     5858
# m1    6 5855 5890 -2921     5843  14.7  2   0.00063 ***
# ---
# Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# Evaluating our hypothesis (2): adding a random slope

```
m2 <- lmer(Rating ~ Lang * BAC + (1 + Lang | SID), data = lls)
anova(m1, m2, refit = FALSE) # random slope necessary? (REML model comparison)
```

```
# Data: lls
# Models:
# m1: Rating ~ Lang * BAC + (1 | SID)
# m2: Rating ~ Lang * BAC + (1 + Lang | SID)
#   npar  AIC  BIC logLik deviance Chisq Df Pr(>Chisq)
# m1     6 5865 5900  -2927     5853
# m2     8 5626 5672  -2805     5610   244  2   <2e-16 ***
# ---
# Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

- This model represents the **appropriate hypothesis test** as it includes the adequate random-effects structure

# Intermezzo: ML or REML model comparison?

- Restricted Maximum Likelihood (REML; default `lmer` fitting method) is appropriate when comparing 2 models differing **exclusively** in the random-effects structure
  - The REML score depends on which fixed effects are in the model, so REML values are not comparable if the fixed effects change
  - As REML is considered to give better estimates for the random effects, it is recommended to fit your final model (for reporting and inference) using REML
- ML is appropriate when comparing models differing **only** in the fixed effects
  - **anova** automatically refits the models with ML, unless **refit=F** is specified
- Never compare 2 models differing in both fixed and random effects
- Only compare models which differ minimally (e.g., only in 1 random slope)

# Result of hypothesis test

```
summary(m2, cor = F)$coef # show only fixed effects of model with random slope
```

#	Estimate	Std. Error	t value
# (Intercept)	0.0988	0.0747	1.322
# LangNL	0.2544	0.0815	3.121
# BAC	0.0994	0.3925	0.253
# LangNL:BAC	-0.9875	0.4265	-2.316

- Note the higher  $|t|$ -values in the model without the random slope:

```
summary(m1, cor = F)$coef # results of model without random slope
```

#	Estimate	Std. Error	t value
# (Intercept)	0.0883	0.0635	1.391
# LangNL	0.2430	0.0358	6.785
# BAC	-0.0592	0.3255	-0.182
# LangNL:BAC	-0.7286	0.1938	-3.759

# Random effects: correlation parameter

```
summary(m2, cor = F)$varcor # show only random effects of model with random slope
```

```
# Groups   Name          Std.Dev. Corr
# SID      (Intercept) 0.653
#          LangNL    0.653   -0.83
# Residual                0.683
```

- Note the absence of the random slope in the random-intercept-only model:

```
summary(m1, cor = F)$varcor # results of model without random slope
```

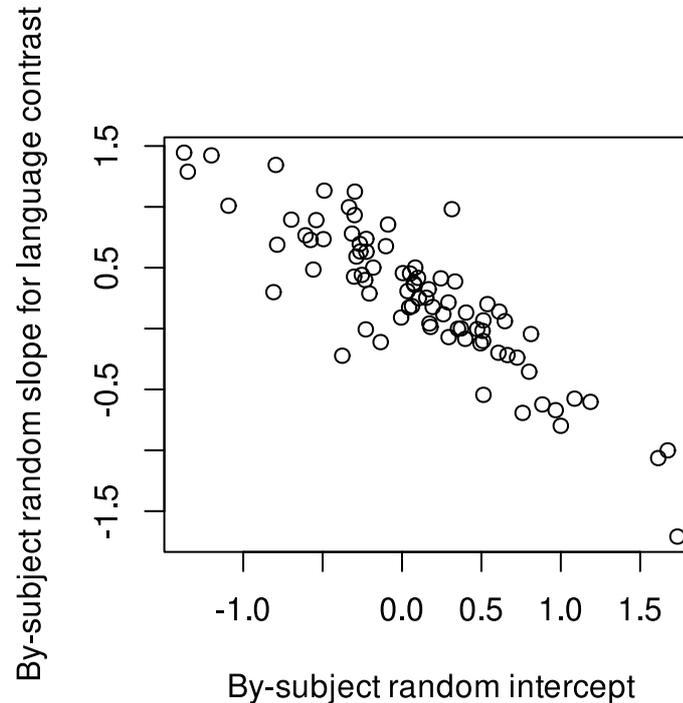
```
# Groups   Name          Std.Dev.
# SID      (Intercept) 0.552
# Residual                0.730
```

# A note about random-effect correlation parameters

- In the previous model a correlation is assumed between random intercept and slope:  $(1+X|RF)$ 
  - Any terms before the  $|$  are assumed to be correlated and a correlation parameter is calculated for each pair
- To exclude the correlation parameter, a simpler model can be specified as follows:  $(1|RF) + (0+X|RF)$  (alternatively:  $(1+X||RF)$ )
  - Model comparison (with `refit=FALSE`) is used to determine the best model
- Factorial predictors (such as `Lang`) should **always** be grouped with the intercept:  $(1+Lang|SID)$ , but not:  $(1|SID) + (0 + Lang|SID)$

# Interpretation of correlation parameter

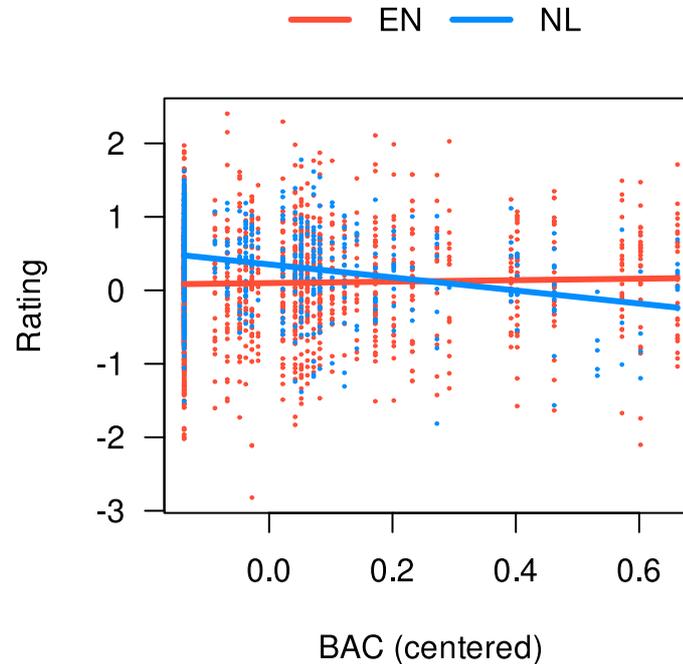
```
plot(coef(m2)$SID[, "(Intercept)"], coef(m2)$SID[, "LangNL"], xlab = "By-subject random intercept",  
     ylab = "By-subject random slope for language contrast")
```



- Interpretation: speakers with higher ratings for English (intercept), have lower ratings for Dutch and vice versa (however: categorical variables are not centered)

# Visualizing the results of the hypothesis test

```
visreg(m2, "BAC", by = "Lang", overlay = T, xlab = "BAC (centered)", ylab = "Rating")
```



- Interpretation as before: a higher BAC has a negative effect on L1, but not on L2

# Alternative summary

```
m2b <- lmer(Rating ~ Lang:BAC + Lang + (1 + Lang | SID), data = lls) # instead of Lang * BAC
summary(m2b)$coef # effect of BAC per language separately
```

#	Estimate	Std. Error	t value
# (Intercept)	0.0988	0.0747	1.322
# LangNL	0.2544	0.0815	3.121
# LangEN:BAC	0.0994	0.3925	0.253
# LangNL:BAC	-0.8882	0.2684	-3.309

```
summary(m2)$coef # original model: effect of BAC for NL vs EN
```

#	Estimate	Std. Error	t value
# (Intercept)	0.0988	0.0747	1.322
# LangNL	0.2544	0.0815	3.121
# BAC	0.0994	0.3925	0.253
# LangNL:BAC	-0.9875	0.4265	-2.316

# Exploratory analysis

- Hypothesis seems partially supported
- But this may be caused by other variables:
  - L2 anxiety
  - L2 self rating
  - (Other control variables)
- It may also be caused by the presence of atypical outliers
- Next step: **exploratory analysis**

# Effect of speaker's sex?

```
m3 <- lmer(Rating ~ Lang * BAC + Sex + (1 + Lang | SID), data = lls)
summary(m3)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.1347	0.0870	1.549
# LangNL	0.2532	0.0815	3.107
# BAC	0.1325	0.3987	0.332
# SexM	-0.0815	0.0973	-0.838
# LangNL:BAC	-0.9801	0.4265	-2.298

```
anova(m2, m3)$P[2] # p-value from anova: Sex not necessary (also not in interaction; not shown)
```

```
# [1] 0.404
```

# Effect of speaker's year of birth?

```
m3 <- lmer(Rating ~ Lang * BAC + BirthYear + (1 + Lang | SID), data = lls)
summary(m3)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.09969	0.07472	1.334
# LangNL	0.25462	0.08171	3.116
# BAC	0.08313	0.39269	0.212
# BirthYear	0.00603	0.00495	1.217
# LangNL:BAC	-0.99645	0.42764	-2.330

```
anova(m2, m3)$P[2] # BirthYear not necessary (also not in interaction with language; not shown)
```

```
# [1] 0.217
```

# Effect of speaker's education?

```
m3 <- lmer(Rating ~ Lang * BAC + Edu + (1 + Lang | SID), data = lls)
summary(m3)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.102	0.0709	1.443
# LangNL	0.255	0.0817	3.115
# BAC	0.200	0.3734	0.536
# Edu	0.112	0.0341	3.288
# LangNL:BAC	-1.034	0.4266	-2.424

```
anova(m2, m3)$P[2] # Education necessary (but not in interaction with language; not shown)
```

```
# [1] 0.00139
```

# Effect of living in an English-speaking country?

```
m4 <- lmer(Rating ~ Lang * BAC + Edu + LivedEN + (1 + Lang | SID), data = lls)
summary(m4)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.0831	0.0732	1.134
# LangNL	0.2526	0.0817	3.091
# BAC	0.2027	0.3708	0.546
# Edu	0.1126	0.0342	3.290
# LivedENY	0.1116	0.1147	0.973
# LangNL:BAC	-1.0368	0.4264	-2.432

```
anova(m3, m4)$P[2] # LivedEN not necessary (also not in interaction with language; not shown)
```

```
# [1] 0.327
```

# Effect of self-rated English proficiency?

```
m4 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN + (1 + Lang | SID), data = lls)
summary(m4)$coef
```

```
#           Estimate Std. Error t value
# (Intercept)  0.1077    0.0644   1.671
# LangNL       0.2491    0.0824   3.021
# BAC          0.1675    0.3401   0.493
# Edu          0.0758    0.0343   2.211
# SelfEN       0.1299    0.0370   3.507
# LangNL:BAC  -1.0967    0.4283  -2.561
```

```
anova(m3, m4)$P[2] # SelfEN necessary
```

```
# [1] 0.00135
```

# Self-rated English proficiency: interaction?

```
m5 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN * Lang + (1 + Lang | SID), data = lls)
summary(m5)$coef
```

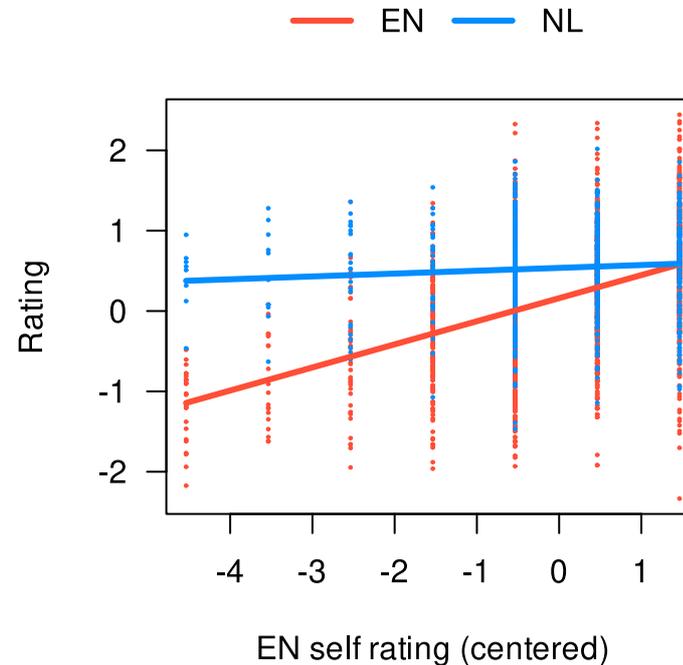
#	Estimate	Std. Error	t value
# (Intercept)	0.1087	0.0612	1.775
# LangNL	0.2474	0.0753	3.285
# BAC	0.0665	0.3236	0.206
# Edu	0.0741	0.0344	2.157
# SelfEN	0.2874	0.0541	5.309
# LangNL:BAC	-0.9385	0.3933	-2.386
# LangNL:SelfEN	-0.2519	0.0630	-3.996

```
anova(m4, m5)$P[2] # interaction necessary
```

```
# [1] 0.000103
```

# Visual interpretation of interaction

```
visreg(m5, "SelfEN", by = "Lang", overlay = T, xlab = "EN self rating (centered)", ylab = "Rating")
```



- Interpretation: a higher self-rated English proficiency is only predictive for the English ratings, not the Dutch ratings (unsurprisingly)

# Effect of L2 count?

```
m6 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN * Lang + L2cnt + (1 + Lang | SID), data = lls)
summary(m6)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.1080	0.0599	1.802
# LangNL	0.2500	0.0755	3.314
# BAC	0.1092	0.3172	0.344
# Edu	0.0635	0.0339	1.875
# SelfEN	0.2787	0.0531	5.246
# L2cnt	0.1012	0.0467	2.168
# LangNL:BAC	-0.9356	0.3937	-2.377
# LangNL:SelfEN	-0.2499	0.0632	-3.955

```
anova(m5, m6)$P[2] # L2cnt necessary
```

```
# [1] 0.028
```

- Note that the variable **Edu** now does not have an absolute  $t$ -value  $> 2$  anymore, but we retain it as it is close to significance ( $p \approx 0.06$  based on **anova**)

# For both English and Dutch, or only English?

```
m7 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN * Lang + L2cnt * Lang + (1 + Lang | SID),  
  data = lls)  
summary(m7)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.1076	0.0601	1.792
# LangNL	0.2501	0.0758	3.300
# BAC	0.1237	0.3183	0.389
# Edu	0.0630	0.0338	1.863
# SelfEN	0.2746	0.0536	5.127
# L2cnt	0.1340	0.0677	1.979
# LangNL:BAC	-0.9640	0.3972	-2.427
# LangNL:SelfEN	-0.2432	0.0642	-3.785
# LangNL:L2cnt	-0.0576	0.0861	-0.668

```
anova(m6, m7)$P[2] # no interaction necessary: L2cnt affects both languages similarly
```

```
# [1] 0.49
```

# Effect of L2 anxiety?

```
m7 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN * Lang + L2cnt + L2anxiety + (1 + Lang |  
  SID), data = lls)  
summary(m7)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.1098	0.0580	1.892
# LangNL	0.2453	0.0755	3.251
# BAC	0.1042	0.3072	0.339
# Edu	0.0606	0.0327	1.854
# SelfEN	0.2026	0.0599	3.384
# L2cnt	0.1202	0.0456	2.638
# L2anxiety	-0.2277	0.0918	-2.481
# LangNL:BAC	-0.9463	0.3931	-2.407
# LangNL:SelfEN	-0.2526	0.0632	-3.998

```
anova(m6, m7)$P[2] # L2anxiety necessary
```

```
# [1] 0.0118
```

# For both English and Dutch, or only English?

```
m8 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN * Lang + L2cnt + L2anxiety * Lang + (1 +  
  Lang | SID), data = lls)  
summary(m8)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.1103	0.0581	1.900
# LangNL	0.2462	0.0756	3.259
# BAC	0.0949	0.3075	0.309
# Edu	0.0607	0.0327	1.857
# SelfEN	0.1721	0.0680	2.531
# L2cnt	0.1206	0.0456	2.644
# L2anxiety	-0.3202	0.1340	-2.389
# LangNL:BAC	-0.9293	0.3940	-2.359
# LangNL:SelfEN	-0.1973	0.0861	-2.292
# LangNL:L2anxiety	0.1653	0.1747	0.946

```
anova(m7, m8)$P[2] # no interaction necessary: L2anxiety affects both languages similarly
```

```
# [1] 0.337
```

# Best model?

- Since SID is the only random-effect factor, and the characteristics are unique per SID, there are no additional random slopes to test
- All fixed-effects were tested (not all n.s. interactions were shown)
- Conclusion: **m7** is the best model!

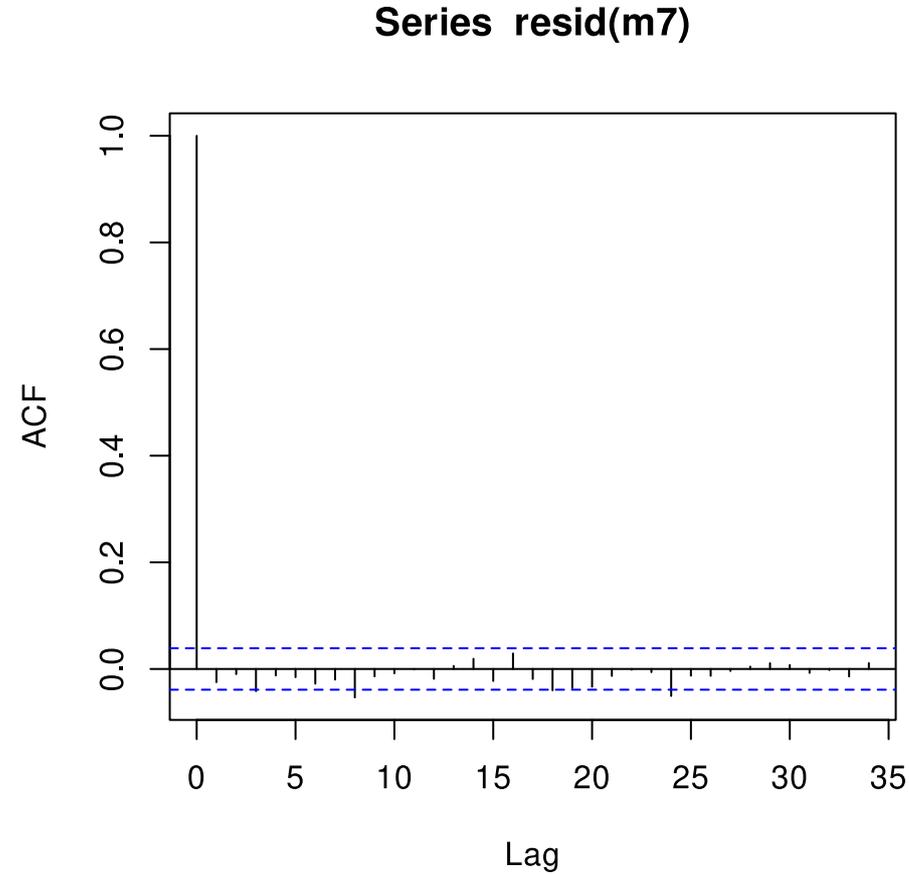
# Testing assumptions: multicollinearity

```
library(car)  
vif(m7) # Should be lower < 5 (for centered numerical variables): OK
```

```
#      Lang      BAC      Edu      SelfEN      L2cnt      L2anxiety      Lang:BAC  
#      1.00      2.32      1.19      3.40      1.08      1.94      2.30  
# Lang:SelfEN  
#      2.34
```

# Testing assumptions: autocorrelation

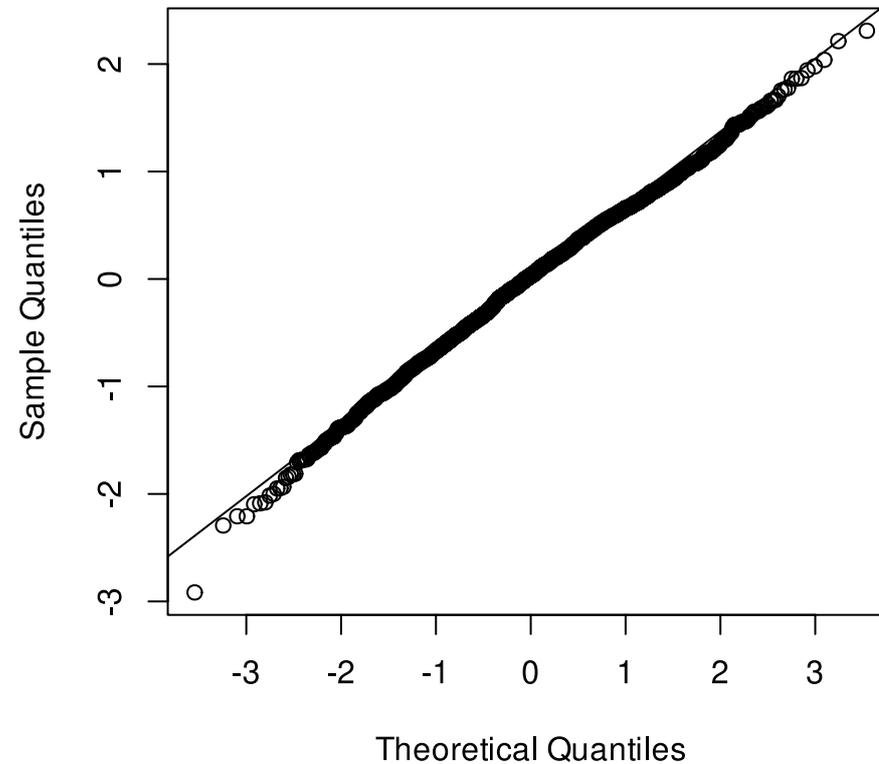
```
acf(resid(m7)) # no autocorrelation
```



# Testing assumptions: normality

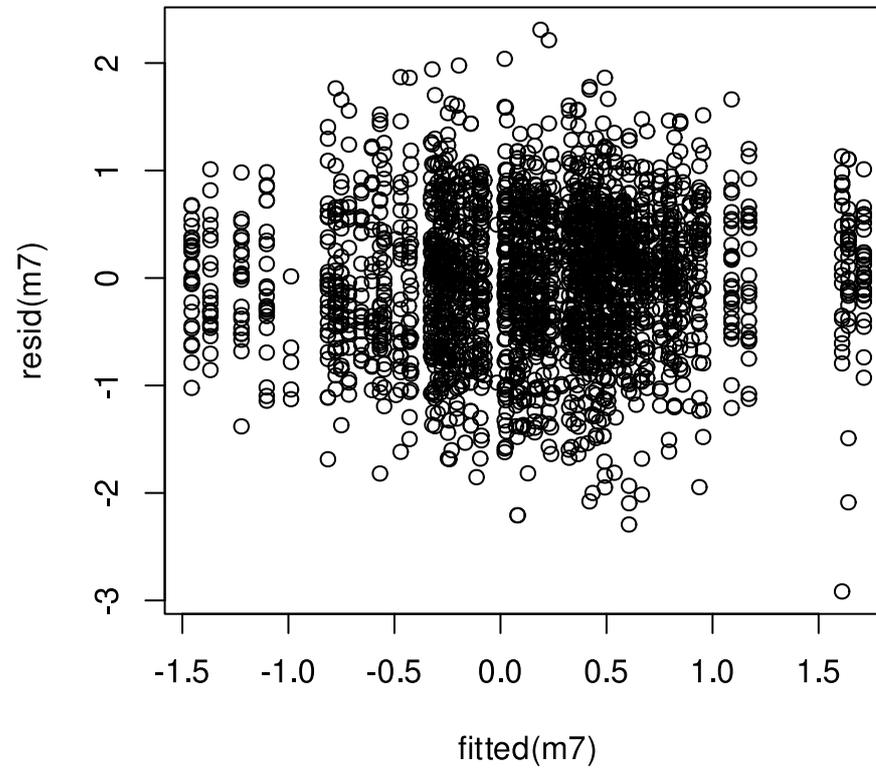
```
qqnorm(resid(m7)) # OK  
qqline(resid(m7))
```

Normal Q-Q Plot



# Testing assumptions: heteroscedasticity

```
plot(fitted(m7), resid(m7)) # some heteroscedasticity
```



# Model criticism: countering outlier influence

```
lls2 <- lls[abs(scale(resid(m7))) < 2.5, ] # trimmed model: 98.5% of original data
m7.2 <- lmer(Rating ~ Lang * BAC + Edu + SelfEN * Lang + L2cnt + L2anxiety + (1 + Lang |
  SID), data = lls2)
summary(m7.2)$coef
```

#	Estimate	Std. Error	t value
# (Intercept)	0.1161	0.0594	1.955
# LangNL	0.2493	0.0758	3.289
# BAC	0.1342	0.3145	0.427
# Edu	0.0627	0.0332	1.887
# SelfEN	0.2055	0.0612	3.360
# L2cnt	0.1245	0.0463	2.687
# L2anxiety	-0.2451	0.0933	-2.626
# LangNL:BAC	-0.9850	0.3948	-2.495
# LangNL:SelfEN	-0.2708	0.0636	-4.260

- The BAC interaction with language remains the same: significantly more negative effect of BAC on ratings for Dutch compared to English

# Question 4

Go to [www.menti.com/ebdd7c](http://www.menti.com/ebdd7c)



## Is model trimming 'cheating'?

---

0	0	0
Yes	No	?



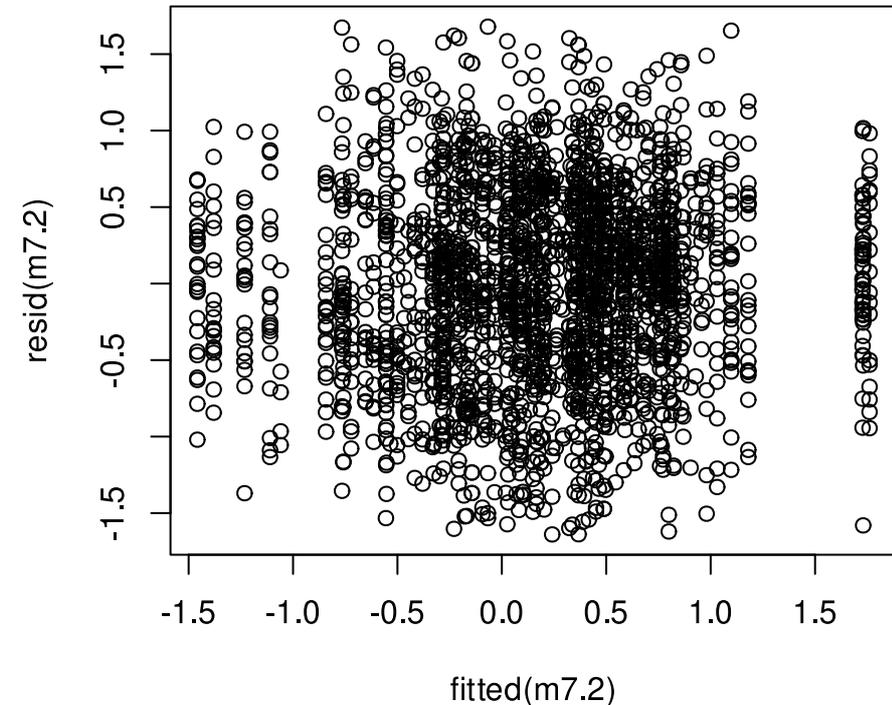
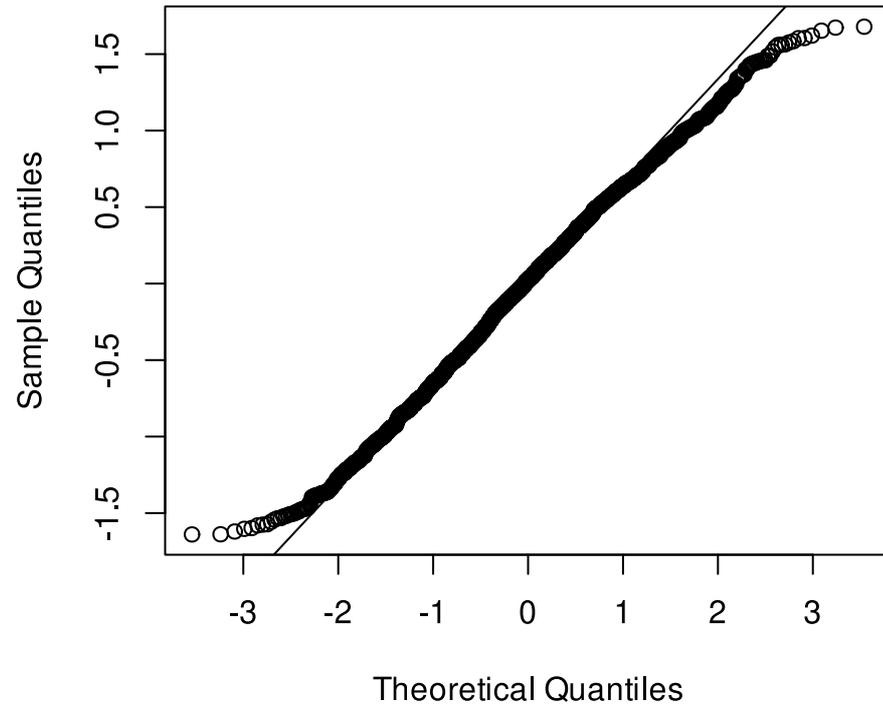
Press **ENTER** to show correct



# Checking the residuals of the trimmed model

```
par(mfrow = c(1, 2))  
qqnorm(resid(m7.2)) # reasonably OK  
qqline(resid(m7.2))  
plot(fitted(m7.2), resid(m7.2)) # better
```

Normal Q-Q Plot



# Bootstrap sampling shows similar results

```
library(boot)
(bsm7.2 <- confint(m7.2, method = "boot", nsim = 1000, level = 0.95))
```

```
#           2.5 % 97.5 %
# .sig01      0.26616 0.3904
# .sig02      0.05719 0.5966
# .sig03      0.23968 0.3649
# .sigma      0.62334 0.6590
# (Intercept) 0.16228 0.3131
# Lang1      -0.19098 -0.0537
# BAC        -0.76876 0.0527
# Edu        -0.00264 0.1271
# SelfEN     -0.01707 0.1694
# L2cnt       0.03094 0.2144
# L2anxiety  -0.42732 -0.0596
# Lang1:BAC   0.09767 0.8765
# Lang1:SelfEN 0.07578 0.1996
```

- Note that the 95% CI of the variable **Edu** does not contain 0 and therefore indicates it is a significant predictor

# Recap

- We have learned:
  - How to interpret mixed-effects regression results
  - How to use **lmer** to conduct mixed-effects regression
  - How to include random intercepts and random slopes in **lmer** and why these are **essential** when you have multiple responses per subject or item
- Associated lab session:
  - <https://www.let.rug.nl/wieling/Statistics/Mixed-Effects/lab>
    - Lab session contains additional information: how to do multiple comparisons, using other optimizers, and conducting logistic regression

# Evaluation

Go to [www.menti.com/ebdd7c](https://www.menti.com/ebdd7c)

Please provide your opinion about this lecture in  at most 3 words/phrases!



# Questions?

Thank you for your attention!

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