Cheat Sheet: generalized linear mixed-effects models

Measurement and Evaluation of HCC Systems

Scenario

Use linear mixed-effects models if you want to test the effect of several variables variables varX1, varX2, ... on a binary outcome variable varY, where the Y and some of the Xs are repeated measurements on the same unit (e.g. multiple conditions or time points per participant, multiple people per group, etc.). This cheat sheet assumes that data is in the long format, with a user id to tie together the repeated measures. (if it is not, use the melt function in the reshape2 package to fix that).

Power analysis

- The power analysis for a linear mixed-effects model is beyond the scope of this course.

Plotting

- For plotting the effects of within-subjects manipulations, see the Plotting sections of the dependent *t* test and repeated and mixed ANOVA cheat sheets.
- For plotting linear effects, see the Plotting section of the regression cheat sheet.

Pre-testing assumptions

- For between-subjects factors, refer to Pre-testing assumptions in the ANOVA and/or factorial ANOVA cheat sheets.
- For linear effects, refer to Pre-testing assumptions in the regression cheat sheet.
- Note that unlike for repeated and mixed ANOVAs, sphericity is not assumed for linear mixedeffects models.
- If your Y variable is has nominal values (e.g. "A" and "B" instead of 0 and 1), you can turn it into a numeric variable by using as.numeric(varY == "B").
- For error bars, use mean_cl_boot instead of mean_cl_normal

(optional) Preparing dummies and/or contrasts

- If one or more of your Xs are nominal variables, you need to create dummy variables or contrasts for them.
- For simple dummies, refer to the regression cheat sheet.
- For contrasts, refer to the ANOVA cheat sheet.

Running the test

- Start with the simplest possible baseline: baseline <- glm(varY ~ 1, data=data, family=binomial)</pre>
- Then add the random intercept for id: random <- glmer(varY ~ 1 + (1|id), data=data, family=binomial)</pre>
- Test whether the random intercept is needed: anova(random, baseline)

If this ANOVA is significant, you indeed need the random intercept. The χ^2 -value and its *p*-value represent the significance of the random intercept.

- Introduce your first X variable: model1 <- update(random, .~. + varX1)</pre>
- (optional) Introduce another X variable (add more Xs one by one if needed): model2 <- update(model1, .~. + varX2)</pre>
- (optional) Introduce interaction effects: model3 <- update(model2, .~. + varX1:varX2)</pre>
- Compare all models: anova(random, model1, model2, model3)
- (optional) Add a random slope for a within-subjects variable (interpretation: is the effect of varX1 different per id?):

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slope1 <- update(model3, .~. + (varX1|id))</pre>
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 Compare with the previous model to see if this random slope improves the model: anova(model3, slope1)

The χ^2 -value and its *p*-value represent the significance of the random slope.

- Get the summary and confidence intervals of your final model: summary(slope1)
- For the correct interpretation of the coefficients, refer to the logistic regression cheat sheet.
- The only additional thing is the random intercept (which has an *SD*), and optionally the random slope (which also has an *SD*) and the correlation between the two. All these measures have a confidence interval as well.

Post-testing assumptions and inspecting outliers

- For post-testing assumptions and inspecting outliers, refer to the regression cheat sheet.

Reporting

- For random intercepts and slopes: The relationship between [varY] and [VarX1 and other within-subjects variables] showed significant variance in intercept across participants, SD = x.xx (95% CI: x.xx, x.xx), $\chi^2(1) = x.xx$, p = .xxx. In addition, the slopes varied across participants, SD = x.xx (x.xx, x.xx), $\chi^2(2) = 38.87$, p = .xxx, the slopes and intercepts were significantly correlated, cor = .xx (.xx, .xx).
- For reporting further results, refer to the logistic regression cheat sheet. Note that there is no model R^2 , and model comparisons are χ^2 tests.