

Multilevel modelling

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Introduction

One of the key data manipulation tasks that must be accomplished prior to estimating several of the multilevel models (specifically contextual models and random coefficient models) is that group-level variables must be “assigned down” to the individual. To make a dataframe containing both individual and group-level variables, one typically begins with two separate dataframes. One dataframe contains individual-level data, and the other dataframe contains group-level data. By combining these two dataframes using a group identifying variable common to both, one is able to create a single data set containing both individual and group data.

Fixed effect in this context is the same coefficient for all subjects. *Random effect* is an effect that varies from group to group - this is usually just as simple as a specific intercept for the group.

The random effects give structure to the error term - if we know that errors are correlated in groups, and thus non-random, we should model this insight.

In R you write (1|group) to show that there should be a specific intercept for each group. You can add another, third, level in the same way: (1|group1) + (1|group2).

These are called mixed models since they have both fixed and random effects.

Examples in R

Look at data

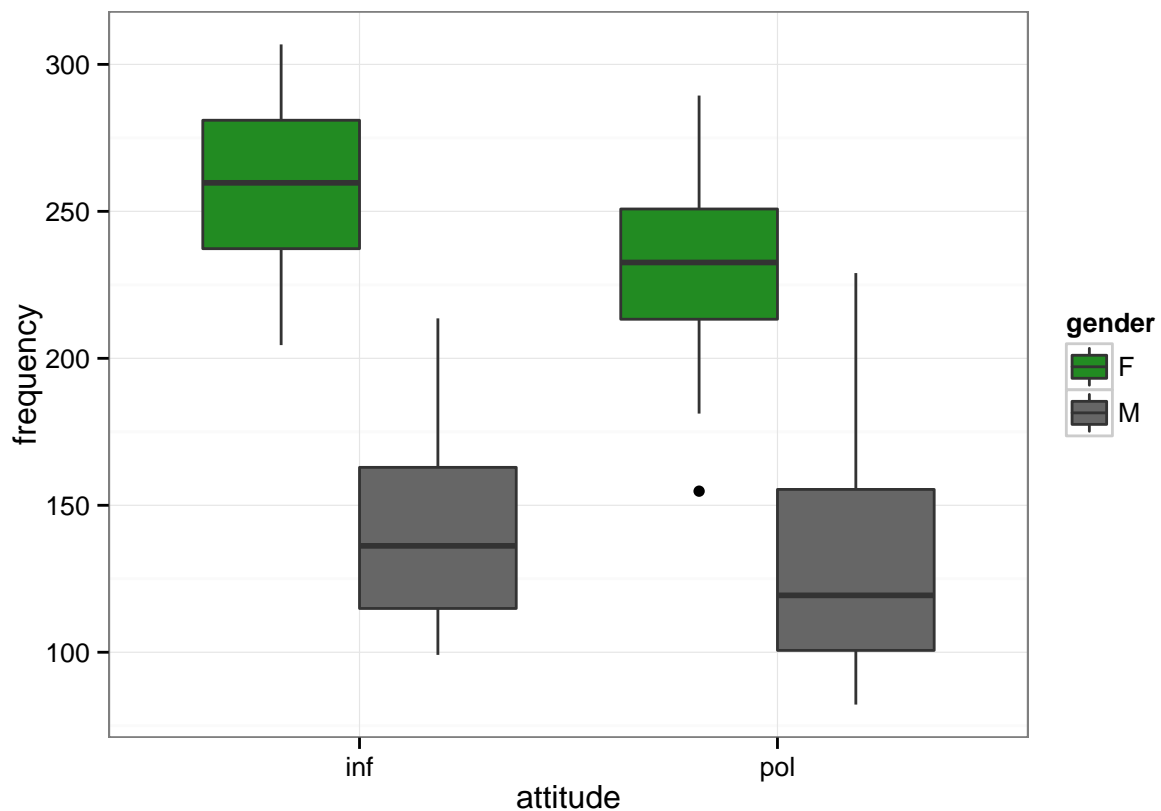
```
library(dplyr)
library(ggplot2)

politeness = read.csv("http://www.bodowinter.com/tutorial/politeness_data.csv")

head(politeness)
```

```
##   subject gender scenario attitude frequency
## 1      F1      F         1      pol      213.3
## 2      F1      F         1      inf      204.5
## 3      F1      F         2      pol      285.1
## 4      F1      F         2      inf      259.7
## 5      F1      F         3      pol      203.9
## 6      F1      F         3      inf      286.9
```

```
ggplot(politeness, aes(x=attitude, y = frequency,
                      fill = gender)) +
  geom_boxplot() +
  theme_bw() +
  scale_fill_manual(values=c("forestgreen", "grey40"))
```



Specifying some simple models

```
library(lme4)
```

```
## Loading required package: Matrix
## Loading required package: Rcpp
```

```
model1 = lmer(frequency ~ attitude + (1|subject),
              data = politeness)
```

```
summary(model1)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: frequency ~ attitude + (1 | subject)
## Data: politeness
##
## REML criterion at convergence: 804.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.2953 -0.6018 -0.2005  0.4774  3.1772
##
## Random effects:
```

```
## Groups Name Variance Std.Dev.
## subject (Intercept) 3982 63.10
## Residual 851 29.17
## Number of obs: 83, groups: subject, 6
##
## Fixed effects:
## Estimate Std. Error t value
## (Intercept) 202.588 26.151 7.747
## attitudepol -19.376 6.407 -3.024
##
## Correlation of Fixed Effects:
## (Intr)
## attitudepol -0.121
```

Compare this to

```
lm_model = lm(frequency ~ attitude, data = politeness)
summary(lm_model)
```

```
##
## Call:
## lm(formula = frequency ~ attitude, data = politeness)
##
## Residuals:
## Min 1Q Median 3Q Max
## -103.488 -62.122 9.044 51.178 105.044
##
## Coefficients:
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 202.59 10.08 20.107 <2e-16 ***
## attitudepol -18.23 14.34 -1.272 0.207
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 65.3 on 81 degrees of freedom
## (1 observation deleted due to missingness)
## Multiple R-squared: 0.01958, Adjusted R-squared: 0.007475
## F-statistic: 1.618 on 1 and 81 DF, p-value: 0.2071
```

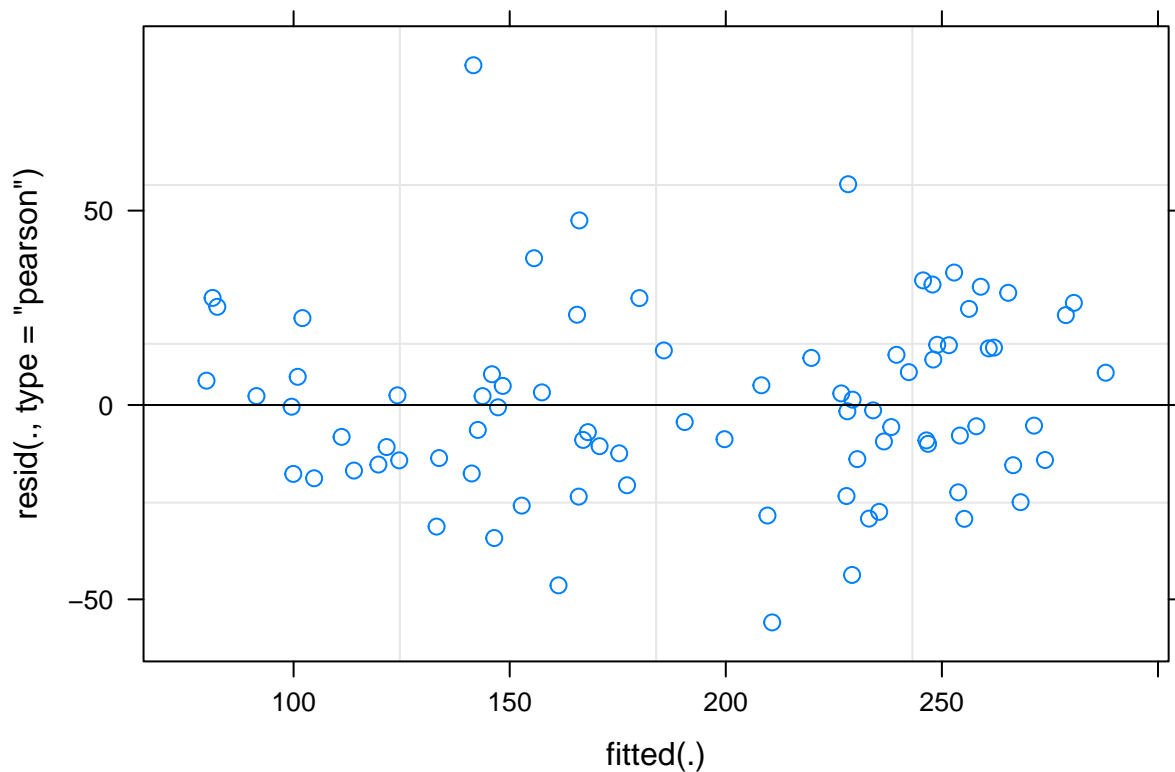
More complex model

```
model2 = lmer(frequency ~ attitude + (1|subject) +
              (1|scenario), data = politeness)
summary(model2)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: frequency ~ attitude + (1 | subject) + (1 | scenario)
## Data: politeness
##
## REML criterion at convergence: 793.5
##
## Scaled residuals:
## Min 1Q Median 3Q Max
```

```
## -2.2006 -0.5817 -0.0639  0.5625  3.4385
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   scenario (Intercept) 219     14.80
##   subject  (Intercept) 4015     63.36
##   Residual              646     25.42
## Number of obs: 83, groups:  scenario, 7; subject, 6
##
## Fixed effects:
##               Estimate Std. Error t value
## (Intercept)   202.588    26.754    7.572
## attitudepol  -19.695     5.585   -3.527
##
## Correlation of Fixed Effects:
##              (Intr)
## attitudepol -0.103
```

```
plot(model2)
```



The standard deviation in the random effects part shows how much variation there is between groups. In model2 we can see that there is a lot more variation between subjects than between the different scenarios in which the pitch was measured.

If we compare the standard linear model to the first lmer model we can see that attitude shifts from being non-significant to being clearly significant. This is both because the coefficient becomes stronger, -19.4 compared to -18.2, and because the standard error is far lower.

You can **plot** the result just like for normal lm models, at least to check that the residuals appear to lack structure.

```
# Get p-values for the fixed effects using normal approximation
coef = summary(model2) %>%
  coef() %>%
  data.frame() %>%
  mutate(pvalue = 1.96 * (1-pnorm(abs(t.value), mean=0,
                                sd=1)))
```

Basically this is just a normal approximation of the t-value and then a check of how likely that result is using the cumulative normal distribution (with mean 0 and sd 1). The more observations we have, the better this approximation will be.

Random slope model

These models have been random intercept models, the variables are assumed to have the same effects but the baseline differs.

There are also random slope models. Here the same variable is assumed to have different effects in different groups - for example, the effect of unemployment on cabinet duration might be more pronounced in some countries.

```
model_rmslopes = lmer(frequency ~ attitude + gender +
                      (1+attitude|subject) +
                      (1+attitude|scenario),
                      data = politeness,
                      REML = FALSE)

summary(model_rmslopes)
```

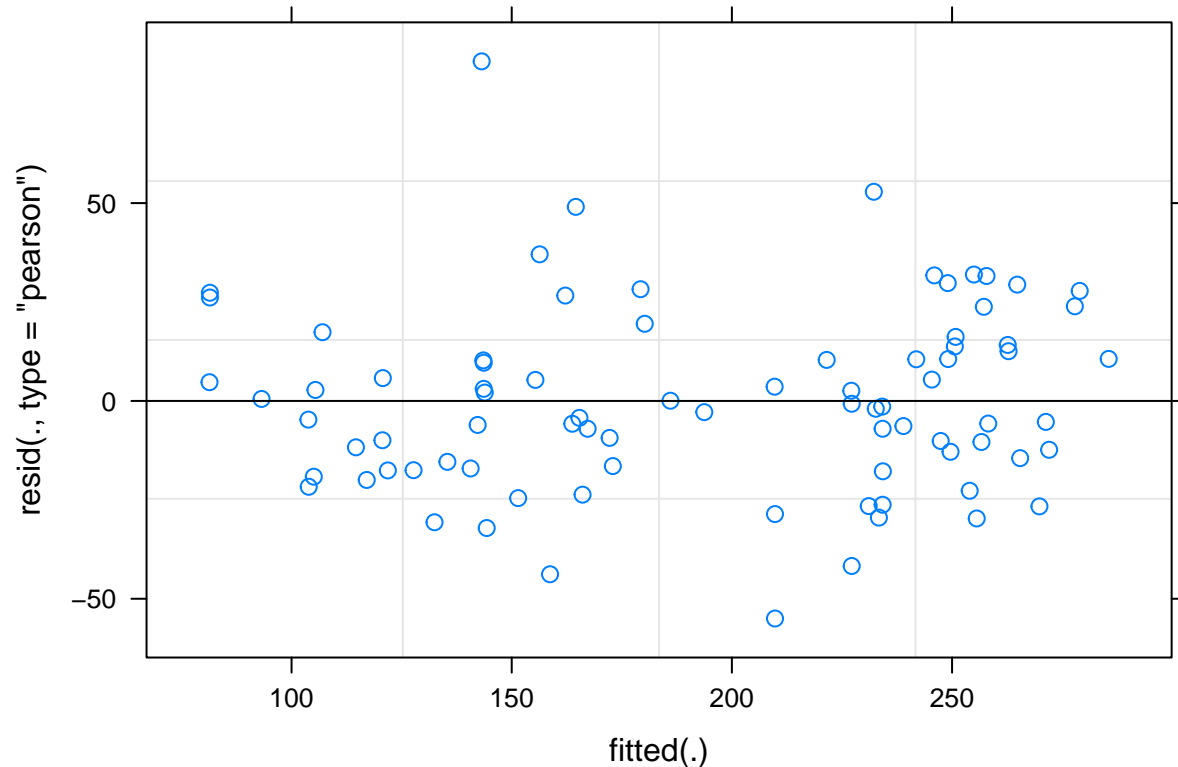
```
## Linear mixed model fit by maximum likelihood ['lmerMod']
## Formula: frequency ~ attitude + gender + (1 + attitude | subject) + (1 +
##      attitude | scenario)
##      Data: politeness
##
##      AIC      BIC    logLik deviance df.resid
##    814.9    839.1   -397.4    794.9      73
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1947 -0.6691 -0.0789  0.5256  3.4252
##
## Random effects:
##  Groups   Name                Variance Std.Dev. Corr
##  scenario (Intercept) 182.082   13.494
##           attitudepol  31.262    5.591   0.22
##  subject  (Intercept) 392.474   19.811
##           attitudepol  1.707    1.307   1.00
## Residual                627.880   25.058
## Number of obs: 83, groups:  scenario, 7; subject, 6
##
## Fixed effects:
```

```
##           Estimate Std. Error t value
## (Intercept)  257.989    13.529  19.069
## attitudepol  -19.747     5.922  -3.334
## genderM      -110.802    17.512  -6.327
##
## Correlation of Fixed Effects:
##           (Intr) atttdp
## attitudepol -0.105
## genderM     -0.647  0.003
```

```
coef(model_rmslopes)
```

```
## $scenario
##   (Intercept) attitudepol  genderM
## 1    245.2603   -20.43832 -110.8021
## 2    263.3012   -15.94386 -110.8021
## 3    269.1432   -20.63361 -110.8021
## 4    276.8309   -16.30132 -110.8021
## 5    256.0579   -19.40575 -110.8021
## 6    246.8605   -21.94816 -110.8021
## 7    248.4702   -23.55752 -110.8021
##
## $subject
##   (Intercept) attitudepol  genderM
## F1    243.8053   -20.68245 -110.8021
## F2    266.7321   -19.17028 -110.8021
## F3    260.1484   -19.60452 -110.8021
## M3    285.6958   -17.91951 -110.8021
## M4    264.1982   -19.33741 -110.8021
## M7    227.3551   -21.76744 -110.8021
##
## attr(,"class")
## [1] "coef.mer"
```

```
plot(model_rmslopes)
```



P-values

```
p_values = summary(model_rmslopes) %>%
  coef() %>%
  data.frame() %>%
  mutate(pvalue = 1.96 * (1 - pnorm(abs(t.value))))
```

The notation “(1+attitude|subject)” means that you tell the model to expect differing baseline-levels of frequency (the intercept, represented by 1) as well as differing responses to the main factor in question, which is “attitude” in this case.

You get p-values for the fixed effects but not for the random slopes.

Multilevel models without random slopes (but with random intercepts) have a too high false positive rate. For this reason you should include all random slopes that are warranted by the data.