Introduction to statistics: Linear mixed models

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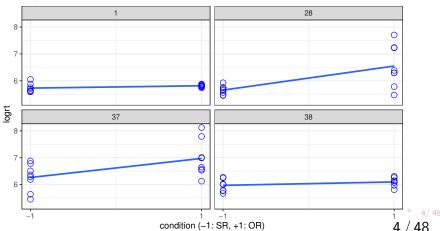
February 10, 2020

As a running example, we will consider subject and objec (SR/OR) relative clause data from English (Grodner and Gibson, Expt 1). First we load the data (not shown).

```
gge1crit<-read.table("data/grodnergibson05data.txt",
                      header=TRUE)
gge1crit$so<-ifelse(gge1crit$condition=="objgap",1,-1)
dat<- gge1crit
dat$logrt<-log(dat$rawRT)
bysubj <- aggregate (logrt "subject + condition,
                   mean, data=dat)
```

The simple linear model (incorrect for these data):

We can visualize the different responses of subjects (four subjects shown):



Given these differences between subjects, you could fit a separate linear model for each subject, collect together the intercepts and slopes for each subject, and then check if the intercepts and slopes are significantly different from zero.

We will fit the model using log reading times because we want to make sure we satisfy model assumptions (e.g., normality of residuals).

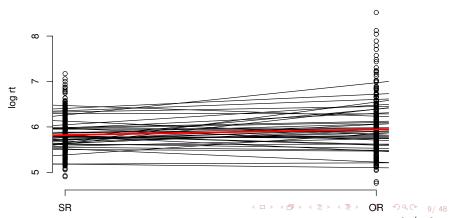
There is a function in the package lme4 that computes separate linear models for each subject: lmList.

```
library(lme4)
## Loading required package: Matrix
lmlist.fm1<-lmList(logrt~so|subject,dat)</pre>
```

Intercept and slope estimates for three subjects:

```
lmlist.fm1$`1`$coefficients
## (Intercept)
                       SO
## 5.769617 0.043515
lmlist.fm1$\^28\^$coefficients
## (Intercept)
                       SO
##
      6.10021 0.44814
lmlist.fm1$\^37\$coefficients
## (Intercept)
                       SO
      6.61699 0.35537
```

One can plot the individual lines for each subject, as well as the linear model m0's line (this shows how each subject deviates in intercept and slope from the model m0's intercept and slopes).



condition

To find out if there is an effect of RC type, you can simply check whether the slopes of the individual subjects' fitted lines taken together are significantly different from zero.

```
t.test(coef(lmlist.fm1)[2])
##
    One Sample t-test
##
##
## data: coef(lmlist.fm1)[2]
## t = 2.81, df = 41, p-value = 0.0076
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 0.017449 0.106585
## sample estimates:
## mean of x
## 0.062017
                                 4□ ト 4団 ト 4 圭 ト 4 圭 ・ 夕 Q @
```

The above test is exactly the same as the paired t-test and the varying intercepts linear mixed model **on aggregated data**:

```
t.test(logrt~condition, bysubj, paired=TRUE) $statistic
##
## 2.8102
## also compare with linear mixed model:
summary(lmer(logrt~condition+(1|subject),
             bysubj))$coefficients[2,]
##
    Estimate Std. Error t value
##
    -0.124033 0.044137 -2.810207
```

- ► The above ImList model we fit is called repeated measures regression. We now look at how to model unaggregated data using the linear mixed model.
- ► This model is now only of historical interest, and useful only for understanding the linear mixed model, which is the modern standard approach.

- ► The linear mixed model does something related to the above by-subject fits, but with some crucial twists, as we see below.
- ▶ In the model shown in the next slide, the statement (1|subject) adjusts the grand mean estimates of the intercept by a term (a number) for each subject.

Notice that we did not aggregate the data here.

```
m0.lmer<-lmer(logrt~so+(1|subject),dat)

Abbreviated output:

Random effects:
Groups Name Variance Std.Dev.
subject (Intercept) 0.09983 0.3160
Residual 0.14618 0.3823

Number of obs: 672, groups: subject, 42

Fixed effects:
```

```
Estimate Std. Error t value
(Intercept) 5.88306 0.05094 115.497
so 0.06202 0.01475 4.205
```

One thing to notice is that the coefficients (intercept and slope) of the fixed effects of the above model are identical to those in the linear model m0 above.

The varying intercepts for each subject can be viewed by typing:

```
ranef(m0.lmer)$subject[,1][1:10]
## [1] -0.1039283  0.0771948 -0.2306209  0.2341978  0.0088
## [7] -0.2055713 -0.1553708  0.0759436 -0.3643671
```

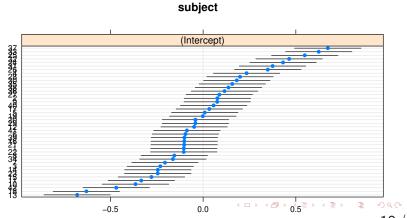
Visualizing random effects

Here is another way to summarize the adjustments to the grand mean intercept by subject. The error bars represent 95% confidence intervals.

```
library(lattice)
print(dotplot(ranef(m0.lmer,condVar=TRUE)))
```

Visualizing random effects

\$subject



The model m0.lmer above prints out the following type of linear model. i indexes subject, and j indexes items.

Once we know the subject id and the item id, we know which subject saw which condition:

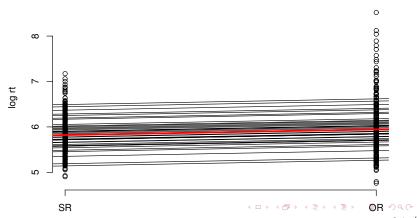
$$y_{ij} = \beta_0 + u_{0i} + \beta_1 \times so_{ij} + \epsilon_{ij} \tag{1}$$

The **only** new thing here is the by-subject adjustment to the intercept.

Note that these by-subject adjustments to the intercept u_{0i} are assumed by lmer to come from a normal distribution centered around 0:

$$u_{0i} \sim Normal(0, \sigma_{u0})$$

- ▶ The ordinary linear model m0 has one intercept β_0 for all subjects, whereas the linear mixed model with varying intercepts m0.lmer has a different intercept $(\beta_0 + u_{0i})$ for each subject i.
- We can visualize the adjustments for each subject to the intercepts as shown below.



condition

Formal statement of varying intercepts linear mixed model

i indexes subjects, j items.

$$y_{ij} = \beta_0 + u_{0i} + (\beta_1 + u_{1i}) \times so_{ij} + \epsilon_{ij}$$
 (2)

Variance components:

- $\blacktriangleright u_0 \sim Normal(0, \sigma_{u0})$
- $ightharpoonup \epsilon \sim Normal(0, \sigma)$

Note that, unlike the figure associated with the lmlist.fm1 model above, which also involves fitting separate models for each subject, the model m0.lmer assumes **different intercepts** for each subject **but the same slope**.

We can have Imer fit different intercepts AND slopes for each subject.

Varying intercepts and slopes by subject

We assume now that each subject's slope is also adjusted:

$$y_{ij} = \beta_0 + u_{0i} + (\beta_1 + u_{1i}) \times so_{ij} + \epsilon_{ij}$$
 (3)

That is, we additionally assume that $u_{1i} \sim Normal(0, \sigma_{u1})$.

```
m1.lmer<-lmer(logrt~so+(1+so||subject),dat)</pre>
```

Random effects:

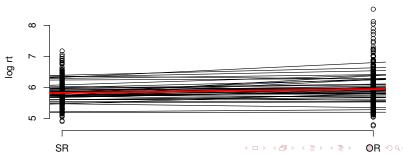
Groups Name Variance Std.Dev. subject (Intercept) 0.1006 0.317 subject.1 so 0.0121 0.110 Residual 0.1336 0.365

Number of obs: 672, groups: subject, 42

Fixed effects: 24/48

These fits for each subject are visualized below (the red line shows the model with a single intercept and slope, i.e., our old model m0):

varying intercepts and slopes for each subject

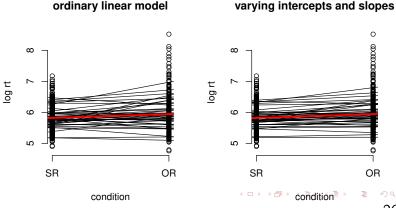


Model type 2: Varying intercepts and slopes model (no correlation)

Linear mixed models

Comparing ImList model with varying intercepts model

Compare this model with the lmlist.fm1 model we fitted earlier:



```
Lecture 6
Linear mixed models
Model type 2: Varying intercepts and slopes model (no correlation)
```

Visualizing random effects

```
print(dotplot(ranef(m1.lmer,condVar=TRUE)))
```

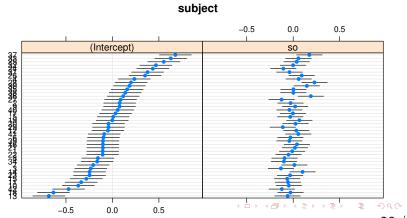
Lecture 6

Linear mixed models

Model type 2: Varying intercepts and slopes model (no correlation)

Visualizing random effects

\$subject



Formal statement of varying intercepts and varying slopes linear mixed model

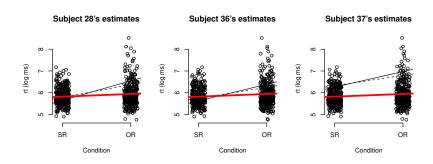
i indexes subjects, j items.

$$y_{ij} = \beta_0 + u_{0i} + \beta_1 \times so_{ij} + \epsilon_{ij} \tag{4}$$

Variance components:

- $\blacktriangleright u_0 \sim Normal(0, \sigma_{u0})$
- $ightharpoonup u_1 \sim Normal(0, \sigma_{u1})$
- $ightharpoonup \epsilon \sim Normal(0, \sigma)$

- ► The estimate of the effect by participant is smaller than when we fit a separate linear model to the subject's data.
- ► This is called shrinkage in linear mixed models: the individual level estimates are shunk towards the mean slope.
- ► The less data we have from a given subject, the more the shrinkage.



The effect of missing data on estimation in LMMs

Let's randomly delete some data from one subject:

```
set.seed(4321)
## choose some data randomly to remove:
rand<-rbinom(1,n=16,prob=0.5)</pre>
```

The effect of missing data on estimation in LMMs

```
dat[which(dat$subject==37),]$rawRT

## [1] 770 536 686 578 457 487 2419 884 3365 233
## [15] 1081 971

dat$deletedRT<-dat$rawRT
dat[which(dat$subject==37),]$deletedRT<-ifelse(rand,NA,dat</pre>
```

The effect of missing data on estimation in LMMs

Now subject 37's estimates are going to be pretty wild:

```
subset(dat,subject==37)$deletedRT

## [1] 770 NA 686 578 NA NA NA NA 3365 233
## [15] NA 971
```

The effect of missing data on estimation in LMMs

```
## original no pooling estimate:
lmList.fm1_old<-lmList(log(rawRT)~so|subject,dat)</pre>
coefs_old<-coef(lmList.fm1_old)</pre>
intercepts_old<-coefs_old[1]
colnames(intercepts_old)<-"intercept"</pre>
slopes_old<-coefs_old[2]
## subject 37's original estimates:
intercepts_old$intercept[37]
## [1] 6.617
slopes_old$so[37]
   [1] 0.35537
```

35/4

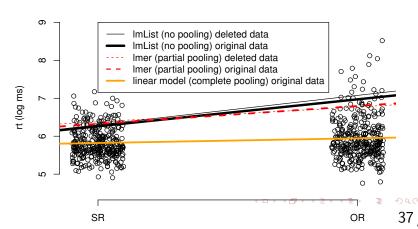
The effect of missing data on estimation in LMMs

```
## on deleted data:
lmList.fm1_deleted<-lmList(log(deletedRT)~so|subject,dat)</pre>
coefs<-coef(lmList.fm1_deleted)</pre>
intercepts<-coefs[1]
colnames(intercepts)<-"intercept"</pre>
slopes<-coefs[2]
## subject 37's new estimates on deleted data:
intercepts$intercept[37]
## [1] 6.6879
slopes$so[37]
   [1] 0.38843
```

Shrinkage in linear mixed models

The effect of missing data on estimation in LMMs

Subject 37's estimates



Shrinkage in linear mixed models

The effect of missing data on estimation in LMMs

- ▶ What we see here is that the estimates from the hierarchical model are barely affected by the missingness, but the estimates from the no-pooling model are heavily affected.
- ► This means that linear mixed models will give you more robust estimates (think Type M error!) compared to no pooling models.
- This is one reason why linear mixed models are such a big deal.

Crossed subjects and items in LMMs

Subjects and items are fully crossed:

igspace Varying intercepts and slopes model, with crossed random effects for subjects and for items

Linear mixed models

Linear mixed model with crossed subject and items random effects.

m2.lmer<-lmer(logrt~so+(1+so||subject)+(1+so||item),dat)</pre>

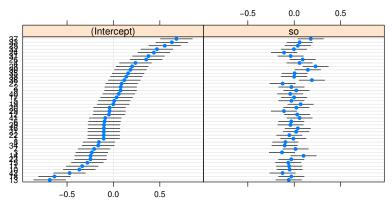
```
Random effects:

Groups Name Variance Std.Dev.
subject (Intercept) 0.10090 0.3177
subject.1 so 0.01224 0.1106
item (Intercept) 0.00127 0.0356
item.1 so 0.00162 0.0402
Residual 0.13063 0.3614
Number of obs: 672, groups: subject, 42; item, 16
```

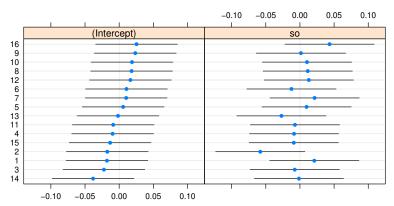
Fixed effects:

```
Estimate Std. Error t value (Intercept) 5.8831 0.0517 113.72 so 0.0620 0.0242 2.56
```

subject



item



Linear mixed model with crossed subject and items random effects.

Linear mixed model with crossed subject and items random effects.

```
Random effects:
```

```
Groups Name Variance Std.Dev. Corr
subject (Intercept) 0.10103 0.3178
so 0.01228 0.1108 0.58
item (Intercept) 0.00172 0.0415
so 0.00196 0.0443 1.00 <= degenerate
Residual 0.12984 0.3603
Number of obs: 672, groups: subject, 42; item, 16
```

Fixed effects:

```
Estimate Std. Error t value (Intercept) 5.8831 0.0520 113.09
```

so 0.0620 0.0247 2.51 3 3 45 45 45 45 45 45 45

Formal statement of varying intercepts and varying slopes linear mixed model with correlation

i indexes subjects, j items.

Lecture 6

$$y_{ij} = \alpha + u_{0i} + w_{0j} + (\beta + u_{1i} + w_{1j}) * so_{ij} + \varepsilon_{ij}$$
(5)

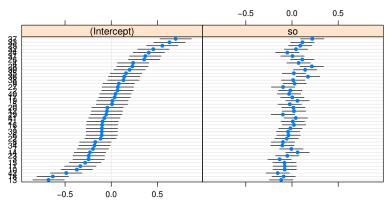
where
$$\varepsilon_{ij} \sim Normal(0, \sigma)$$
 and

$$\Sigma_{u} = \begin{pmatrix} \sigma_{u0}^{2} & \rho_{u}\sigma_{u0}\sigma_{u1} \\ \rho_{u}\sigma_{u0}\sigma_{u1} & \sigma_{u1}^{2} \end{pmatrix} \quad \Sigma_{w} = \begin{pmatrix} \sigma_{w0}^{2} & \rho_{w}\sigma_{w0}\sigma_{w1} \\ \rho_{w}\sigma_{w0}\sigma_{w1} & \sigma_{w1}^{2} \end{pmatrix}$$

$$(6)$$

$$\begin{pmatrix} u_0 \\ u_1 \end{pmatrix} \sim \mathcal{N}\left(\begin{pmatrix} 0 \\ 0 \end{pmatrix}, \Sigma_u \right), \quad \begin{pmatrix} w_0 \\ w_1 \end{pmatrix} \sim \mathcal{N}\left(\begin{pmatrix} 0 \\ 0 \end{pmatrix}, \Sigma_w \right)$$
 (7)

subject



These are degenerate estimates

item

