STA 303/1002-Methods of Data Analysis II Sections L0101& L0201, Winter 2018

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Linear Mixed Models

- ► Learning Objectives
 - Define fixed and random effects
 - Write out the models used and the assumptions for inference
 - Develop a statistical toolbox for analyzing linear mixed models
 - ▶ Interpret the respective R outputs
- ► Reference: SJS, Chapter 10

Example I: Orthodontics Growth Data

- Study conducted at Department of Orthodontics from North Carolina Dental School
- ► Followed growth of 27 children (16 males, 11 females)
- ▶ Measured at ages 8, 10, 12 and 14
- Response: Distance (in mm) from the centre of the pituitary to the pterygomaxillary fissure
- ► Interest: Model distances in terms of age and sex
- What are the fixed effects?
- ▶ What are the random effects?

Example I: Orthodontics Growth Data

```
Grouped Data: distance ~ age | Subject
 distance age Subject Sex
     26.0 8
                M01 Male
1
     25.0 10 MO1 Male
    29.0 12 MO1 Male
    31.0 14 MO1 Male
     21.5 8 MO2 Male
5
     22.5 10
                MO2 Male
Grouped Data: distance ~ age | Subject
  distance age Subject
                       Sex
     21.0
           8 F01 Female
65
66
     20.0 10 F01 Female
     21.5 12 F01 Female
67
68 23.0 14 F01 Female
69 21.0 8 F02 Female
70
     21.5 10 F02 Female
```

Example I: Mixed Model

$$\begin{aligned} Distance_{ijk} &= \beta_0 \\ &+ \beta_1 \mathbf{I}_{[sex=male],j} \\ &+ \beta_2 \mathbf{I}_{[age=10],k} + \beta_3 \mathbf{I}_{[age=12],k} + \beta_4 \mathbf{I}_{[age=14],k} \\ &+ \beta_5 \mathbf{I}_{[sex=male],j} * \mathbf{I}_{[age=10],k} \\ &+ \beta_6 \mathbf{I}_{[sex=male],j} * \mathbf{I}_{[age=12],k} \\ &+ \beta_7 \mathbf{I}_{[sex=male],j} * \mathbf{I}_{[age=14],k} \\ &+ u_{ij} \\ &+ \epsilon_{ijk} \end{aligned}$$

where

- Distance_{ijk}: distance at time k on subject i in treatment j
- \triangleright u_{ij} : random effect due to subject i of sex j
- $ightharpoonup \epsilon_{ijk}$: random error

Example II: Carbohydrates in Diabetes

- ▶ Diet study on n=71 persons with Type 2 diabetes
- ► Each person was assigned to 1 of 3 treatment (diet) groups:
 - I) HG: high GI (glycemic index)
 - II) LG: low GI
 - III) HM: high in monosaturated fats
- ► Traced for 6 months: measurements taken at 0, 3 and 6 months

Example II: Data

► The first 10 observations:

				-	
Obs	id	diet	hdl1	hdl2	hdl3
_1	1	LG	0.97	1.14	1.45
2	3	HG	0.85	0.85	0.84
3	5	HM	1.11	1.36	1.25
4	6	HM	0.95	0.99	0.96
5	8	LG	0.78	0.80	0.72
6	9	HG	0.71	0.76	0.78
7	10	LG	0.58	0.69	0.72
8	11	HM	1.24	1.24	1.31
9	12	HM	0.93	1.18	0.98
10	13	HG	1.65	1.23	1.24

- ▶ Variables of interest: ID#, Diet, Time
- ▶ Outcome of interest: HDL- level of "good" cholesterol
- ► Aim: Is there a diet* time interaction? Do differences among diets change over time?

Example II: Mixed Model

$$Y_{ijk} = \beta_0$$

$$+ \beta_1 \mathbf{I}_{[diet=HG],j} + \beta_2 \mathbf{I}_{[diet=HM],j}$$

$$+ \beta_3 \mathbf{I}_{[time=1],k} + \beta_4 \mathbf{I}_{[time=2],k}$$

$$+ \beta_6 \mathbf{I}_{[diet=HG],j} * \mathbf{I}_{[time=1],k} + \beta_7 \mathbf{I}_{[diet=HG],j} * \mathbf{I}_{[time=2],k}$$

$$+ \beta_8 \mathbf{I}_{[diet=HM],j} * \mathbf{I}_{[time=1],k} + \beta_9 \mathbf{I}_{[diet=HM],j} * \mathbf{I}_{[time=2],k}$$

$$+ u_{ij}$$

$$+ \epsilon_{ijk}$$

- $ightharpoonup Y_{ijk}$: response at time k on subject i in treatment j
- \triangleright u_{ij} : random effect due to subject i under diet j
- $ightharpoonup \epsilon_{ijk}$: random error

Variance-Covariance Structure

Assume:

Subjects are independent and

- within u_{ij} are i.i.d. $N(0, \sigma_{ii}^2)$
 - $ightharpoonup \epsilon_{ijk}$ are i.i.d. $N(0, \sigma_e^2)$
 - $ightharpoonup u_{ij}$ and ϵ_{ijk} are independent

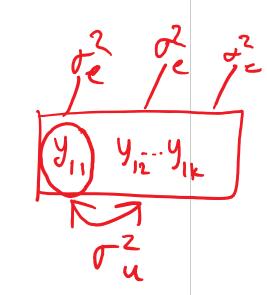
Therefore:

▶ Different subjects, $i \neq I$:

$$Cov(Y_{ijk}, Y_{lmn}) = 0$$

n subjects, kobspersubj. Total: nk observations

Variance-Covariance Structure



- Same subject at the <u>same time</u> (i = l, j = m, k = n): $Var(Y_{ijk}) = Cov(Y_{ijk}, Y_{lmn}) =$
- Same subject at different time $(k \neq n)$: $Cov(Y_{ijk}, Y_{ijn}) = Cov(u_{ij}, u_{ij}) \longrightarrow Var(u_{ij}).$ $+ Cov(u_{ij}, e_{ijn}) + Cov(e_{ijk}, u_{ij})$ $+ Cov(e_{ijk}, e_{ijn}) \longrightarrow Var(u_{ij}).$ $+ Cov(e_{ijk}, e_{ijn}) \longrightarrow Var(u_{ij}).$

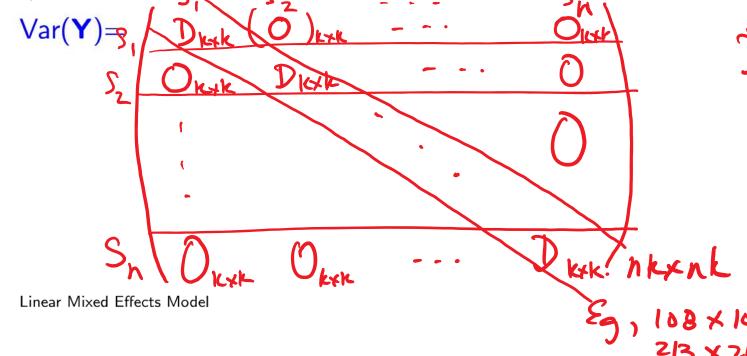
Compound Symmetry Variance-Covariance Structure

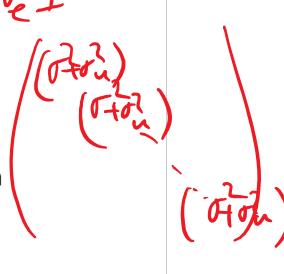
Intraclass Correlation Coefficient:

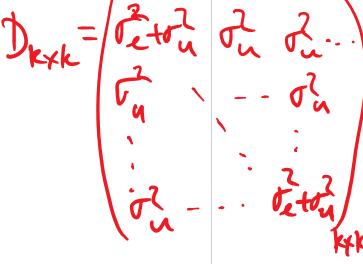
$$\rho_{IC} = \frac{Cov(Y_{ijk}, Y_{ijn})}{\sqrt{Var(Y_{ijk})Var(Y_{ijn})}} = \frac{\sigma_u^2}{\sigma_u^2 + \sigma_e^2}$$

the correlation is the same for every pair of observations on the same subject

the variance is the same for all observations







Mixed model for repeated measures data

MIXED Model:

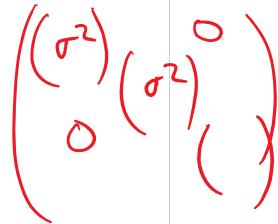
$$\mathbf{Y} = \underbrace{f(\mathbf{X}, \boldsymbol{\beta})}_{\text{fixed effects}} + \underbrace{u}_{\text{random effects}} + \underbrace{\epsilon}_{\text{random noise}}$$

Part	Properties	Eg.I	Eg.II
Response, Y	continuous, $Y \sim Normal$	HDL	Distance
Fixed effects, X	continuous or categorical	diet, time	sex, age
Random effect, u	$\mathbf{u} \sim \mathcal{N}(0, \sigma_u^2)$	subject	subject
Random error, ϵ	$\mathbf{e} \sim N(0, \sigma_e^2), \ u \perp e$		

 Random effect induces a random intercept (different intercept for different subjects)

Restricted Maximum Likelihood Estimation

- ► Restricted Maximum Likelihood (or Residual ML): gives unbiased estimators of variance-covariance parameters
- **b** based on the notion of separating the likelihood used for estimating Σ from that used for estimating β
- Steps:
 - (1) Estimate variance-covariance parameters using maximum likelihood estimation
 - (2) Use the estimates from (1) and estimate β 's of fixed effects using generalized least squares (GLS) where:
 - ▶ GLS is least squares optimization with adjustment since $Var(\mathbf{Y}) = V \neq \sigma^2 \mathbf{I}$
 - We get $\widehat{\beta} = (X^\top V^{-1} X)^{-1} X^\top V^{-1} Y$ and $Var(\widehat{\beta}) = (X^\top V^{-1} X)^{-1}$
 - (3) Repeat steps (1) and (2) until convergence



Within-subject Covariance structures

▶ Between-subject: Recall that we assumed that subjects are independent, so between-subject covariance is 0, i.e.,

$$Cov(Y_{ijk}, Y_{lmn}) = 0 \text{ if } i \neq I$$

•	Within-su	ubject: 3 + 17			
	Туре	Interpretation	# cov. parameters	3	
	CS	-same variances and common	2		Eg, k=4
		covariance —			
	UN	-different variances and	(t*(t+1))/2	k (k+1)	=4(s)
		difference covariances		2	2
	AR(1)	-same variances, covariance	2		= 10
		decrease exponentially with			, ,
		distance			

Within-subject Covariance structures

Compound Symmetry [2]: same variance and common covariances

$$D_{CS} = \begin{bmatrix} \sigma_u^2 + \sigma_\epsilon^2 & \sigma_u^2 & \sigma_u^2 \\ \sigma_u^2 & \sigma_u^2 + \sigma_\epsilon^2 & \sigma_u^2 \\ \sigma_u^2 & \sigma_u^2 & \sigma_u^2 + \sigma_\epsilon^2 \end{bmatrix}$$

▶ Unstructured [t(t+1)/2]: different variances and different covariances

$$D_{UN} = \begin{bmatrix} \sigma_1^2 & \sigma_{12}^2 & \sigma_{13}^2 \\ \sigma_{12}^2 & \sigma_2^2 & \sigma_{23}^2 \\ \sigma_{13}^2 & \sigma_{23}^2 & \sigma_3^2 \end{bmatrix}$$

 Auto-Regressive, lag1 [2]: same variances, covariances decrease exponentially

$$D_{AR(1)} = \sigma^2 \begin{bmatrix} 1 & \rho & \rho^2 \\ \rho & 1 & \rho \\ \rho^2 & \rho & 1 \end{bmatrix}$$

Comparing models

▶ Using likelihood-based criteria (due to MLE): compare models with same *Y* and *X*'s but different covariance structures

```
    AIC= -2 Res log L + 2(# of covariance parameters)
    BIC= -2 Res log L+(# of covariance parameters)log(n), where n=# of subjects
    G² = -2 Res log(LR)
```

▶ Using \underline{t} and \underline{F} tests (due to GLS): check relevance of fixed effects (test β 's)