

# STAT 770 Nov. 18 Lecture 23

## GLMs with Random (Intercept) Effects

Reading and Topics for this and next lecture: Chapter 13 Sections 13.1 – 13.3 and Handout 1(ii) from course web-page.

- (1) Random effects model for clustered GLM data
- (2) Matched Pairs as an Example of Clusters
- (3) Other Examples of GLMMs – Binary outcome models, Item-response models
- (4) GLMM likelihood – cluster means, within-cluster correlations, and predictors
- (5) Strategies for ML and approximate ML calculation

**next class**

## What is a GLMM ?

General form of GLMM with cluster-level random intercepts:

$$Y_{ij} \sim f(y | \theta_{ij}, \underline{X}_{ij}, \epsilon_i) = h(y) \exp(\theta'_{ij} y - c(\theta_{ij})) , \quad \epsilon_i \stackrel{iid}{\sim} F(\cdot, \sigma)$$

$$\nabla_{\theta} c'(\theta_{ij}) = \mu_{ij} = g^{-1}(\underline{X}'_{ij} \beta + \epsilon_i)$$

$i = 1, \dots, m$  indexes **cluster**, with cluster random-effect  $\epsilon_i$

$j = 1, \dots, n_i$  indexes observation within cluster  $i$ ,

conditionally *iid* given  $\epsilon_i$

- $\beta \in \mathbb{R}$ ,  $\sigma > 0$  parameters are shared across cluster
- $\epsilon_i$  a random cluster-specific **offset**

## Examples

**Idea:** cluster combines observations with common experimental setting

**Examples:** families, schools, clinical centers within which units share some common unmodeled feature, modeled as independent variate differing across cluster

If  $m$  were small and offsets  $\epsilon_i$  of direct interest, they would be viewed as non-random fixed-effect parameters.

Dimension of GLMM parameter  $(\beta, \sigma)$  is  $p + 1$ .

Model dimension with fixed effects =  $p + m$ .

## Additional Keywords for Models with Random Effects

**Empirical Bayes Models:** these GLMMs can be interpreted as a way to fit/predict cluster predictive-scores  $X'_{ij}\beta + \epsilon_i$  for individual  $(i, j)$ : **empirical** because of the identical model relationships across cluster, **Bayes** because of the independence of observations (identical within cluster) conditional on 'prior' effects  $\epsilon_i$ .

**Meta-Analysis:** think of the clusters (eg clinical centers) as separate (small but not too tiny) studies; one might report estimates  $\hat{\beta}_i, \widehat{\text{Var}}(\hat{\beta}_i)$  from those studies and analyze those as estimates of common  $\beta$ , with  $\epsilon_i$  modeling differences between studies. GLMM is what one would fit to the unified dataset if it were available.

## Special Logistic Mixed Models

Logit-Normal Mixed-model:

$$Y_{ij} \sim \text{Binom}(r_{ij}, \text{plogis}(\underline{X}'_{ij}\beta + \epsilon_i)) \text{ given } \epsilon_i, \quad \epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

Binary Matched Pairs:  $n_i \equiv 2$ ,  $\underline{X}_{ij} = \underline{x}_i$

similarly, Litter-Studies or Family-studies

Rasch Item-Response Model:  $n_i \equiv \nu \geq 2$ ,  $\underline{X}_{ij} = \underline{X}_i \in \mathbb{R}^\nu$   
dummy-column for factor with levels  $j = 1, \dots, \nu$ , so  $\underline{X}'_{ij}\beta \equiv \beta_j$ ,

$$\text{logit } P(Y_{ij} = 1 | \epsilon_i) = \beta_j + \epsilon_i$$

$j$  indexes study subject with skill-level  $\beta_j$ ,  $i$  item with difficulty  $\epsilon_i$

## Within-cluster Correlation

General Formula:  $\text{var}(W) = E(\text{var}(W|Z)) + \text{var}(E(W|Z))$

Consequence: after using the formula for  $W = W_1 + W_2$  and  $W = W_1 - W_2$  and subtracting 2nd from 1st, dividing by 4:

$$\text{cov}(W_1, W_2) = E(\text{cov}(W_1, W_2 | Z)) + \text{cov}(E(W_1 | Z) \cdot E(W_2 | Z))$$

Fix cluster index  $i$  and  $\underline{X}_{ij}$ 's (by conditioning on them if random)

$$\begin{aligned}\text{cov}(Y_{i,j_1}, Y_{i,j_2}) &= E(\text{cov}(Y_{i,j_1}, Y_{i,j_2} | \epsilon_i)) + \text{cov}(E(Y_{i,j_1} | \epsilon_i) \cdot E(Y_{i,j_2} | \epsilon_i)) \\ &= \text{cov}(E(Y_{i,j_1} | \epsilon_i) \cdot E(Y_{i,j_2} | \epsilon_i)) > 0\end{aligned}$$

**Next slide presents an argument why this inequality is true.**

## Proof of Positive Within-Cluster Correlation

This argument applies to scalar random-intercept GLMM with  $g, c'$  strictly increasing.

- $Y \sim f(y, \theta) = h(y) \exp(\theta y - c(\theta))$  is **stochastically increasing** as function of  $\theta$ , i.e. , for  $\theta_1 < \theta_2$  and all  $y$ ,  $F(y, \theta_1) \geq F(y, \theta_2)$
- $Y$  is stochastically increasing in  $\theta$  if and only if  $E_\theta(h(Y)) \nearrow$  in  $\theta$  for any monotone increasing function  $h$

**Pf. for GLMM**  $\frac{\partial}{\partial \theta} \int h(y) f(y, \theta) dy = \int (y - c'(\theta)) h(y) f(y, \theta) dy > 0$   
because  $h(y) > h(c'(\theta)) > h(z)$  whenever  $y > c'(\theta) > z$ .

- In GLMM,  $g_j(\epsilon_i) \equiv E(Y_{i,j} | \epsilon_i) \nearrow$  in  $\epsilon_i$  (varies with  $\underline{X}_{ij}$ )
- $\text{cov} = E[g_{j_1}(\epsilon_i) g_{j_2}(\epsilon_i)] - [E(g_{j_1}(\epsilon_i)) \cdot E(g_{j_2}(\epsilon_i))]$   
 $= \int (g_{j_1}(t) - E(g_{j_1}(\epsilon_i))) g_{j_2}(t) f_\epsilon(t) dt > 0$

## Unit-level Prediction (Logit-Normal Model)

Fix  $i, j$ ,  $\eta = \underline{X}'_{i,j} \beta$ :

$$P(Y_{ij} = 1) = E(\text{plogis}(\eta + \epsilon_i)) = \int \frac{e^{\eta + \sigma z}}{1 + e^{\eta + \sigma z}} \phi(z) dz$$

**Remark.** If  $\underline{X}_{ij} = \underline{X}_i$  depends only on  $i$ , not  $j$ ,

- the displayed quantity is the inverse of a modified link-function  $g_\sigma^*$  if  $\sigma$  is fixed
- the last inequality on the previous slide is the Cauchy-Schwarz inequality



## GLMM Likelihood

Consider logit-normal model, and for simplicity restrict to case where all  $Y_{ij}$  are binary, and cluster  $i$  has  $n_i$  unit observations.

Then

$$\begin{aligned} P(Y_{i,j} = y_j, 1 \leq j \leq n_i) &= E\left(P(Y_{i,j} = y_j, 1 \leq j \leq n_i \mid \epsilon_i)\right) \\ &= \int \exp\left(\sum_{j=1}^{n_i} y_j (\underline{X}_{ij}^{tr} \beta + \sigma z)\right) \left[\prod_{j=1}^{n_i} (1 + e^{\underline{X}_{ij}^{tr} \beta + \sigma z})\right]^{-1} \phi(z) dz \end{aligned}$$

We will consider strategies for maximizing in  $\beta, \sigma$  the sum of log's over  $i$ , mostly in the simpler case where all covariates  $\underline{X}_{i,j} = \underline{X}_i$  are at cluster level

## Software Comments

`glmer` in R package `lme4`

`PROC NLINMIX` or `GLIMMIX` in SAS

EM, MCMC and **Adaptive Gaussian Quadrature** ideas for  
logLik maximization