# STAT3016/4116/7016 Introduction to Bayesian Data Analysis

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Linear and generalized linear mixed effects models

**Example:** Recall the school data from Chapter 8. Let's now examine the relationship between math scores and another variable, socioeconomic status (SES). Perhaps the relationship between mathscore and SES varies from school to school.

- Ordinary least squares model?
- ► Hierarchical Model? hierarchical model: y\_[ij] ~ iid normal (theta\_j, sigma^2) within group model theta\_j ~ iid normal (mu, tau^2) between group model
- Bayesian Hierarchical Model?

now for hierarchical linear regression model, use betaTx to replace y {ii} and beta to replace theta i

# Least squares analysis of mathscore data when the sample size is small,

the variability is large math score 10 15 20 25 30 10 15 20 25 30 SES

**Priors** 

$$\theta \sim MVN(\mu_0, \Lambda_0)$$

$$oldsymbol{\Sigma} \sim \mathit{inverse} - \mathit{Wishart}(\eta_0, S_0^{-1})$$

$$\sigma^2 \sim inverse - gamma(\nu_0/2, \nu_0 \sigma_0^2/2)$$

#### Sampling Model

$$\mathbf{Y}_{j} \sim MVN(\mathbf{X}_{j}\beta_{j}, \sigma^{2}\mathbf{I})$$

$$\beta_i \stackrel{\text{iid}}{\sim} MVN(\theta, \Sigma)$$

(j=1,...,m; m groups)

Full conditional distributions of  $\beta_j$ 

$$Var[\beta_j|\mathbf{y}_j, \mathbf{X}_j, \sigma^2, \boldsymbol{\theta}, \Sigma] = (\Sigma^{-1} + \mathbf{X}_j^T \mathbf{X}_j/\sigma^2)^{-1}$$

$$E[\beta_j|\mathbf{y}_j,\mathbf{X}_j,\sigma^2,\theta,\Sigma] = (\Sigma^{-1} + \mathbf{X}_j^T\mathbf{X}_j/\sigma^2)^{-1}(\Sigma^{-1}\theta + \mathbf{X}_j^T\mathbf{y}_j/\sigma^2)$$

Full conditional distributions of  $\theta$ 

$$Var[\theta|\beta_1,...,\beta_m] = (\Lambda_0^{-1} + m\Sigma^{-1})^{-1}$$

$$E[\theta|\beta_1,...,\beta_m] = (\Lambda_0^{-1} + m\Sigma^{-1})^{-1}(\Lambda_0^{-1}\mu_0 + m\Sigma^{-1}\bar{\beta})$$

$$(\bar{\boldsymbol{\beta}} = \frac{1}{m} \sum_{j} \boldsymbol{\beta}_{j})$$

#### Full conditional distributions of $\Sigma$

$$\Sigma | m{ heta}, m{eta}_1, ..., m{eta}_m \sim \mathit{Inverse} - \mathit{Wishart}(\eta_0 + m, [\mathbf{S}_0 + \mathbf{S}_{ heta}]^{-1})$$
 where  $\mathbf{S}_{ heta} = \sum_{j=1}^m (m{eta}_j - m{ heta}) (m{eta}_j - m{ heta})^T$ 

#### Full conditional distributions of $\sigma^2$

$$\sigma^2 \sim \textit{Inverse} - \textit{Gamma}([
u_0 + \sum_j \textit{n}_j]/2, [
u_0 \sigma_0^2 + \textit{SSR}]/2)$$

where 
$$SSR = \sum_{j=1}^{m} \sum_{i=1}^{n_j} (y_{i,j} - \beta_j^T \mathbf{x}_{i,j})^2$$
.

# A hierarchical regression model - example

#### Mathscore data example

$$\mu_{0} = \frac{1}{m} \sum_{j=1}^{m} \hat{\beta}_{j,OLS};$$

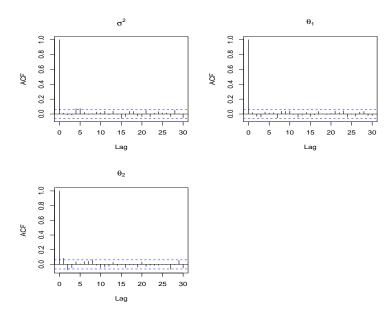
$$\Lambda_{0} = \mathbf{S}_{0} = Cov(\hat{\beta}_{j,OLS});$$

$$\eta_{0} = p + 2 = 4;$$

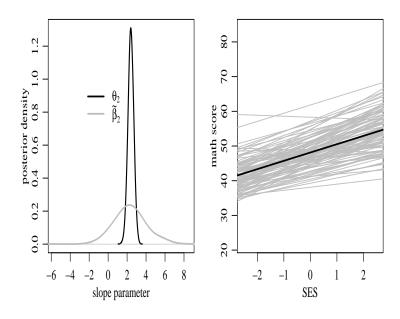
$$\sigma_{0}^{2} = \frac{1}{m} \sum_{j=1}^{m} \hat{\sigma}_{j,OLS}^{2}; \ \nu_{0} = 1$$

$$S=10000, \text{ thin}=10$$

# A hierarchical regression model - example



# A hierarchical regression model - example



#### Generalised Linear Mixed effects models

$$\beta_1,...,\beta_m \stackrel{\text{iid}}{\sim} \text{MVN}(\boldsymbol{\theta},\boldsymbol{\Sigma})$$

$$p(\mathbf{y}_j|X_j\beta_j,\gamma) = \prod_{i=1}^{n_j} p(y_{i,j}|\beta_j^T\mathbf{x}_{i,j},\gamma)$$

 $\gamma$ : scale parameter; mean of y is a function of  $\boldsymbol{\beta}_j^T \mathbf{x}_{i,j}$ 

#### Generalised Linear Mixed effects models

#### A Metropolis-Gibbs algorithm

- ▶ Gibbs steps for  $\theta$ ,  $\Sigma$
- Metropolis step for  $\beta_j$ 
  - 1. Sample  $\beta_j^* \sim \text{MVN}(\beta_j^{(s)}, V_j^{(s)})$
  - 2. Compute

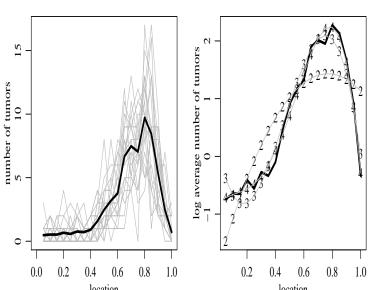
$$r = \frac{p(\mathbf{y}_j | X_j, \beta_j^*) p(\beta_j^* | \boldsymbol{\theta}^{(s)}, \Sigma^{(s)})}{p(\mathbf{y}_j | X_j, \beta_j^{(s)}) p(\beta_j^{(s)} | \boldsymbol{\theta}^{(s)}, \Sigma^{(s)})}$$

3. Sample  $u \sim \text{uniform}(0,1)$ . Set  $\beta_j^{(s+1)}$  to  $\beta_j^*$  if u < r and to  $\beta_j^{(s)}$  if u > r.

 $V_j^{(s)}$  is the proposal variance; perhaps set equal to a scaled version of  $\Sigma^{(s)}$ 

#### Analysis of tumour location data

$$(m = 21; n_j = 20)$$



#### Analysis of tumour location data

Model:  $Y_{x,j}$ : tumor count of mouse j at location x, where

 $Y_{x,j} \sim \operatorname{Pois}(\exp(f_j(x)))$ , and

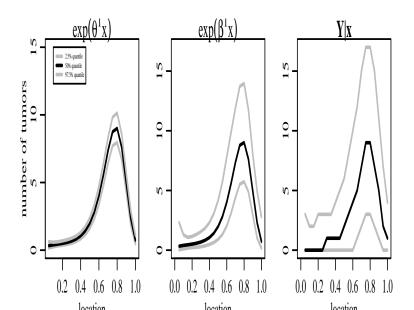
$$f_j(x) = \beta_{0,j} + \beta_{1,j}x + \beta_{2,j}x^2 + \dots + \beta_{p-1,j}x^{p-1} \ (x \in [0,1])$$

A fourth-degree polynomial was fit for  $f_j$  based on visual inspection of graphs of the data.

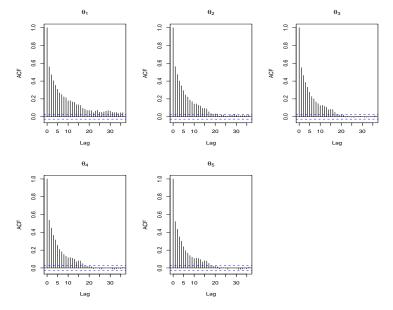
#### Unit information priors

- Prior guess for  $\beta_j$ 's: regress  $\log(y_{1,j}+1/n)...\log(y_{n,j}+1/n)$  on  $(\mathbf{x}_1,...,\mathbf{x}_{20})$  (where  $\mathbf{x}_i=(1,x_i,x_i^2,x_i^3,x_i^4)$  for  $x\in(0.05,0.10,....,0.95))\to \text{obtain estimates }\tilde{\boldsymbol{\beta}}_j$ 's.
- Set  $\mu = \frac{1}{m} \sum_{j=1} \tilde{\beta}_j$ , and a prior covariance matrix equal to the sample covariance matrix of the  $\tilde{\beta}_j$ 's. Set  $S_0$  equal to this sample covariance matrix and  $\eta_0 = p + 2 = 7$  (diffuse prior on variance of  $\beta$ 's).

#### Analysis of tumour location data



#### Analysis of tumour location data - some MCMC diagnostics



### Parameter expansion

Suppose a group level variance parameter happens to be estimated near zero. Then in the updating step for the group level coefficients,  $\beta_1, ..., \beta_J$ , these parameters will be pooled towards their common mean.

But the updating step for the group level variance parameter is based on the coefficients  $\beta_1,...,\beta_J$ , and so will be estimated close to zero again, and the algorithm can get stuck.

<u>Solution</u>: Rescale the  $\beta_j$ 's by multiplying the entire vector of group level coefficients  $\beta$  by a constant.

### Parameter expansion - tumor location example

The parameter expanded version of the tumor location model is:

 $Y_{x,j}$ : tumor count of mouse j at location x, where  $Y_{x,j} \sim \operatorname{Pois}(\exp(f_j(x)))$ , and  $f_j(x) = \alpha \omega_{0,j} + \alpha \omega_{1,j} x + \alpha \omega_{2,j} x^2 + \alpha \omega_{3,j} x^3 + \alpha \omega_{4,j} x^4 \ (x \in [0,1])$ 

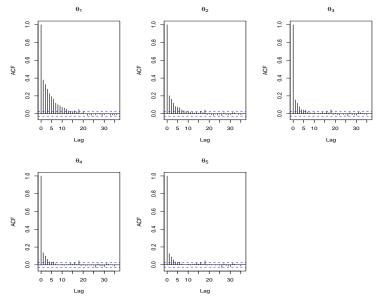
Here,  $\alpha$  is a redundant multiplicative working parameter. The regression parameters of interest are now  $\beta_j = \alpha \omega_j$ , with group level standard deviation of  $\sigma_\beta = |\alpha|\sigma_\omega$ .

# Parameter expansion

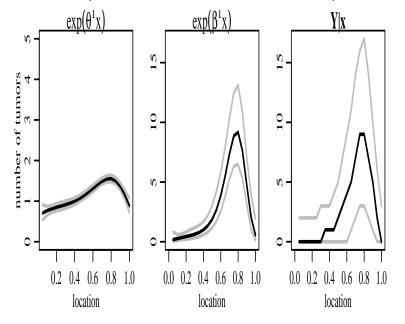
#### Why does it work?

- We transform from  $\beta$  to  $\omega$  via multiplaction by  $\alpha$  that rescales and potentially decorrelates the components of  $\beta$  (equivalent to rotating the parameter space)
- See Gelman et al. (2004) for theoretical derivations to show the increase in speed of convergence under parameter expansion.
- ▶ The likelihood function  $p(y_j|\theta,\sigma_\beta)$  obtained by integrating out the  $\omega_j$ 's in the parameter expanded model is the same as that obtained by integrating out the  $\beta_j$ 's in the original model. So we can fit either model to obtain a Monte Carlo sample from the same posterior distribution  $p(y_j|\theta,\sigma_\beta)$ .
- ▶ The added parameter  $\alpha$  is not necessarily of any statistical interest.

# Parameter expansion - Tumor location example - MCMC diagnostics



# Parameter expansion - Tumor location example - results



# Exercise - a nonlinear hierarchical growth curve model

The following table displays data on the growth of five orange trees over time. The response  $y_{ij}$  is the trunk circumference recorded at time  $x_j$ , j = 1, ..., 7 and i = 1, ..., 5.

Reponse for Tree Number					
X	1	2	3	4	5
118	30	33	30	32	30
484	58	69	51	62	49
664	87	111	75	112	81
1004	115	156	108	167	125
1231	120	172	115	179	142
1372	142	203	138	209	174
1582	145	203	140	214	177

# Exercise - a nonlinear hierarchical growth curve model

One assumes  $y_{ij}$  is normally distributed with mean  $\eta_{ij}$  and variance  $\sigma^2$ , where the means satisfy the non-linear growth model

$$\eta_{ij} = \frac{\phi_{1i}}{1 + \phi_{i2} \exp(\phi_{i3} x_j)}$$

Suppose one reexpresses the parameters as the real-valued parameters:

$$\theta_{i1} = \log \phi_{i1}; \ \theta_{i2} = \log(\phi_{i2} + 1); \theta_{i3} = \log(-\phi_{i3}), \ i = 1, ..., 5$$

Let  $\theta_i$  represent the vector of growth parameters for the  $\mathbf{i}^{th}$  tree.

# Exercise - a nonlinear hierarchical growth curve model

To reflect a prior belief in similarity in the growth patterns of the five trees, one assumes that  $\{\theta_i, i=1,...,5\}$  are a random sample from a multivariate normal distribution with mean vector  $\mu$  and variance covariance matrix  $\Omega$ .

For the hyperprior distribution, assume  $\Omega^{-1}$  follows a Wishart distribution with parameters R and 3, where R is a diagonal matrix with diagonal elements 0.1. Assume  $\mu$  is multivariate normal with mean vector  $\mu_0$  (equal to the zero vector) and variance-covariance matrix M. Let  $M^{-1}$  be a diagonal matrix with diagonal elements  $1.0 \times 10^{-6}$ .

Implement a MCMC algorithm to approximate the posterior distribution of all parameters. Obtain 90% posterior interval estimates for all parameters.