
36-463/663: Multilevel and Hierarchical Models

Multilevel Models in lmer and jags

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1

Outline

- Quick review: Bayesian Statistics, MCMC and JAGS
- Example 1: Minnesota Radon – Intercept Only
 - What's new?
 - What is $\hat{R}$?
- Example 2: Mn Radon: Level 1 predictor “floor”, Level 2 predictor “log(uranium)”
- Example 3: CD4 levels in HIV-positive youth

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2

Bayesian Statistics and MCMC

- Our slogan(s)
 - $(\text{posterior}) \propto (\text{likelihood}) \times (\text{prior})$
 - $(\text{posterior}) \propto (\text{level 1}) \times (\text{level 2})$
 - $(\text{posterior}) \propto (\text{level 1}) \times (\text{level 2}) \times (\text{level 3})$
 - etc.
- Often there are no formulas for posterior, so we have to simulate draws from the posterior
- MCMC is a general method for doing posterior draws
 - Calculate complete conditionals
 - Invent methods for sampling from complete conditionals
 - Stitch together samples into a Markov Chain
 - Summarize Markov Chain with histograms, medians, credible intervals, etc.

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3

Bayesian Statistics and JAGS

- JAGS¹ automates MCMC
 - Describe “slogan” to JAGS in R-like language
 - JAGS figures out complete conditionals
 - JAGS invents MCMC sampling methods
 - JAGS generates the random draws for you
- JAGS is just the latest implementation of the BUGS² platform (declarative language and AI system for constructing MCMC). Others:
 - WinBUGS
 - OpenBUGS

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1. Just Another Gibbs Sampler
2. Bayesian inference Using Gibbs Sampling

4

Multilevel Models in JAGS

■ MLM form

$$\begin{aligned}y_i &= \alpha_{0j[i]} + \epsilon_i, \\ \epsilon_i &\sim N(0, \sigma^2) \\ \alpha_{0j} &= \beta_0 + \eta_j, \\ \eta_j &\sim N(0, \tau^2)\end{aligned}$$

■ Hierarchical form

Level 1: $y_i \sim N(\alpha_{0j[i]}, \sigma^2)$

Level 2: $\alpha_{0j} \sim N(\beta_0, \tau^2)$

■ BUGS/rube form

```
model {  
  # LEVEL 1  
  for (i in 1:N) {  
    log.radon[i] ~  
      dnorm(a0[county[i]], prec.y)  
  }  
  # LEVEL 2  
  for (j in 1:MAX.COUNTY=85) {  
    a0[j] ~ dnorm(b0, prec.cty)  
  }  
  
  # PRIORS/LEVEL 3:  
  b0 ~ dnorm(0, PREC.b0=1e-6)  
  prec.y ~ dgamma(ALPHA=0.001, BETA=0.001)  
  prec.cty ~ dgamma(ALPHA=0.001, BETA=0.001)  
  
  # CONVERT PRECISION TO VARIANCES  
  var.y <- 1/prec.y  
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7

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Look at the
Distributions
section (pp 56ff
in manual14.pdf)

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8

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Have to add priors
to all free parameters

Example 1: Minnesota Radon – Intercept Only

- MLM:
$$y_i = \alpha_{0j[i]} + \epsilon_i,$$
$$\epsilon_i \sim N(0, \sigma^2)$$
$$\alpha_{0j} = \beta_0 + \eta_j,$$
$$\eta_j \sim N(0, \tau^2)$$
- Hierarchical:
 - Level 1:** $y_i \sim N(\alpha_{0j[i]}, \sigma^2)$
 - Level 2:** $\alpha_{0j} \sim N(\beta_0, \tau^2)$
- Demonstration in R and rube()/JAGS...
- (comparison with lmer also)

What's new?

- `summary(rube.object)`: point estimates and CI's for "some" parameters. Others available in
 - `rube.object$mean`
 - `rube.object$sd`
 - `rube.object$median`
 - `rube.object$sims.list`
 - etc
- `p3(rube.object)`: interactive graphical summaries

What's new?

- `rube()`/WinBUGS/JAGS automatically
 - Runs 3 separate MCMC chains
 - Runs each MCMC chain for 2000 steps, and throws away the first 1000 steps as “burn-in”
 - Thins each chain by 1/3 to reduce autocorrelation
 - *You can change this when you run `rube()`; see pp. 25ff. of the “`rube.pdf`” documentation.*
- `rube()`/WinBUGS/JAGS reports an “Rhat” statistic for each parameter estimated
 - Rhat is a ratio of between-chain to within-chain variation
 - *When the chain is converged, $Rhat < 1.2$. Otherwise, the chain hasn't run long enough yet.*

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13

What is $\hat{R}$?

- Suppose we have M chains:

Chains				Means	Variances
$\theta^{(1;1)},$	$\theta^{(1;2)},$	$\dots,$	$\theta^{(1;N)}$	$\bar{\theta}_1$	W_1
$\vdots$	$\vdots$	$\ddots$	$\vdots$	$\vdots$	$\vdots$
$\theta^{(M;1)},$	$\theta^{(M;2)},$	$\dots,$	$\theta^{(M;N)}$	$\bar{\theta}_M$	W_M
Grand mean				$\bar{\theta}$	

- Define

$$W = \frac{1}{M(N-1)} \sum_{m=1}^M \sum_{n=1}^N (\theta^{(m;n)} - \bar{\theta}_m)^2 = \frac{1}{M} \sum_{m=1}^M W_m$$

= Average within-chain variance

$$B = \frac{M}{M-1} \sum_{m=1}^M (\bar{\theta}_m - \bar{\theta})^2$$

= Between-chain variance, inflated for sample size

$$V = \frac{M-1}{M} W + \frac{1}{M} B$$

= Pooled variance estimate,

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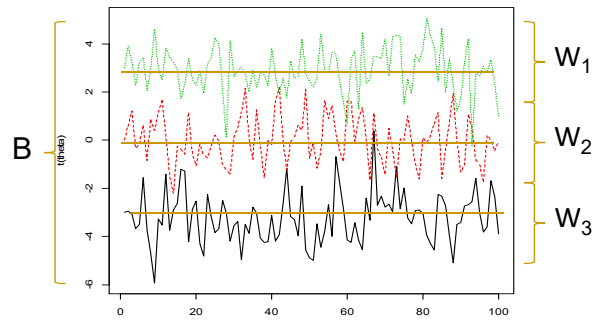
14

What is $\hat{R}$?

■ Separated chains:

$$\begin{aligned} W &= \frac{1}{N} \sum_{n=1}^N W_n = 1.02 \\ B &= \frac{M}{N-1} \sum_{n=1}^N (\bar{\theta}_n - \bar{\theta})^2 = 938.53 \\ V &= \frac{M-1}{M} W + \frac{1}{M} B = 10.40 \end{aligned}$$

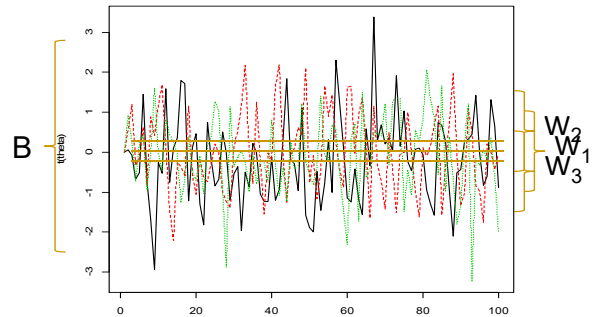
$$\hat{R} = \sqrt{V/W} = 3.19$$



■ Converged chains:

$$\begin{aligned} W &= \frac{1}{N} \sum_{n=1}^N W_n = 1.02 \\ B &= \frac{M}{N-1} \sum_{n=1}^N (\bar{\theta}_n - \bar{\theta})^2 = 1.32 \\ V &= \frac{M-1}{M} W + \frac{1}{M} B = 1.03 \end{aligned}$$

$$\hat{R} = \sqrt{V/W} = 1.00$$



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15

Example 2: Mn Radon: Level 1 predictor “floor”, Level 2 predictor “log(uranium)”

■ MLM:

$$\begin{aligned} y_i &= \alpha_{0j[i]} + \alpha_{1j[i]}(\text{floor})_i + \epsilon_{ij[i]}, \quad \epsilon_{ij} \sim N(0, \sigma^2) \\ \alpha_{0j} &= \beta_{00} + \beta_{01} \log(\text{uranium}_j) + \eta_{0j}, \quad \eta_{0j} \sim N(0, \tau_0^2) \\ \alpha_{1j} &= \beta_{10} + \eta_{1j}, \quad \eta_{1j} \sim N(0, \tau_1^2) \end{aligned}$$

■ Hierarchical:

$$\text{Level 1: } y_i \sim N(\alpha_{0j[i]} + \alpha_{1j[i]}(\text{floor})_i, \sigma^2)$$

$$\text{Level 2: } \alpha_{0j} \sim N(\beta_{00} + \beta_{01} \log(\text{uranium}_j), \tau_0^2)$$

$$\alpha_{1j} \sim N(\beta_{10}, \tau_1^2)$$

■ Demonstration in R and rube()/JAGS...

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16

Example 3: CD4 in HIV-positive youth

- See R handout, and demonstration in class

Summary

- JAGS automates MCMC
 - Specify $(\text{posterior}) \propto (\text{level 1}) \times (\text{level 2}) \times \dots$
in an R-like language
 - JAGS designs and runs the MCMC for you
- rube() makes JAGS less painful
 - Gelman & Hill use R2WinBUGS(), we will use rube()
- Partial summaries of parameter estimates, and good initial graphs, come easy in rube().
 - $\hat{R}_{\text{hat}} < 1.2$ is a handy “convergence diagnostic”