

and how it can be implemented in BUGS.

- Discuss Bayesian inference for non-linear and non-Gaussian mixed models and samples using MCMC.
- In particular consider Monte Carlo methods and how to obtain Monte software).

models (to provide knowledge of the likelihood function for such models) to review methods for computation of the likelihood function for such models.

- Discuss likelihood-based inference for non-linear and non-Gaussian mixed

Purpose of workshop

Data example: cherries

$$Y_{sIB^b} = \begin{cases} 0 & \text{if bud is dead} \\ 1 & \text{if bud is fresh} \end{cases}$$

$b = 1, 2, \dots, BUD$.

$s = 1, \dots, 5$ STOCK (variety), $t = 1, \dots, 4$ TREE, $B = 1, 2, 3$ BRANCH,

Logistic regression but observations on same tree/branch may be correlated....
Are some stocks more sensitive to night frost than others?

3

1

Ramius Waggepetresen
Department of Mathematics
Aalborg University

Monte Carlo methods for hierarchical models

- Monte Carlo chain Monte Carlo using BUGS
- Bayesian inference using BUGS
- Analysis of sensitivity to specification of priors
- Non-normal random effects
- ?

Lecture IV

- Markov chain Monte Carlo using BUGS
- Bayesian inference using BUGS
- Analysis of sensitivity to specification of priors
- Non-normal random effects
- ?

Lecture III

- Markov chain Monte Carlo
- maximization of likelihood (derivatives)
- importance sampling
- importance sampling
- ?

Lecture II

- Computation of likelihood function
- Generalized linear mixed models
- Data examples
- Data examples
- ?

Lecture I

- Computation of likelihood function
- Generalized linear mixed models
- Data examples
- Data examples
- ?

Outline

No analytical solution.

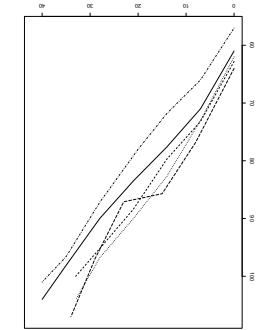
$$\text{Likelihood function } u = (u_1, \dots, u_n) = \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)}$$

$$\text{Conditional density: } f_{U|Y}(u|y) = \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} = \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)}$$

$$\begin{aligned} d\theta &= ((\theta_i u_i) / (1 + \exp(\theta_i u_i))) d\theta_i \\ \log(p_j) / (1 - p_j) &= \theta_j + \theta_i \\ Y_{il}|U_l = u_l &\sim \text{Bern}(p_j) = \text{binomial}(1, p_j), l = 1, \dots, n \\ U_l &\sim N(0, \tau^2), j = 1, \dots, n \end{aligned}$$

Non-normal example: Logistic regression with random effects (simple exp. design)

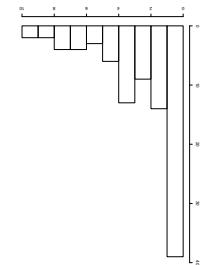
Growth curve with random coefficient?



Y_{lt} weight of pig p at time t .

Data example: growth curves for pigs

Non-nested random effects REPL, REPL*WEK and REPL*Z*L (=PEN)



Histogram of observations:

$r = 1, \dots, 6$ REPL.

$Z = 0, 1$, treatment $l = 0, 1$, week $w = 1, \dots, 4$ WEK, and replicate

Y_{lwtr} number of aggressions under shower within 48 minutes for treatment

$$\begin{aligned} Y &= X\beta + Z\gamma + \epsilon \\ \epsilon &\sim N(0, \tau^2) \\ \text{multivariate normal.} \\ \text{Likelihood function } (\theta, \sigma^2, \tau): &= \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \prod_{i=1}^n \frac{\exp(-\theta_i u_i)}{\exp(\theta_i u_i + \theta_i)} \\ T(\theta) &= (\theta, \sigma^2, \tau) = \int \exp(-\theta_i u_i) \exp(-\theta_i u_i) \exp(-\theta_i u_i) d\theta_i \end{aligned}$$

$$\begin{aligned} Y &= X\beta + Z\gamma + \epsilon \\ \epsilon &\sim N(0, \tau^2) \\ \text{General formulation:} \\ Y_{lwtr} &= \epsilon_{lwtr} + U_{lwtr} \end{aligned}$$

$$\begin{aligned} Y_{lwtr} &= \epsilon_{lwtr} + U_{lwtr} \\ U_{lwtr} &\sim N(0, \tau^2) \\ U_{lwtr} &\sim N(0, \tau^2) \quad \epsilon_{lwtr} \sim N(0, \sigma^2) \text{ independent} \\ Y_{lwtr} &\sim N(0, \tau^2 + \sigma^2) \end{aligned}$$

Normal linear mixed model (cherries design):

Data example: behavioral data for pigelets

Implemented in SAS macro `GLIMMIX`.

Asymptotic results require increasing number of observations for each random effect.

PQL estimates less accurate than ML.

$$f(y; \theta) = f(y|u; \beta, \phi) f(\phi|u; \tau).$$

PQL estimates θ and u maximize joint density

$$\theta = (\beta, \tau, \phi)$$

Penalized quasilikelihood

The computational problem

Likelihood integrable except when $Y|U$ is normal and $g(u) = u$.

- $Y|U$ Bernoulli with $g(x) = \log(x/(1-x))$ and $\phi = 1$

- $Y|U$ Poisson with $g(x) = \log(x)$ and $\phi = 1$

Examples:

$$X|U \sim n = \Omega|u; \beta, \phi$$

Conditional distribution parameterised by u and dispersion parameter ϕ

$$g(u) = \mathbb{E}[X|U]$$

Link function

$$X|U + \beta Z = u$$

Linear predictor

$$U \sim N^d(\tau)$$

Generalized linear mixed models

Example (growth curve for pigs):

Nonlinear normal mixed models

$$X = m(t; U^d, a, q, c) + e^d$$

$$m(t; U^d, a, q, c) = \exp(a - q \exp(-t(c + U^d)))$$

$$U^d \sim N(0, \tau^d) \text{ and } e^d \sim N(0, \sigma^d) \text{ independent}$$

Example (growth curve for pigs):

Nonlinear normal mixed models

and Z .

May factorize into lower-dimensional integrals depending on structure of $\Sigma(\tau)$

$$\int \prod_{i=1}^n \int_{\Omega_i} f(\phi, \beta, u_i; \tau) du_i \dots d\Omega = (\theta) \Omega$$

n -dimensional integral:

$$X|U \sim n = \Omega|u; \beta, \phi \quad \text{independent}$$

$$\beta Z + \beta X = u$$

$$((\tau) \Sigma(0, 0, N^d)) = \Omega$$

The computational problem

$$\zeta((\theta)^{O_{MS}}T - (\theta^{\circ}_{:w\Omega}|\tilde{h})f) \sum_{M=1}^{L=u} \frac{L-M}{L}$$

$W \cdots I = u \cdot (g \cdot u \cap h) f$

Estimate $\text{Var}_f(y|U_1; \beta)$ using empirical variance estimate based on

Often $\text{Var}(y|U_1; \beta)$ is large so large M is needed.

$$\text{Var}(L^{SMC}(\theta)) = \frac{M}{\prod_i M_i}$$

Monte Carlo variance:

where $U_m \sim N(0, \tau_z)$ independent.

$$(\theta) \cdot_{\text{uM}} T = (\theta) \circ_{\text{MS}} T \approx (\theta) \cdot_{\text{L}} T = np(\zeta) \cdot_{\text{L}} n \cdot_{\text{L}} (\theta) \int^{\mathbb{H}} = (\theta) T$$

Simple Monte Carlo:

NB: $\cdot ((n)_\mu \delta / [1 - n]) N \approx \hbar = \lambda |U$ os $((n)\delta)dx \propto (\hbar)f/(n)f(n|\hbar)f = (\hbar|n)f$

Also possible for higher dimensions.

$$\int_{-\frac{2\pi}{n}}^{\frac{2\pi}{n}} \left| \sum_{k=1}^{n-1} e^{ikx} \right|^2 dx = n \int_{-\frac{2\pi}{n}}^{\frac{2\pi}{n}} \left(\sum_{k=1}^{n-1} e^{ikx} \right)^2 dx = n \int_{-\frac{2\pi}{n}}^{\frac{2\pi}{n}} \left(\sum_{k=1}^{n-1} e^{ikx} \right) \left(\sum_{j=1}^{n-1} e^{-ijx} \right) dx =$$

i.e. $\exp(g(u))$ proportional to density for $N(u, -I/g''(u))$.

$$((n)_\mu b -)_{\zeta}(\eta - n)\frac{\zeta}{1} - (\eta)b = (\eta)_\mu b_{\zeta}(\eta - n)\frac{\zeta}{1} + (\eta)_\nu b(\eta - n) + (\eta)b = (n)b$$

Laylor expansion around u :

Let $y(n) = \log f(n)$ and choose n so $y(n) = \arg \max y(n)$

Laplace approximation

- Simple Monte Carlo (small dimensions).
 - Importance sampling.

Possibilities:

$$np(\omega^n)f(\vartheta^n|\kappa)f\int^{\mathbb{H}} = (\theta)T$$

Compute

One-dimensional case

8

$$\text{...} \cdot np(\cdot n) f \left[\text{...} np(\cdot m n) f \prod^m \right] \left[\text{...} lz np(\cdot lz n) f \prod^{lz} \right] \left[(\cdot n \cdot \cdot m n \cdot \cdot lz n) f \prod^{mlz} \right] 6^{\frac{mlz}{2}} \int \prod^l = (\theta) T$$

Behavioral: 9 dimensional

$$\int \prod_{i=1}^{B,q} \left[f_{\theta}(y_i | x_i) \right] \left[f_{\theta}(x_i | \phi_i) \right] = \text{Product of marginal posteriors}$$

Cherries: 4 dimensional

$${}^d n p({}^d n) f \left[({}^d n | {}^{id} \hat{n}) f \prod_{d_n}^{\mathbb{I}=t} \right] {}^u \int \coprod_u^{\mathbb{I}=d} = {}^u n p \dots {}^u n p \left[({}^d n | {}^{id} \hat{n}) f \prod_{d_n}^{\mathbb{I}=t} ({}^d n) f \right] \coprod_u^{\mathbb{I}=d} {}^u \int = (\theta) T$$

Growth curves: one dimensional

Examples:

same importance sampling distribution used).
 NB coincides with derivatives of Monte Carlo approximation of likelihood if
 Replace $u(\theta_m)$ and $i(\theta_m)$ by Monte Carlo estimates.

$$\begin{aligned} h &= X \left| \frac{\partial}{\partial \theta} \log f(y; \theta) \right| + (\theta; \Omega; h) f(y; \theta; \Omega; h) - \frac{\partial}{\partial \theta} \log f(y; \theta; \Omega; h) = (\theta) n \frac{\partial}{\partial \theta} - = (\theta) i \\ h &= X \left| \frac{\partial}{\partial \theta} \log f(y; \theta; \Omega; h) \right| = (\theta) n \frac{\partial}{\partial \theta} \end{aligned}$$

$$\begin{aligned} &:: \\ &= (\theta) n \frac{\partial}{\partial \theta} + (\theta) n \frac{\partial}{\partial \theta} = (\theta) n \frac{\partial}{\partial \theta} \\ &= (\theta) n \frac{\partial}{\partial \theta} \end{aligned}$$

Newton-Raphson iteration:

$$0 = (\theta) T \log \frac{\partial}{\partial \theta} = (\theta) n$$

Want to solve

Maximization of likelihood

19

So we can estimate ratio $L(\theta)/L(\theta^0)$ where $L(\theta^0)$ is unknown constant.
 This suffices for finding MLE:
 $\arg \max_{\theta} L(\theta) = \arg \max_{\theta} \frac{L(\theta)}{L(\theta^0)}$

$$\begin{aligned} h &= X \left| \frac{(\theta^0) T \log f(y; \theta^0; \Omega; h) f}{(\theta^0) T \log f(y; \theta^0; \Omega; h) f} \right| \mathbb{E} = \frac{(\theta^0) T}{(\theta^0) T} \Leftrightarrow h = X \left| \frac{(\theta^0) T \log f(y; \theta^0; \Omega; h) f}{(\theta^0) T \log f(y; \theta^0; \Omega; h) f} \right| \mathbb{E} (\theta^0) T = \\ &= np(\theta^0; h; n) f \frac{(\theta^0) T \log f(y; \theta^0; \Omega; h) f}{(\theta^0) T \log f(y; \theta^0; \Omega; h) f} \int (\theta^0) T = np(n) h \frac{(n) h}{(\theta^0) T \log f(y; \theta^0; \Omega; h) f} \int = (\theta^0) T \\ &\text{Then} \\ &(\theta^0) T / (\theta^0; n) f (\theta^0; n; h) f = (\theta^0; h; n) f = (n) h \end{aligned}$$

Possibility: Consider fixed θ^0 :

18

Simulation straightforward,
 so small variance.

$$\frac{h(n) f(y; \theta^0; \Omega; h) f}{f(y; \theta^0; \Omega; h) f} \approx \text{const}$$

Use $h(\cdot)$ density for $N(u, -1/g''(u))$ or $t_a(u, -1/g''(u))$ -distribution:

$$\begin{aligned} \text{Laplace: } U | \Lambda &\approx N(u, -1/g''(u)) \\ \text{const} &= 1/L(\theta) \end{aligned}$$

$$f(y; \theta^0; \Omega; h) f = \text{const} f(u; \theta^0; \Omega; h) f \Leftrightarrow (\theta^0; h; n) f = \text{const}$$

Possibility: Note

17

$\text{Var} L_{IS, h}(\theta) > \infty$ if $f(y; a; \theta) f(a; \theta) / h(a)$ bounded (i.e. use $h(\cdot)$ with heavy tails).
 Find h so $\text{Var} \frac{f(y; \theta) f(\theta; \Omega; h)}{h(\Lambda)} f(\Lambda; \theta; \Omega; h)$ small.

$$\begin{aligned} L(\theta) &\approx L_{IS, h}(\theta) \quad \text{where } V_m h(\Lambda_m) f(\Lambda_m; \theta; \Omega; h) f(\Lambda_m; \theta; \Omega; h) \sum_m^M \frac{W}{1} = (\theta) L(\theta) \\ &\text{where } V \sim h(\cdot). \\ \frac{(\Lambda) h}{(\theta) f(\Lambda; \theta; \Omega; h) f} \mathbb{E} &= np(n) h \frac{(n) h}{(\theta) f(\Lambda; \theta; \Omega; h) f} \int = np(\theta) f(\theta; \Omega; h) f \int = (\theta) T \end{aligned}$$

$h(\cdot)$ probability density on \mathbb{R} .

Importance sampling

Very small probability $< \epsilon$ that a simulation from $\text{Unif}([0, 1]^n)$ falls in $\times_{i=1}^n [y_i - \epsilon/2, y_i + \epsilon/2] \cup [0, 1]^n$.

I.e. $(U_1, \dots, U_n) | X = y_1, \dots, X_n = y_n$ lives on set of volume $< \epsilon^n$.

$$(U_1, \dots, U_n) | X = y_1, \dots, X_n = y_n \sim \text{Unif}(\times_{i=1}^n [y_i - \epsilon/2, y_i + \epsilon/2] \cup [0, 1]^n)$$

Then

$$X | U_i = u_i \sim \text{Unif}([\theta_i - \epsilon/2, \theta_i + \epsilon/2])$$

and

$$(U_1, \dots, U_n) \sim \text{Unif}([0, 1]^n)$$

Suppose

Course of dimensionality for a toy example

$$\mathbb{E}[y|u] f \sim \frac{1}{M} \sum_{m=1}^M \frac{W}{L} = (\theta = X | U) \mathbb{E}$$

of $U|Y = y$ or importance sampling.

Compute Monte Carlo estimate of $\mathbb{E}(U|Y = y)$ using conditional simulations

is minimal for $\hat{X} = \mathbb{E}(X|Y)$.

$$\mathbb{E}(X - \hat{X})^2$$

I.e.

conditional expectation $\mathbb{E}(X|Y)$.

Minimum mean square error predictor \hat{X} of X given observation of Y is

Consider pair of random variables (X, Y) .

Further use of Monte Carlo: Prediction of U

Instead we can use MCMC (later).

Extremely small accept probability in high dimensions (course of dimensionality).

Typically small probability for accept (cf. simple Monte Carlo Method).

2. Return V if $W \leq f(y|V; \tau_0)/B$; otherwise go to 1.

1. Generate $V \sim f(\cdot; \tau_0)$ and $W \sim \text{Unif}(0, 1)$

Rejection sampling:

$$\text{where } B = \sup_u f(y|u; \tau_0).$$

$$(f(y|\theta_m) f(u; \tau_0)) \geq B f(u; \tau_0)$$

Note

Conditional simulation of $U|Y = y$

where $U_m \sim h(\cdot)$.

$$\mathbb{E}^{\theta_m} [\log f(y, U; \theta; \tau_0)] \approx \frac{1}{M} \sum_{m=1}^M \frac{W}{\log f(y, U_m; \theta)} f(y|U_m; \tau_0) / L(\theta_m)$$

or importance sampling

$$\mathbb{E}^{\theta_m} [\log f(y, U; \theta; \tau_0)] \approx \frac{1}{M} \sum_{m=1}^M \frac{W}{\log f(y, U_m; \theta)} f(y|U_m; \theta) / L(\theta_m)$$

MCEM (Monte Carlo EM): use Monte Carlo approximation

$$\mathbb{E}^{\theta_m} [\log f(y, U; \theta; \tau_0)] \approx \mathbb{E}[y|U = y]$$

Suppose θ_m is current parameter value. Obtain θ_{m+1} by maximizing

EM-algorithm:

$$\mathbb{E}(\bar{U}_m) \approx \frac{1}{M} \sum_{i=1}^M U_i$$

I.e. for large m , $\bar{U}_m \sim \pi(\cdot)$ and distribution of $\bar{U}_m \leftarrow \pi(\cdot)$
so that

$$U_1, U_2, U_3, \dots \quad (U_m = (U_1, \dots, U_m))$$

Generate ergodic Markov chain
Markov chain Monte Carlo:

Suppose $U = (U_1, \dots, U_n) \sim \pi(\cdot)$ where $\pi(\cdot)$ is a complicated probability distribution.

MCMC

A: additive genetic relationship matrix (depending on pedigree).

$$(U_1, \dots, U_n, \bar{U}_1, \dots, \bar{U}_n) \sim N(0, G \otimes A)$$

$$Y_i | U_i = u_i, \bar{U}_i = \bar{u}_i \sim N(u_i + u_i, \exp(\bar{u}_i + u_i))$$

U_i, \bar{U}_i random genetic effects influencing size and variability of Y_i :
 Y_i size of j th litter of i th pig.

Example: quantitative genetics (Sorensen and Waagepeterson 2003)

Observations X_i are weed counts at spatial locations (x_i, y_i) $i = 1, \dots, 270$
 $X_i | U_i$ is Poisson where U_i random effect associated with (x_i, y_i) (soil properties).
 U_i is Markov chain Monte Carlo:
 $\text{Cov}(U_i, U_j) = \tau^2 \exp(-d_{ij}/\alpha)$

where d_{ij} is distance between (x_i, y_i) and (x_j, y_j)

$$\text{Cov}(U_i, U_j) = \tau^2 \exp(-d_{ij}/\alpha)$$

NB: high dimension $n > 6000$.

p : coefficient of genetic correlation between U_i and U_j .

$$G = \begin{bmatrix} p & & & \\ & p & & \\ & & p & \\ & & & p \end{bmatrix}$$

Can not be factorized into lower dimensional densities.

$$f(u_1, \dots, u_n | y) \propto \prod_i f(y_i | u_1, \dots, u_n)$$

Then n -dimensional conditional density for $U | Y = y$:

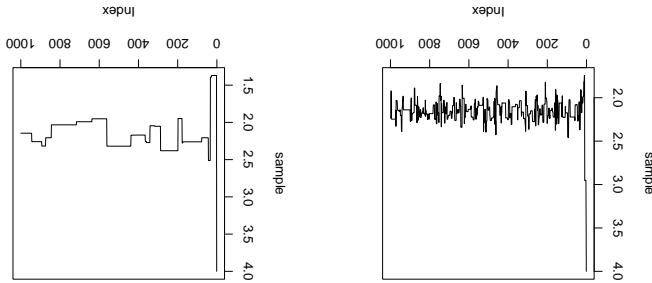
$$\text{Cov}(U_i, U_j) \neq 0 \text{ for all } i, j$$

with

$$U = (U_1, \dots, U_n) \sim N^u(0, \Sigma)$$

Consider

Conditional simulation for high-dimensional U : Markov chain Monte Carlo



$$\sigma^2_{\text{prop}} = 0.2 \text{ (accept rate } 24\%) \quad \sigma^2_{\text{prop}} = 40 \text{ (accept rate } 1\%)$$

Plots of U_1, U_2, U_3, \dots :

Convergence of Markov chains for simple example

NB: need only know π up to constant of proportionality.

$$= \frac{\prod_{i=1}^{10} f(y_i | \pi) \exp(u_m + \beta) f(u_m; \tau_2)}{\prod_{i=1}^{10} f(y_i | \pi) \exp(V_{m+1} + \beta) f(V_{m+1}; \tau_2) / L(\theta)}$$

out):

Random walk Metropolis ratio (normalising constant $L(\theta) = f(y; \theta)$ cancels

where $f(y_i | \lambda)$ density for Poisson distribution of intensity λ .

$$\pi(u|y) = \prod_{i=1}^{10} f(y_i | \pi) \exp(u + \beta) f(u; \tau_2) / L(\theta)$$

Simple example (Exercise 1):

$$\min\{1, \frac{\pi(u_m + 1|u_m)}{\pi(u_m|u_{m+1})}\} = \min\{1, \frac{\pi(u_m)}{\pi(u_{m+1})}\}$$

so Metropolis-Hastings ratio reduces to Metropolis ratio:

$$(a|n)b = (n|a)b$$

Then

where σ^2_{prop} is the proposal variance.

$$V_{m+1} \sim N(u_m, \sigma^2_{\text{prop}})$$

Ex: random walk Metropolis:

Under mild conditions of irreducibility and aperiodicity this produces an ergodic Markov chain with stationary distribution given by $\pi(\cdot)$.²⁹

$$\text{accept } U_{m+1} = V_{m+1}, \text{ otherwise } U_{m+1} = u_m.$$

$$\min\{1, \frac{\pi(u_m|u_{m+1})}{\pi(u_{m+1}|u_m)}\}$$

2. With probability

1. Conditional on $U_m = u_m$ generate proposal $V_{m+1} \sim q(\cdot|u_m)$.

Given initial state U_1 generate U_2, U_3, \dots as follows:

defined for all $u \in \mathbb{R}^p$ and easy to sample.

$$q(a|u), a \in \mathbb{R}^n$$

Basic ingredient: proposal density

Joint updating Metropolis-Hastings algorithm:

Repeat for $i = 1, \dots, n$

$$\text{accept } U_{m+1} = V_{m+1}, \text{ otherwise } U_{m+1} = u_m.$$

$$\min\{1, \frac{\pi(u_m|_{I+mu})\Lambda^i(u_m|_{I+mu})}{(\Lambda(u_m|_{I+mu})\Lambda^i(u_m|_{I+mu}))}\}$$

2. With probability

$$\Lambda_{m+1} = (u_m, \dots, u_{m-1}, \Lambda_{m+1}, u_{m+1}, \dots, u_m)$$

1. Conditional on $U_m = u_m$ generate $V_{m+1} \sim q_i(\cdot|u_m)$ and let

Update of i th component:

Update one component in each iteration.

Single-site Metropolis-Hastings

Implementation of MCMC using BUGS (Bayesian analysis using

Example:

Model specification in BUGS: hierarchical/directed acyclic graph (DAG).

Gibbs sampling).

DAG:

BUGS code

```
y2 ~ dnorm(u,sigma2rec)
y1 ~ dnorm(u,sigma2rec)
u ~ dnorm(0.0,tau2rec)
sigma2rec <- 1/0.05
```

```
tau2rec <- 1
```

```
model f
```

```
BUGS code
```

DAG:

BUGS code

$$Y_1, Y_2 | U = u, \tau^2, \sigma^2 \sim N(u, \sigma^2), \text{ conditionally independent}$$

$$U | \tau^2, \sigma^2 \sim N(0, \tau^2)$$

$$\tau^2 = 1 \quad \sigma^2 = 0.05$$

No need to choose a proposal variable.

so all proposals are accepted.

$$1 = \frac{(\bar{u}_m|_{I+mu})\pi(\bar{u}_m|_{I+mu})\pi(\bar{u}_m|_{I+mu})}{(\bar{u}_m|_{I+mu})\pi(\bar{u}_m|_{I+mu})\pi(\bar{u}_m|_{I+mu})} = \frac{\pi(u_m|_{I+mu})\Lambda^i(u_m|_{I+mu})}{\pi(u_m|_{I+mu})\Lambda^i(u_m|_{I+mu})}$$

Then $b(\Lambda^i u_m|_{I+mu}) = (\Lambda^i u_m|_{I+mu})$ and

Gibbs sampler: $\Lambda^i u_m|_{I+mu} = U^i|_{I+mu}$, $i \neq i$.

$$\min\{1, \frac{\pi(u_m|_{I+mu})\Lambda^i(u_m|_{I+mu})}{\pi(u_m|_{I+mu})\Lambda^i(u_m|_{I+mu})}\} = \min\{1, \frac{\pi(u_m|_{I+mu})}{\pi(u_m|_{I+mu})}\}$$

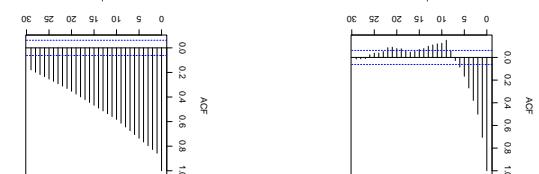
Random walk Metropolis: $\Lambda^i u_m|_{I+mu} \sim N(u_m, \sigma^{prop})$ and

Examples

so small autocorrelation advantageous.

$$\text{Var} \frac{1}{M} \sum_{m=1}^M U_m = \frac{\text{Var} U}{M} = \sum_{m=1}^M \sum_{n=1}^M d(|m - n|)$$

Note:



$\sigma^{prop} = 0.2$ (quick mixing) $\sigma^{prop} = 40$ (slow mixing)

Plot of autocorrelation $p(k) = \text{Corr}(U_m, U_{m+k})$:

Autocorrelation/mixing:

$$U^{st} \sim N^a(0, \tau_{TB}^2) \quad U^{stB} \sim N(0, \tau_{TB}^2) \text{ independent}$$

Mean 1 and variance 1000 for $\Gamma(0.001, 0.001)$.

$$\text{If } 1/\tau_2 \sim \Gamma(a, \beta) \text{ then density of } \tau = (1/\tau_2)^{-1/2} \text{ is}$$

$$\frac{\Gamma(a)}{\beta^a} \exp(-\omega a - \beta \exp(-\omega)) \approx \text{const if } a \text{ and } \beta \text{ small}$$

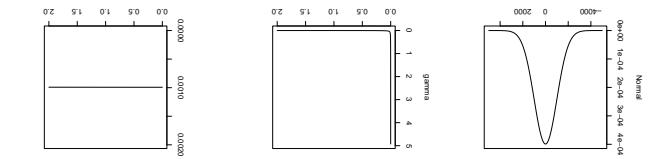
$$\text{If } 1/\tau_2 \sim \Gamma(a, \beta) \text{ then density of } \omega = -\log(\tau_2) \text{ is}$$

$$((\mathcal{H})_1 - \delta)/((\mathcal{H})_1 - \delta) x f = (\mathcal{H})_1 x f$$

Then

Suppose X has density f_X and $Y = g(X)$.

Transformation



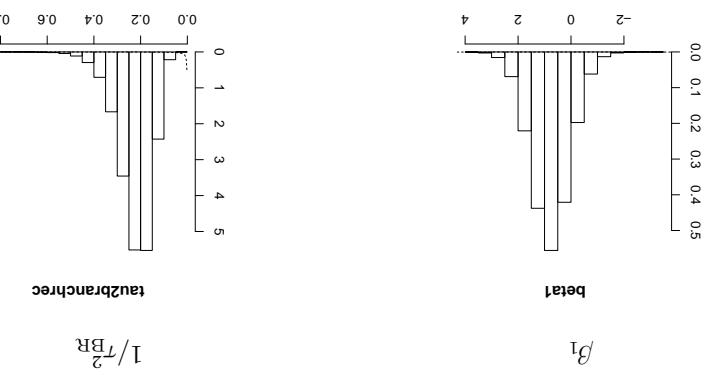
$$N(0, 10^6) \quad \Gamma(0.001, 0.001) \quad \text{Prior for } \log(\tau_{BR}^2)$$

$$1/\tau_{TR}, 1/\tau_{BR}^2 \sim \Gamma(0.001, 0.001) \quad \beta_s \sim N(0, 10^6)$$

Priors:

Example: Bayesian analysis analyses for cherries

Posterior mean 0.77 and variance 0.52 Posterior mean 0.23 and variance 0.005



Note: if (θ_m, U_m) sample from $p(\theta, u|y)$ then θ_m sample from $p(\theta|y)$.

Compute posterior expectations, variances etc. using samples from $p(\theta, u|y)$.

$$d(\theta)d(\theta|n)f(\theta|n, u|y)\propto f(y|\theta)d(\theta)$$

(demarginalize)

Likelihood $L(\theta)$ given data y is unknown so consider augmented posterior

$$d(\theta|y) = \frac{f(y)}{f(y|\theta)p(\theta)} \propto L(\theta)p(\theta)$$

Posterior distribution (knowned obtained by observing data y):

Introduce prior $p(\theta)$ for unknown parameters.

Some posterior results for cherries

Bayesian inference for hierarchical model

Priors and posteriors for β_1 and $1/\tau_{BR}^2$:

$$1/\tau_{BR}^2$$

$$\beta_1$$

tau_2branching

39

37

Hence, computation of $L(\theta)$ avoided.

Posterior mean 0.77 and variance 0.52 Posterior mean 0.23 and variance 0.005

- heterogeneity: a case study. *Genetic Research*. to appear.
- Sorensen, D. and Waagepeterson, R. (2003). Normal linear models with genetically structured variance. *Biometrika*, 90, 162–170.
- Robert, C. and Casella, G. (1999). *Monte Carlo Statistical Methods*. Springer-Verlag, New York.
- McQuillich, C. E. (1992). Maximum likelihood algorithms for generalized linear mixed models. *J. Am. Statist. Ass.* 87, 657–699.
- Geyer, C. J. and Thompson, E. A. (1992). Constrained Monte Carlo maximum likelihood for dependent data. *Journal of the Royal Society of Statistics Series B* 54: 254–272.
- Bavans, M. and Swartzi, T. (1995). Methods for approximating integrals in statistics with special emphasis on Bayesian integration problems. *Statistica Sinica* 10: 280–286.
- Christensen, O. F. and Waagepeterson, R. (2002). Bayesian prediction of spatial count data using generalized linear mixed models. *Biometrika* 89, 265–285.
- Booth, J. G. and Hobert, J. P. (1999). Maximizing generalized linear mixed model likelihoods with an automated Bayesian EM algorithm. *J. R. Statist. Soc. B* 61: 280–286.
- Geyer, C. J. and Thompson, E. A. (1992). Monte Carlo maximum likelihood for generalized linear mixed models. *Journal of the Royal Society of Statistics Series B* 54: 657–699.
- Robert, C. and Casella, G. (1999). *Monte Carlo Statistical Methods*. Springer-Verlag, New York.
- Sorensen, D. and Waagepeterson, R. (2003). Normal linear models with genetically structured variance. *Biometrika*. to appear.

Suppose we have sample θ_m from $p_1(\theta|y)$ and we want to compute

$$\mathbb{E}_2[h(\theta|y)] = \int h(\theta)p_2(\theta|y)d\theta$$

$$p_2(\theta|y) = f(y|\theta)p_2(\theta)/f_2(y)$$

and

$$p_1(\theta|y) = f(y|\theta)p_1(\theta)/f_1(y)$$

Consider two priors $p_1(\theta)$ and $p_2(\theta)$ which yields posteriors

Sensitivity to prior specification

References

$$\begin{aligned} & \left(\frac{d}{d\theta} \log p_1(\theta) \right)_{\theta=\theta_m} = \frac{1}{M} \sum_{m=1}^M \frac{\partial}{\partial \theta} \log p_1(\theta_m) \approx \theta \frac{\partial}{\partial \theta} \log p_1(\theta) \Big|_{\theta=\theta_m} \\ & \text{Moreover:} \\ & \theta \frac{\partial}{\partial \theta} \log p_1(\theta) \approx \frac{1}{M} \sum_{m=1}^M \frac{\partial}{\partial \theta} \log p_1(\theta_m) \Big|_{\theta=\theta_m} \\ & \theta \frac{\partial}{\partial \theta} \log p_1(\theta) = \theta \frac{\partial}{\partial \theta} \log \left(\frac{f_1(y)}{f_2(y)} \right) = \theta \frac{\partial}{\partial \theta} \log \left(\frac{f_1(y| \theta)}{f_2(y| \theta)} \right) = \theta \frac{\partial}{\partial \theta} \log \left(\frac{h(\theta|y)}{h(\theta)} \right) = \theta \mathbb{E}_2[h(\theta|y)] - \mathbb{E}_2[h(\theta)] \end{aligned}$$

Reuse sample using importance sampling:

Plot of density for $\tau = (1/\tau^2)^{-1/2}$ when $1/\tau^2 \sim \Gamma(0.001, 0.001)$

