Bayesian Regression

Bayesian Modeling for Socio-Environmental Data

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Outline

- Understand Bayesian inference using familiar examples.
- Appreciate one-to-one relationship between math and JAGS code.
- Learn some programming tricks.

Notation

$$\underbrace{g(\boldsymbol{\theta}, x)}_{\text{deterministic model}}$$

Embedding the deterministic model

The joint distribution in a simple Bayesian model is the product of the likelihood and the priors.

$$[\theta, \sigma^2] \propto [y \mid \theta, \sigma^2] [\theta] [\sigma^2]$$

A deterministic model of an ecological process is embedded in the likelihood like this. . .

$$\left[\boldsymbol{\theta},\sigma^{2}\right]\propto\left[\boldsymbol{y}\mid\boldsymbol{g}(\boldsymbol{\theta},\boldsymbol{x}),\sigma^{2}\right]\left[\boldsymbol{\theta}\right]\left[\sigma^{2}\right]$$

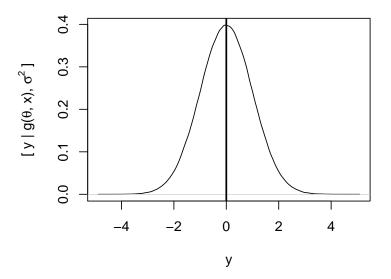
The general set up for a simple Bayesian regression model

$$\mu_i = \underbrace{g(\boldsymbol{\theta}, x_i)}_{\text{deterministic model}}$$

$$y_i \sim \underbrace{f(\mu_i, \sigma^2)}_{\text{stochastic model}}$$

$$\left[\boldsymbol{\theta}, \sigma^2\right] \propto \left[y \mid g(\boldsymbol{\theta}), \sigma^2\right] \left[\boldsymbol{\theta}\right] \left[\sigma^2\right]$$

A graphical representation



You don't have to be normal!

Data (y-values)	Distribution	Mean function	Link
continuous, real valued	normal	$\mu = \beta_0 + \beta_1 x$	NA
discrete, strictly positive	Poisson	$\mu = \mathrm{e}^{eta_0 + eta_1 x}$	$\log(\mu) = \beta_0 + \beta_1 x$
0 or 1	Bernoulli	$\mu = \frac{\exp(\beta_0 + \beta_1 x)}{\exp(\beta_0 + \beta_1 x) + 1}$	$\operatorname{logit}\left(\mu\right) = \operatorname{log}\left(\frac{\mu}{1-\mu}\right) = \beta_0 + \beta_1 x$
0 – 1	beta	$\mu = \frac{\exp(\beta_0 + \beta_1 x)}{\exp(\beta_0 + \beta_1 x) + 1}$	$logit(\mu) = log(\frac{\mu}{1-\mu}) = \beta_0 + \beta_1 x$
continuous, strictly positive	lognormal or gamma	$\mu = \mathrm{e}^{\beta_0 + \beta_1 x}$	$\log\left(\mu\right) = \beta_0 + \beta_1 x$

Lots of flexibility as a modeler

Continent-wide Adélie penguin population dynamics

$$\begin{array}{lcl} z_{s,y} & \sim & \operatorname{lognormal}(z_{s,y} \mid g(\beta_1,\beta_2,z_{s,y-1}),\sigma_p) \\ g(\theta) & = & \operatorname{log}(z_{s,y-1}e^{\beta_1+\beta_2\operatorname{wsic}_{s,y}+\beta_3\operatorname{ssic}_{s,y}}) \end{array}$$

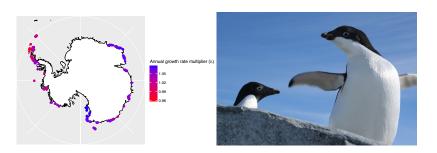


Photo c/o Heather J. Lynch

Normal data, continuous and real valued

```
[\beta_0, \beta_1, \sigma \mid \mathbf{y}] \propto \prod_{i=1}^{N} \operatorname{normal}(y_i \mid g(\beta_0, \beta_1, x_i), \sigma^2) \times \operatorname{normal}(\beta_0 \mid 0, 1000) \operatorname{normal}(\beta_1 \mid 0, 1000) \times \operatorname{uniform}(\sigma \mid 0, 100)
g(\beta_0, \beta_1, x_i) = \beta_0 + \beta_1 x_i
```

```
b0 ~ dnorm(0, .001)
b1 ~ dnorm(0, .001)
sigma ~ dunif(0, 100)
tau <- 1/sigma^2
for (i in 1:length(y)){
   mu[i] <- b0 + b1 * x[i]
   y[i] ~ dnorm(mu[i], tau)
}</pre>
```

Poisson, discrete and positive

```
[\beta_0, \beta_1 \mid \mathbf{y}] \propto \prod \text{Poisson}(y_i \mid g(\beta_0, \beta_1, x_i))
                                      \times \text{normal}(\beta_0 \mid 0, 1000) \text{ normal}(\beta_1 \mid 0, 1000)
           g(\beta_0, \beta_1, x_i) = e^{\beta_0 + \beta_1 x_i}
b0 \sim dnorm(0, .001)
b1 \sim dnorm(0, .001)
for(i in 1:length(y)){
   log(mu[i]) \leftarrow b0 + b1 * x[i]
   y[i] ~ dpois(mu[i])
```

or

}

```
mu[i] \leftarrow exp(b0 + b1 * x[i])
y[i] ~ dpois(mu[i])
```

Bernoulli, data 0 or 1 (aka logistic)

```
[\beta_0, \beta_1 \mid \mathbf{y}] \propto \prod \text{Bernoulli}(y_i \mid g(\beta_0, \beta_1, x_i))
                                         \times normal(\beta_0 \mid 0, 2.56) normal(\beta_1 \mid 0, 2.56)
             g(\beta_0, \beta_1, x_i) = \frac{e^{\beta_0 + \beta_1 x_i}}{e^{\beta_0 + \beta_1 x_i} + 1}
b0 \sim dnorm(0, .368)
b1 \sim dnorm(0, .368)
for(i in 1:length(y)){
   logit(p[i]) \leftarrow b0 + b1 * x[i]
       y[i] ~ dbern(p[i])
```

or

```
p[i] <- inv.logit(b0 + b1 * x[i])
y[i] ~ dbin(p[i])</pre>
```

lognormal, data continuous and > 0 (log link)

$$\begin{split} \left[\beta_0,\beta_1,\sigma\mid\mathbf{y}\right] &\propto & \prod_{i=1}^N \operatorname{lognormal} \left(y_i\mid \operatorname{log} \left(g\left(\beta_0,\beta_1,x_i\right)\right),\sigma^2\right) \\ &\times \operatorname{normal} \left(\beta_0\mid 0,1000\right) \operatorname{normal} \left(\beta_1\mid 0,1000\right) \\ &\times \operatorname{uniform} \left(\sigma\mid 0,5\right) \\ g\left(\beta_0,\beta_1,x_i\right) &= & e^{\beta_0+\beta_1x_i} \end{split}$$

Talk about the interpretation of σ .

```
b0 ~ dnorm(0, .001)
b1 ~ dnorm(0, .001)
sigma ~ dunif(0, 5)
tau <- 1/sigma^2
for(i in 1:length(y)){
   mu[i] <- exp(b0 + b1 * x[i])
   y[i] ~ dlnorm(log(mu[i]), tau)
}</pre>
```

lognormal, data continuous and > 0 (not log link)

```
 [\beta_0, \beta_1, \sigma \mid \mathbf{y}] \propto \prod_{i=2}^{N} \operatorname{lognormal}(y_i \mid \operatorname{log}(g(\beta_0, \beta_1, y_{i-1})), \sigma^2) 
 \times \operatorname{normal}(\beta_0 \mid 0, 1000) \operatorname{normal}(\beta_1 \mid 0, 1000) 
 \times \operatorname{uniform}(\sigma \mid 0, 5) \operatorname{uniform}(y_1 \mid 1, 1E6) 
 g(\beta_0, \beta_1, y_{i-1}) = y_{i-1}e^{\beta_0 + \beta_1 y_{i-1}}
```

```
b0 ~ dnorm(0, .001)
b1 ~ dnorm(0, .001)
sigma ~ dunif(0, 5); tau <- 1/sigma^2
y[1] ~ dunif(1, 1E6)
for(i in 2:length(y)){
  mu[i] <- y[i-1] * exp(b0 + b1 * y[i-1])
  y[i] ~ dlnorm(log(mu[i]), tau)
}</pre>
```

lognormal, data continuous and > 0 (max function)

```
[\beta_0, \beta_1, \sigma \mid \mathbf{y}] \propto \prod_{i=1}^{N} \operatorname{lognormal}(y_i \mid \log(g(\beta_0, \beta_1, x_i)), \sigma^2) \\ \times \operatorname{normal}(\beta_0 \mid 0, 1000) \operatorname{normal}(\beta_1 \mid 0, 1000) \\ \times \operatorname{uniform}(\sigma \mid 0, 5) \\ g(\beta_0, \beta_1, x_i) = \max(\beta_0 + \beta_1 x_i, 0.0001)
```

```
b0 ~ dnorm(0, .001)
b1 ~ dnorm(0, .001)
sigma ~ dunif(0, 5)
tau <- 1/sigma^2
for(i in 1:length(y)){
   mu[i] <- b0 + b1 * x[i]
   y[i] ~ dlnorm(log(max(mu[i],.000001)), tau)
}</pre>
```

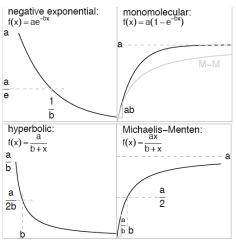
lognormal, data continuous and > 0 (priors)

```
 [\beta_0, \beta_1, \sigma \mid \mathbf{y}] \propto \prod_{i=1}^{N} \operatorname{lognormal}(y_i \mid \operatorname{log}(\mathbf{g}(\beta_0, \beta_1, x_i)), \sigma^2) 
 \times \operatorname{uniform}(\beta_0 \mid .001, 1000) \operatorname{uniform}(\beta_1 \mid .001, 1000) 
 \times \operatorname{uniform}(\sigma \mid 0, 5) 
 \mathbf{g}(\beta_0, \beta_1, x_i) = \beta_0 + \beta_1 x_i
```

Is this really a good idea?

```
b0 ~ dunif(.001, 1000)
b1 ~ dunif(.001, 1000)
sigma ~ dunif(0, 5)
tau <- 1/sigma^2
for(i in 1:length(y)){
   mu[i] <- b0 + b1 * x[i]
   y[i] ~ dlnorm(log(mu[i]), tau)
}</pre>
```

Nonlinear regression



Figures c/o Bolker, B. 2008. Ecological Models and Data in R. Princeton University Press, Princeton, NJ. USA.

Centering and standardizing

The remainder of the slides apply to all of the general linear models, but I will use a simple linear for normally distributed data as an example.

Centering predictor data

$$y_i = \beta_0 + \beta_1 (x_i - \bar{x})$$

Why complicate things...?

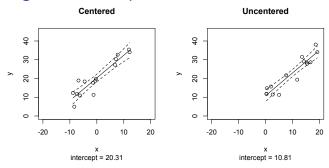
- To reduce autocorrelation in MCMC chain and speed convergence.
- To make the intercept more easily interpretable.

Centering predictor data

```
\begin{split} \left[\beta_{0},\beta_{1},\sigma\mid\boldsymbol{y}\right] &\propto & \prod_{i=1}^{N}\operatorname{normal}\left(y_{i}\mid\boldsymbol{g}\left(\beta_{0},\beta_{1},x_{i},\bar{x}\right),\sigma^{2}\right) \\ &\times\operatorname{normal}\left(\beta_{0}\mid0,1000\right)\operatorname{normal}\left(\beta_{1}\mid0,1000\right) \\ &\times\operatorname{uniform}\left(\sigma\mid0,100\right) \\ \boldsymbol{g}\left(\beta_{0},\beta_{1},x_{i}\right) &= & \beta_{0}+\beta_{1}\left(x_{i}-\bar{x}\right) \end{split}
```

```
b0 ~ dnorm(0, .001)
b1 ~ dnorm(0, .001)
sigma ~ dunif(0, 100)
tau <- 1/sigma^2
xBar <- mean(x)
for (i in 1:length(y)){
   mu[i] <- b0 + b1 * (x[i] - xBar)
   y[i] ~ dnorm(mu[i], tau)
}
b0_UC <- b0 - b1 * xBar</pre>
```

Recovering uncentered parameters



$$B_0 = \beta_0 - \beta_1 \bar{x}$$

$$B_1 = \beta_1$$

- For this to work properly, all of the coefficients in the model must be added.
- Slopes will not be the same if there is an interaction term or quadratic. In these cases, back transforming is not simple.

Standardizing predictor data

$$y_i = \beta_0 + \beta_1 \left(\frac{x_i - \bar{x}}{\sigma_x} \right)$$

Why complicate things...?

- To reduce autocorrelation in MCMC chain and speed convergence.
- To make the intercept more easily interpretable.
- To make parameters more easily comparable.

Standardizing predictor data

```
[\beta_0, \beta_1, \sigma \mid \mathbf{y}] \propto \prod_{i=1}^{\infty} \operatorname{normal}(y_i \mid g(\beta_0, \beta_1, x_i, \bar{x}, \sigma_x), \sigma^2) \\ \times \operatorname{normal}(\beta_0 \mid 0, 1000) \operatorname{normal}(\beta_1 \mid 0, 1000) \\ \times \operatorname{uniform}(\sigma \mid 0, 100) \\ g(\beta_0, \beta_1, x_i) = \beta_0 + \beta_1 \left(\frac{x_i - \bar{x}}{\sigma_x}\right) \\ b0 \sim \operatorname{dnorm}(0, .001) \\ b1 \sim \operatorname{dnorm}(0, .001) \\ \operatorname{sigma} \sim \operatorname{dunif}(0, 100)
```

tau <- 1/sigma^2
xBar <- mean(x)
xSD <- sd(x)</pre>

for (i in 1:length(y)){

y[i] ~ dnorm(mu[i], tau)

 $mu[i] \leftarrow b0 + b1 * ((x[i] - xBar)/xSD$

Recovering unstandardized parameters

$$B_0 = \beta_0 - \frac{\beta_1 \bar{x}}{\sigma_x}$$

$$B_1 = \frac{\beta_1}{\sigma_x}$$

- This only works if there are not squared values or interactions.
- Generally, I back-transform predictions not parameters. (How?)