

①

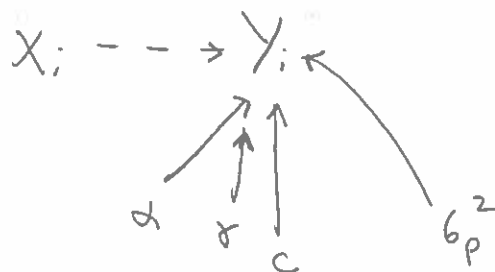
# Simple Bayes

$y_1, x_1$	$y_4, x_4$
$y_2, x_2$	etc
$y_3, x_3$	

$$g(\alpha, \gamma, c, x_i) = \frac{\alpha (x_i - c)}{\frac{\alpha}{\gamma} + (x_i - c)}$$

height increment

$x_i$ 's  
are measured  
perfectly



Interpretation of  
 $b^2$ : process variance  
because  $y_i$  are  
measured perfectly

$$[\alpha, \gamma, c, b^2 | \underline{y}] \propto \prod_{i=1}^n [y_i | g(\alpha, \gamma, c, x_i), b^2] [\alpha] [\gamma] [c] [b_p^2]$$

← put these here

$$y_i \sim \text{normal}(g(\alpha, \gamma, c, x_i), b^2)$$

$$\alpha \sim \text{gamma}\left(\frac{35^2}{4.25^2}, \frac{35}{4.25^2}\right)$$

$$\gamma \sim \text{Uniform}(0, 10)$$

$$b_p^2 \sim \text{uniform}(0, 50) \quad \leftarrow \text{mention inverse gamma}$$

talk about alternative notations for likelihood?

$$[y_i | \mu_i, b^2]$$

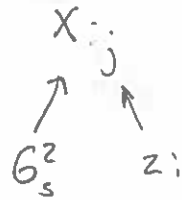
$$\mu_i = \alpha$$

~~$$[y_i | \theta]$$~~

$$\theta = (\gamma, \alpha, c, b^2)$$

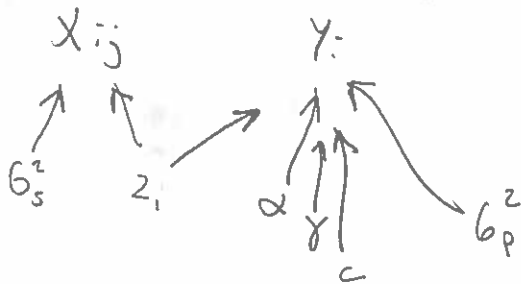
$$[y | \alpha, \gamma, c, b^2]$$

- ② ~~the~~ - Errors in the  $x$ 's. We have  $j=1, \dots, 8$  replicate measures of light each tree
- How would we estimate the mean light seen by the tree?



where  $z_i$  is the mean

We now attach this to our model



Talk about  $G_s^2$  at individual tree level

$$[G_s^2, \underline{z}, \alpha, \gamma, c, G_p^2 \mid \underset{\substack{\uparrow \\ \text{vector}}}{\underline{x}}, \underline{y}] \propto \prod_{i=1}^n \prod_{j=1}^8 [y_i \mid g(\alpha, \gamma, c, x_{ij}), z_i] [x_{ij} \mid z_i, G_s^2]$$

matrix

note!  $z_i$  has

$$\times [z_i] [G_s^2] [\alpha] [\gamma] [c] [G_p^2]$$

Same priors as before

except we now add

$$z_i \sim \text{uniform}(0, 100)$$

$$G_s^2 \sim \text{uniform}(0, 100)$$

① Ask about spread  $G_s^2$

② Show alternative priors

③ Show no replication

3

Errors in the  $Y_i$

Calibration equation — model the tree mass  $M = ah^b$

Single plot

$\begin{bmatrix} x, y & x, y & x, y \\ x, y & \text{etc} \end{bmatrix}$

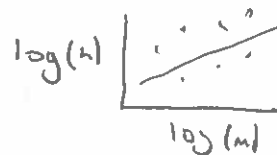
Mass =  $m$   
Height =  $h$

$$m = ah^b$$

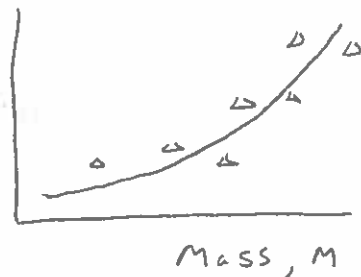
$$h = \left(\frac{m}{a}\right)^{\frac{1}{b}}$$

the tree from its current height not height inc.

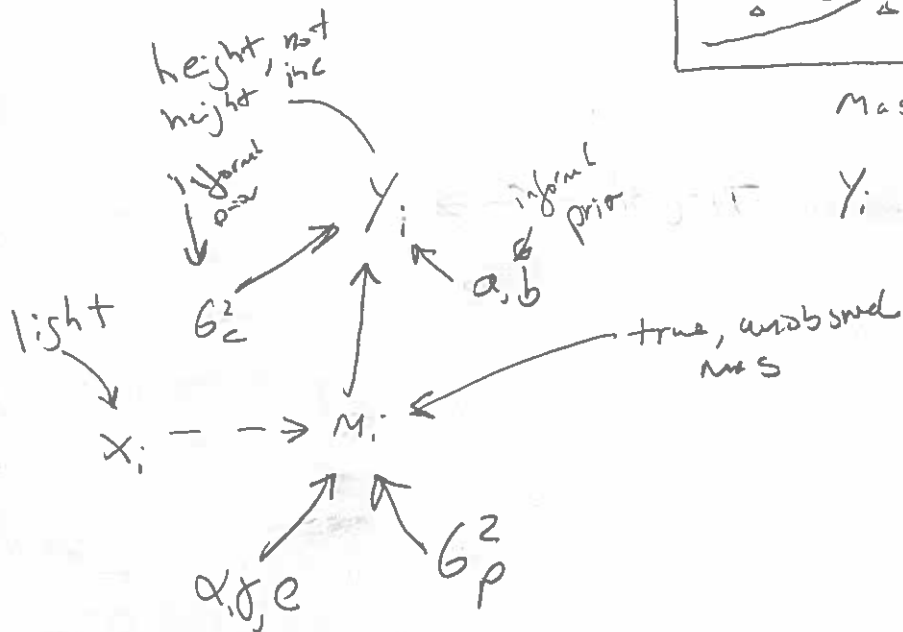
$\sigma_c^2$  is scatter



height,  $h$



$Y_i$  = height of  $i$ th tree



$$[m, a, b, \sigma_c^2, \alpha, \delta, \epsilon, \sigma_p^2 | Y] = \prod_{i=1}^n [Y_i | (M_i/a)^{\frac{1}{b}}, \sigma_c^2] [M_i | g(\alpha, \delta, \epsilon, X_i), \sigma_p^2]$$

$$\times [a] [b] [\sigma_c^2] [\alpha] [\delta] [\epsilon] [\sigma_p^2]$$

choices of distributions

$$\log(Y_i) \sim \text{normal}(\log((M_i/a)^{\frac{1}{b}}), \sigma_c^2)$$

$$Y_i \sim \text{lognormal}(\log((M_i/a)^{\frac{1}{b}}), \sigma_c^2)$$

$$Y_i \sim \text{gamma}\left(\frac{(M_i/a)^{\frac{2}{b}}}{\sigma_c^2}, \frac{(M_i/a)^{\frac{1}{b}}}{\sigma_c^2}\right)$$

Be careful here!

on log scale will not work! unless regression is done on log scale

what could you use to model mass?

④

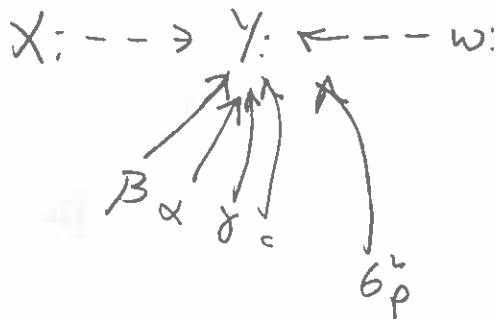
Differences due to treatment

$$g(\alpha, \delta, c, \beta) = \frac{\alpha(X_i - c)}{\frac{\alpha}{\gamma} + (X_i - c)} + \beta w_i$$

treatment for  $i$ th tree

$w_i = 0$  if control

$w_i = 1$  if treated

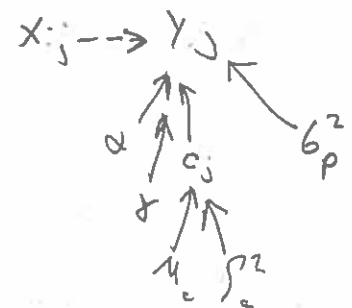
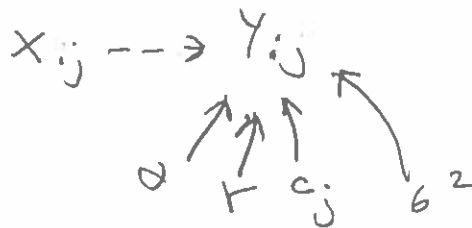


Differences <sup>in shade tolerance</sup> between species

$i$  = individual  
 $j$  = species

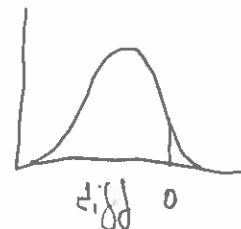
< 4 species

> 4 species



posterior distribution of difference

$$\text{diff} = c_1 - c_2 \quad [\text{diff}]$$



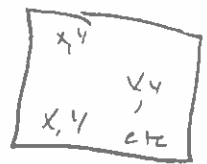
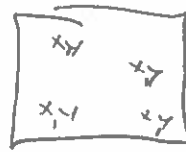
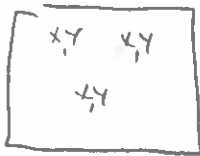
5

Mult-level model including effect of locations

$j = 1, \dots, S$  sites

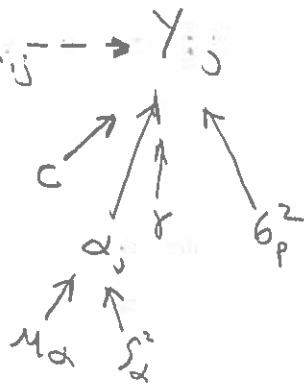
4 knowledge differences...

Put on board

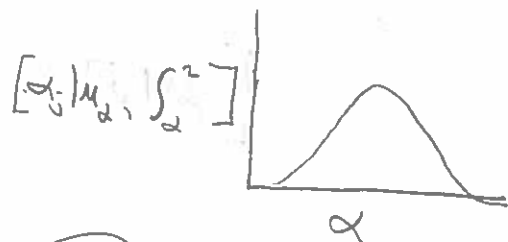


What are the options for modeling  $\alpha_j$ ?

- ① pooled
- ② individual



ith plant at location plot j



Talk about borrowing strength

model

note vector

$$[\alpha_j, \gamma, c, \mu_\alpha, \sigma_\alpha^2, b_p^2 | \underline{y}] \propto \prod_{i=1}^{n_j} \prod_{j=1}^S [y_{ij} | g(\alpha_j, \gamma, c, x_{ij}), b_p^2] [\alpha_j | \mu_\alpha, \sigma_\alpha^2] \times [\gamma] [c] [b_p^2] [\sigma_\alpha^2] [\mu_\alpha]$$

Note no prior on  $\alpha_j$

$$y_{ij} \sim \text{normal}(g(\alpha_j, \gamma, c, x_{ij}), b_p^2)$$

$$\gamma \sim \text{unif}(0, 10)$$

$$b_p \sim \text{unif}(0, 50)$$

$$\sigma \sim \text{unif}(0, 50)$$

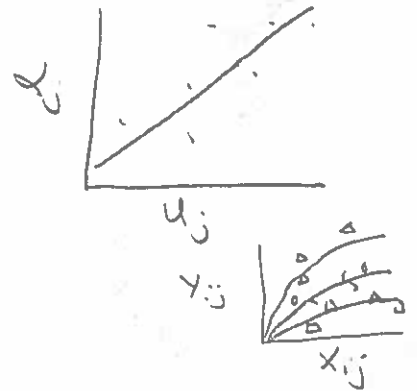
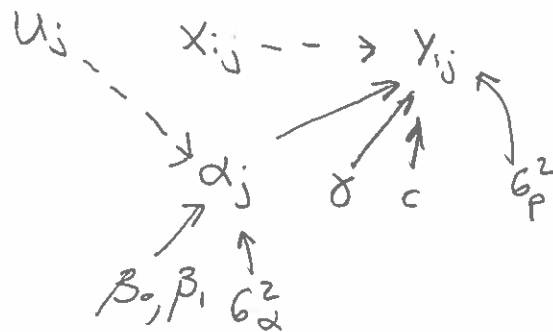
$$\mu_\alpha \sim \text{gamma}(.001, .001) \text{ or } \mu_\alpha \sim \text{gamma}\left(\frac{35^2}{4.25^2}, \frac{35}{4.25^2}\right)$$

Ask about  $b_p^2$ , how to model at plot scale

could make the broader

6

Multi-level with location data, covariates for each plot, say soil ~~water~~ water — add  $u_j$  to diagram to represent site data, assume normal perfectly

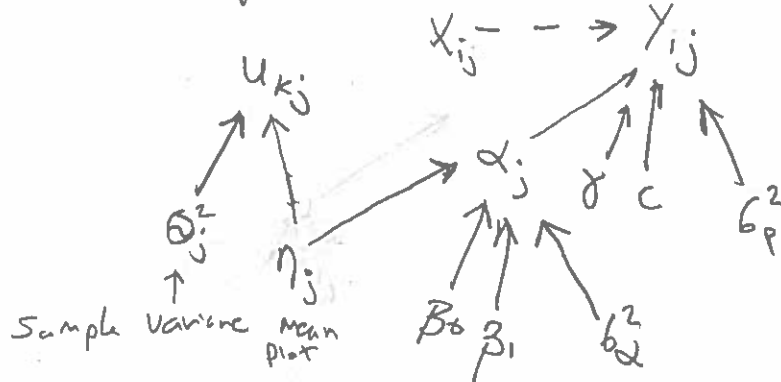


$$[\alpha, \gamma, c, \beta_0, \beta_1, \epsilon^2, \epsilon^2_p | \underline{y}] \propto \prod_{i=1}^{n_j} \prod_{j=1}^J [y_{ij} | g(\alpha_j, \gamma, c) \epsilon^2] [\alpha_j | \beta_0 + \beta_1 u_j, \epsilon^2_\alpha]$$

$$X [\gamma] [c] [\epsilon^2_p] [\beta_0] [\beta_1] [\epsilon^2_\alpha]$$

↑ ↑  
normal priors

What if we have replicate measures of soil water? (samples, error in covariate)



Talk about spatial interpolation

not  $x_{ij}$  on  $y_{ij}$

$$[\eta, \epsilon^2, \beta_0, \beta_1, \alpha, \gamma, c, \epsilon^2_p | \underline{u}, \underline{y}] \propto \prod_{i=1}^{n_j} \prod_{j=1}^J \prod_{k=1}^{10} [y_{ij} | g(\alpha_j, \gamma, c, x_{ij}) \epsilon^2_p] [\alpha_j | \beta_0 + \beta_1 \eta_j, \epsilon^2_\alpha]$$

$$[\epsilon^2_\eta] [\eta_j] [\beta_0] [\beta_1] [\epsilon^2_\alpha] [\gamma] [c] [\epsilon^2_p]$$