

①

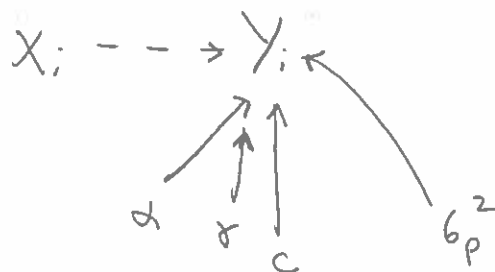
Simple Bayes

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?

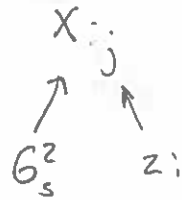
$$[y_i | \mu_i, b^2] \quad \mu_i = \alpha$$

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

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

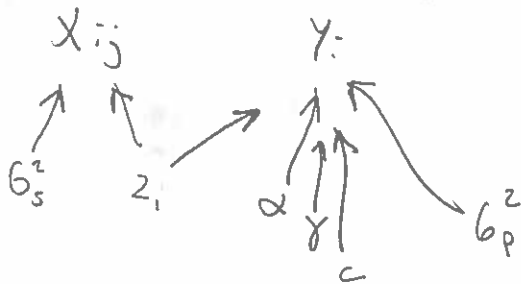
$$[y | \alpha, \gamma, c, b^2]$$

- ② ~~the same~~ - 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

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

note! z_i has \downarrow

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

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Errors in the Y_i

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

Single plot

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

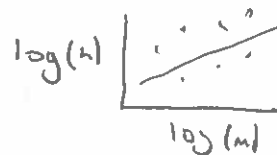
Mass = m
Height = h

$$m = ah^b$$

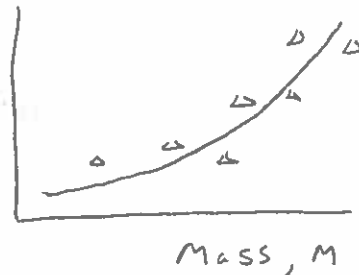
$$h = \left(\frac{m}{a}\right)^{\frac{1}{b}}$$

the tree from its current height not height inc.

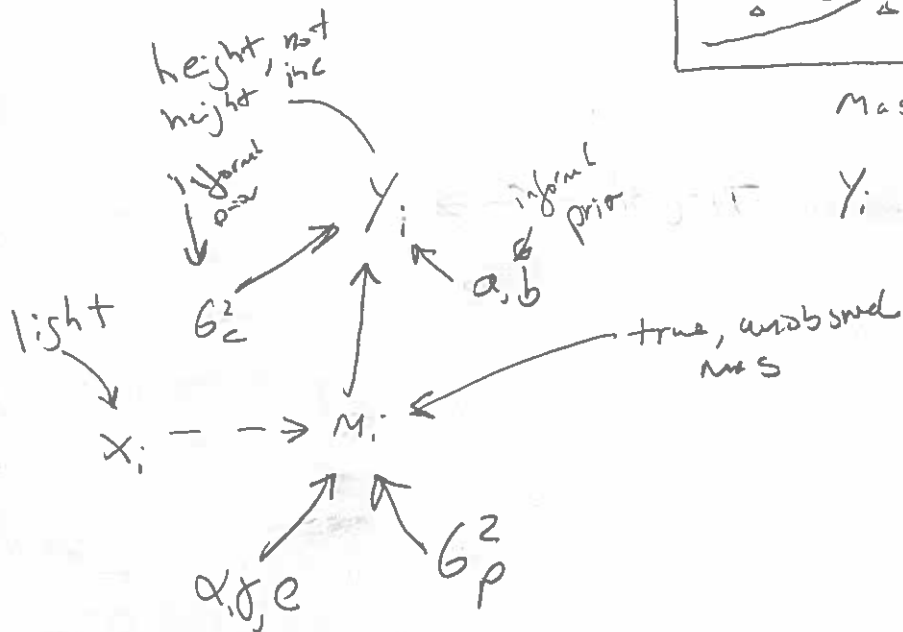
σ_c^2 is scatter



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

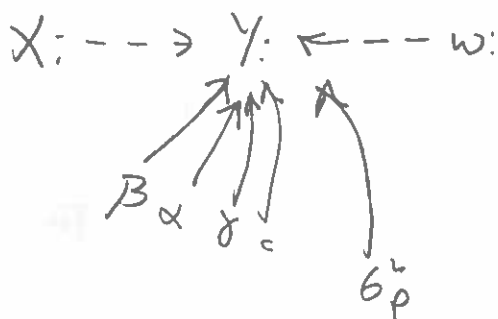
Differences due to treatment

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

treatment for i th tree

$w_i = 0$ if control

$w_i = 1$ if treated

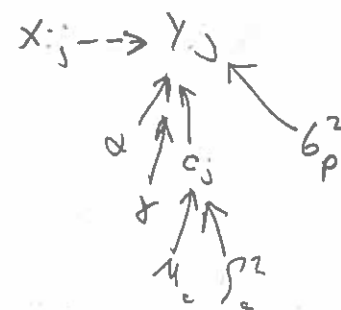
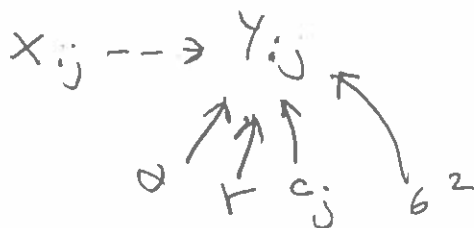


Differences ^{in shade tolerance} between species

i = individual
 j = species

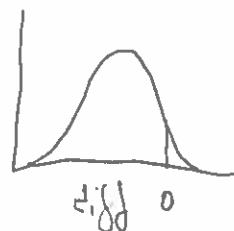
< 4 species

> 4 species



posterior distribution of difference

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



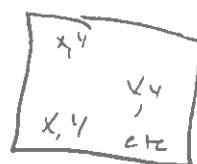
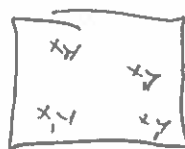
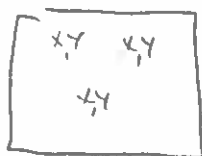
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Mult-level model including effect of locations

$j = 1, \dots, S$ sites

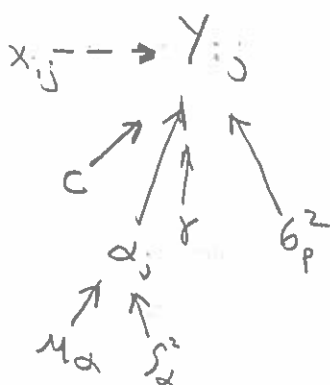
4 knowledge differences...

Put on board

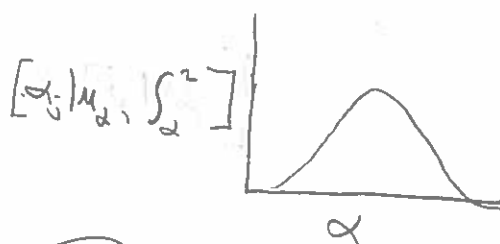


What are the options for modeling α_j ?

- ① pooled
- ② individual



ith plant at location plot j



Talk about borrowing strength

model

note vector

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

Note no prior on α_j

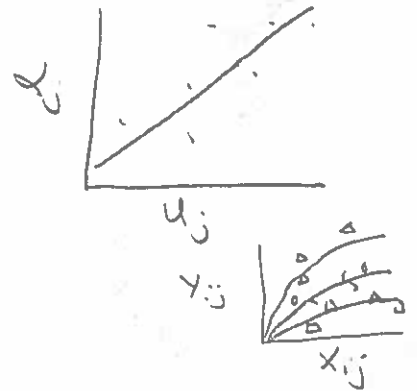
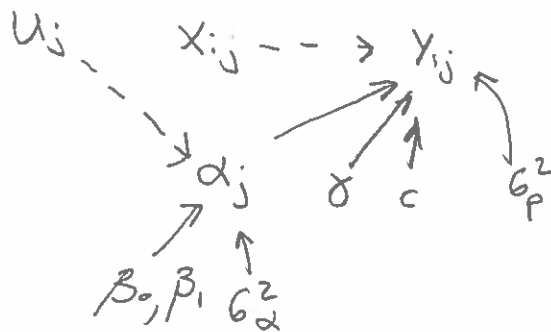
$$\begin{aligned} y_{ij} &\sim \text{normal}(g(\alpha_j, r, c, x_{ij}), b^2_p) \\ r &\sim \text{unif}(0, 10) \\ b_p &\sim \text{unif}(0, 50) \\ S &\sim \text{unif}(0, 50) \\ \mu_\alpha &\sim \text{gamma}(.001, .001) \text{ or } \mu_\alpha \sim \text{gamma}\left(\frac{35^2}{4.25^2}, \frac{35}{4.25^2}\right) \end{aligned}$$

Ask about b^2_p , how to model at plot scale

could make the broader

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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

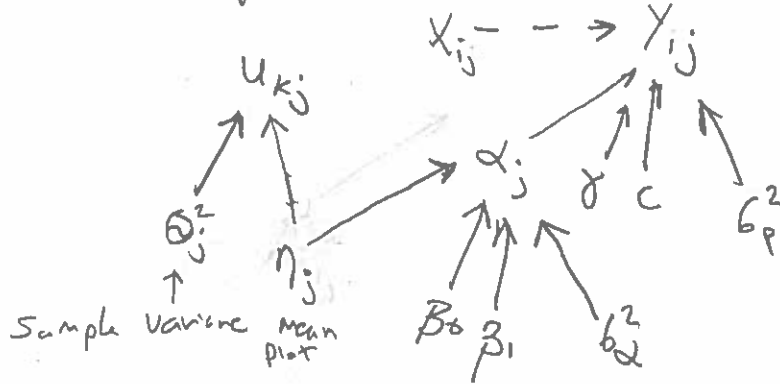


$$[\alpha, \gamma, c, \beta_0, \beta_1, \epsilon_\alpha^2, \epsilon_p^2 | \underline{y}] \propto \prod_{i=1}^{n_j} \prod_{j=1}^J [y_{ij} | g(\alpha_j, \gamma, c) \epsilon_p^2] [\alpha_j | \beta_0 + \beta_1 u_j, \epsilon_\alpha^2]$$

$$X [\gamma] [c] [\epsilon_p^2] [\beta_0] [\beta_1] [\epsilon_\alpha^2]$$

↑ ↑
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}

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

$$[\theta_j^2] [\eta_j] [\beta_0] [\beta_1] [\epsilon_\alpha^2] [\gamma] [c] [\epsilon_p^2]$$