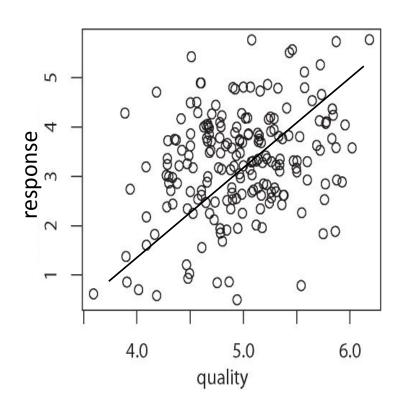
Characterizing Uncertainty

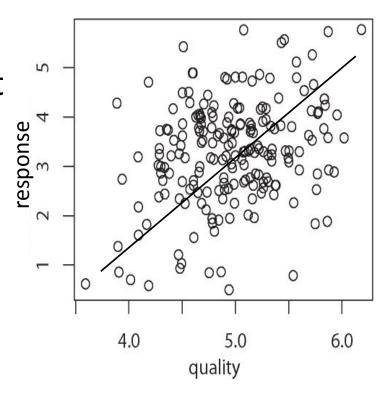
$$y_i \sim \beta_0 + \beta(x_i) + \epsilon_i$$



Classic Assumptions

- Error in Y is measurement error
- Normally distributed error
- Observations are independent

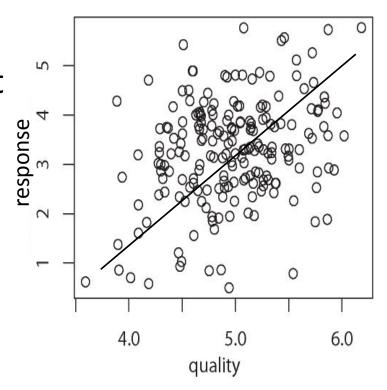
$$y_i \sim \beta_0 + \beta(x_i) + \epsilon_i$$



Classic Assumptions

- Error in Y is measurement error
- Normally distributed error
- Observations are independent
- Homoskedasticity

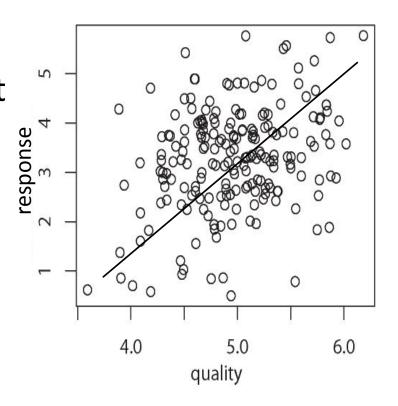
$$y_i \sim \beta_0 + \beta(x_i) + \epsilon_i$$

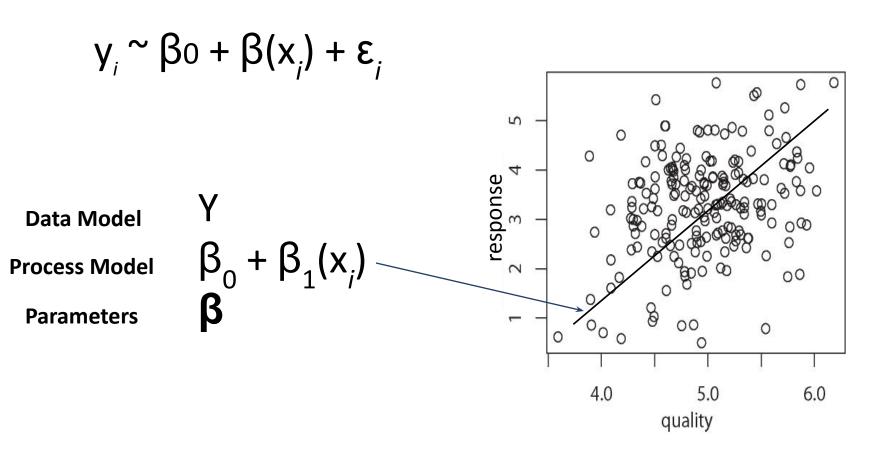


Classic Assumptions

- Error in Y is measurement error
- Normally distributed error
- Observations are independent
- Homoskedasticity
- No error in X variables

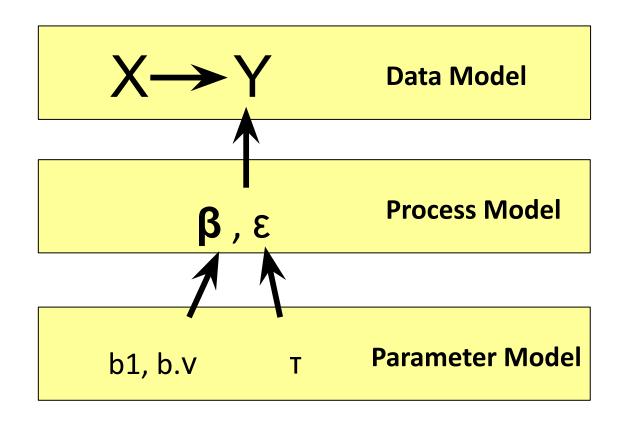
$$y_i \sim \beta_0 + \beta(x_i) + \varepsilon_i$$

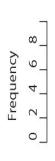


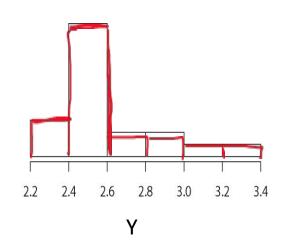


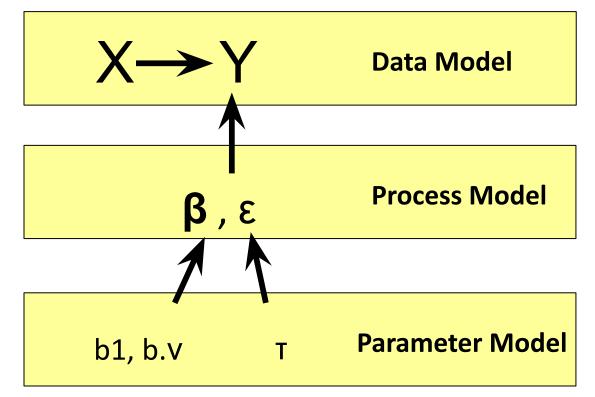
Linear Model – Graph Notation

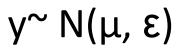
$$y_i \sim \beta_0 + \beta(x_i) + \epsilon_i$$

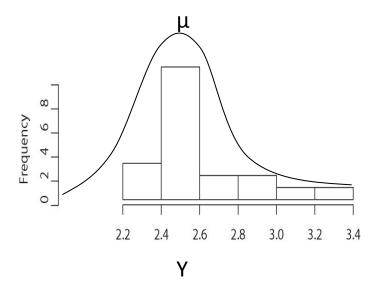




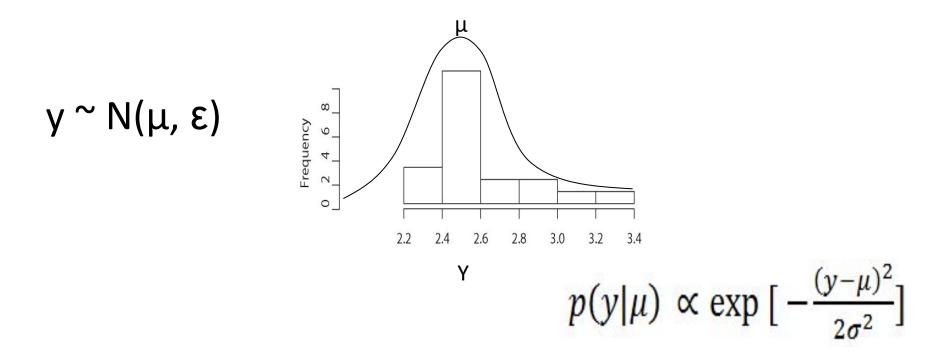


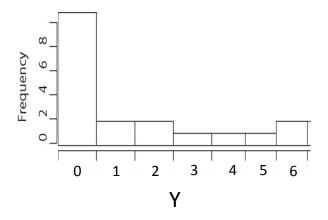


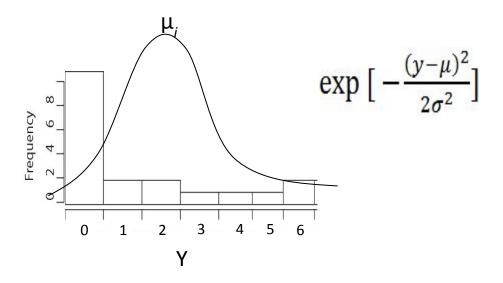


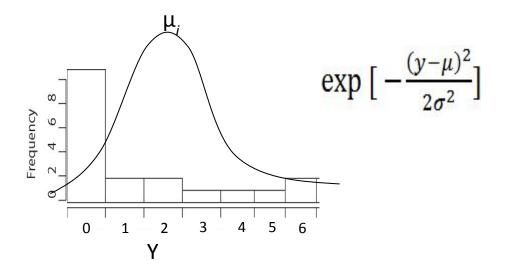


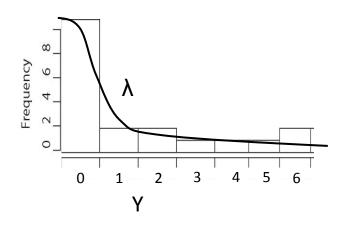
Expected relationship between data samples and the range of all possible data is described by a probability distribution.



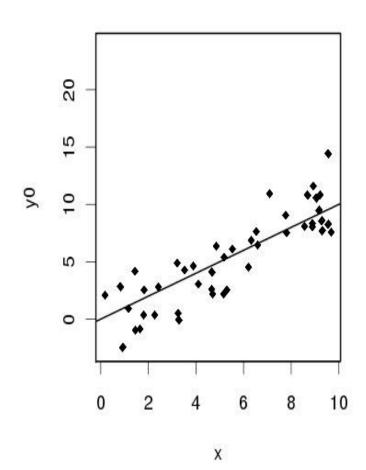


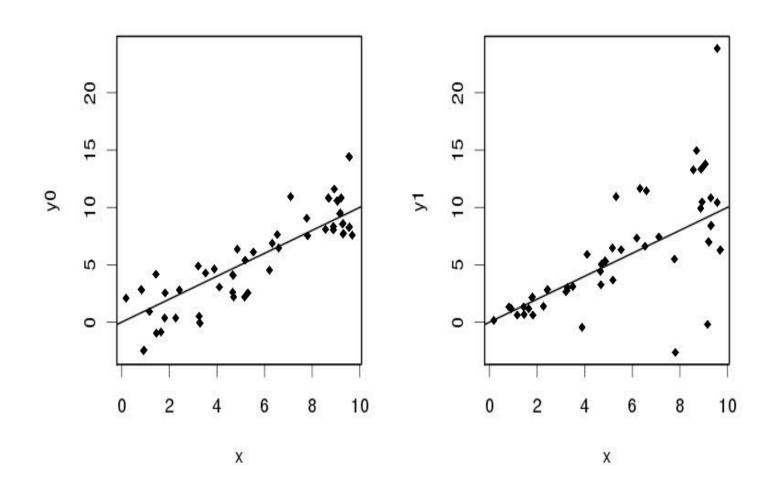


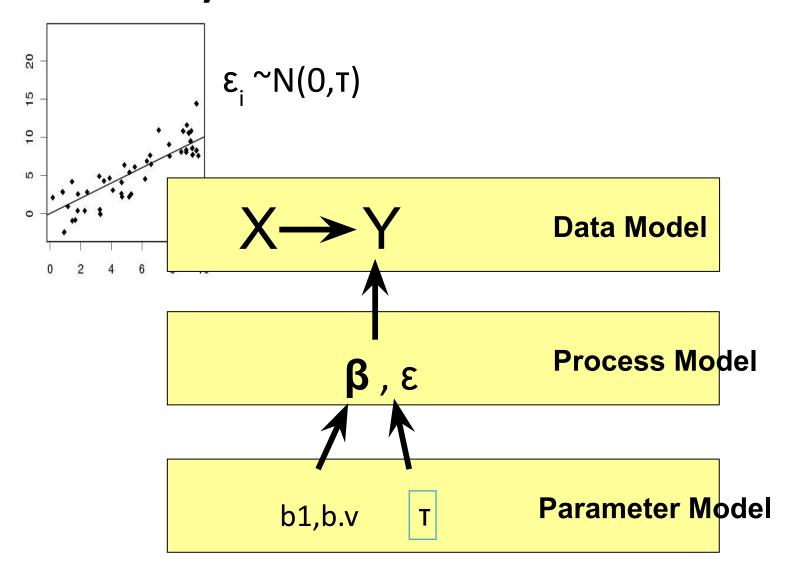




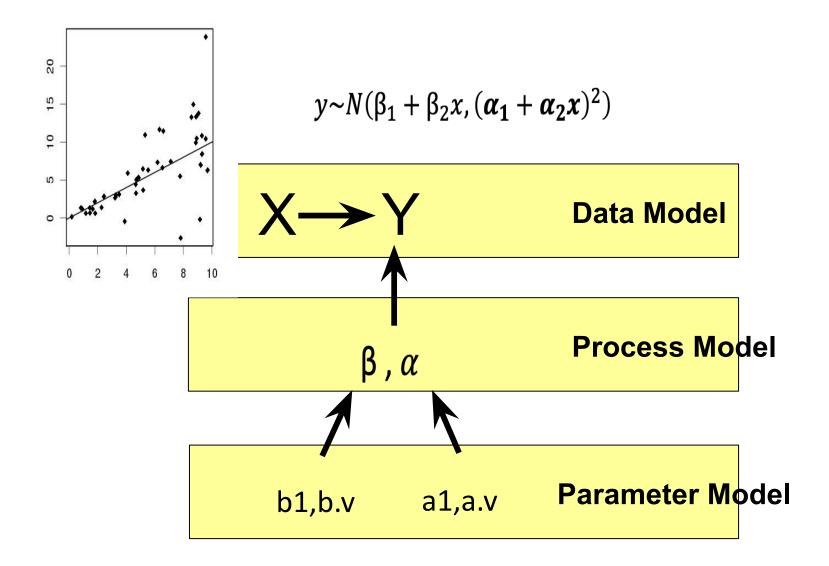
$$\frac{\lambda_{,}}{v!}^{y}e^{-\lambda}$$







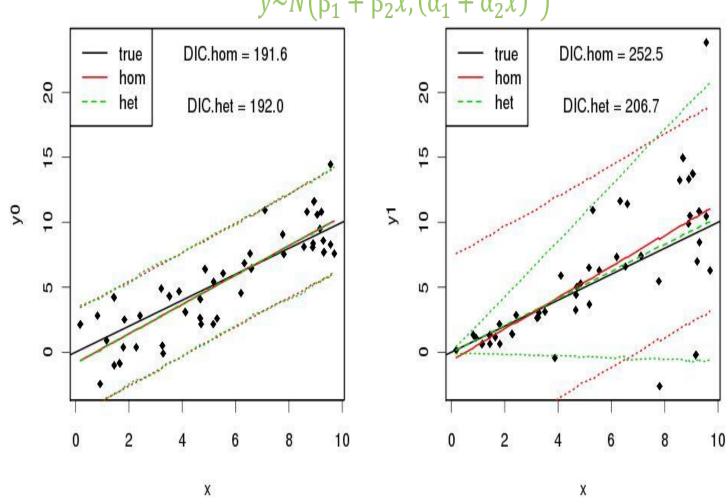
Uncertainty & Variance: Heteroskedasticity

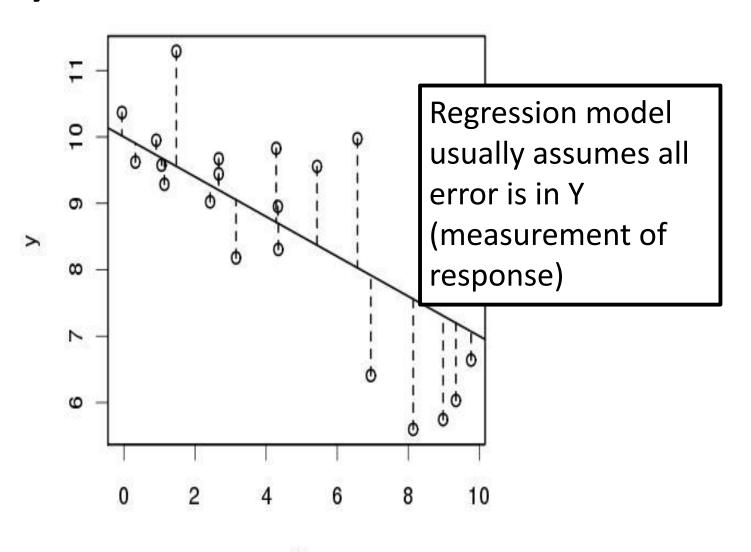


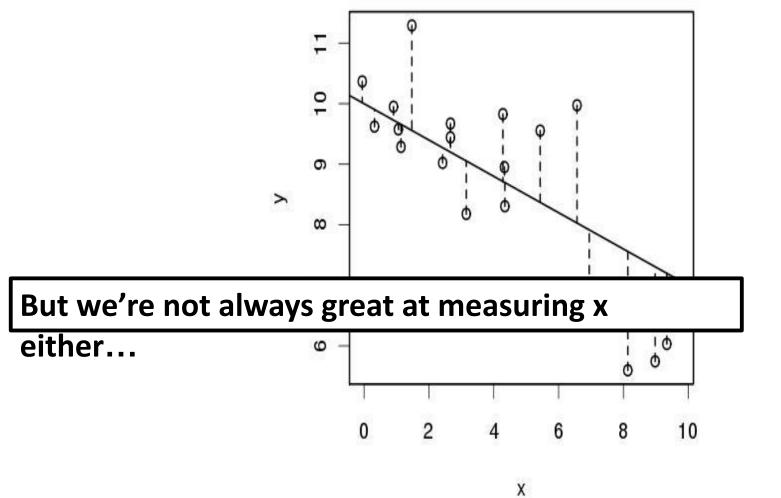
Uncertainty & Variance: Heteroskedasticity

$$y \sim N(\beta_1 + \beta_2 x, s^2)$$

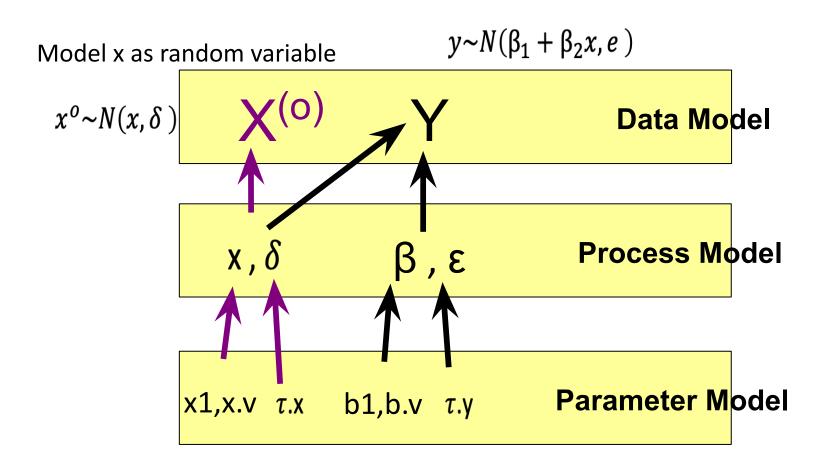
$$y \sim N(\beta_1 + \beta_2 x, (\alpha_1 + \alpha_2 x)^2)$$

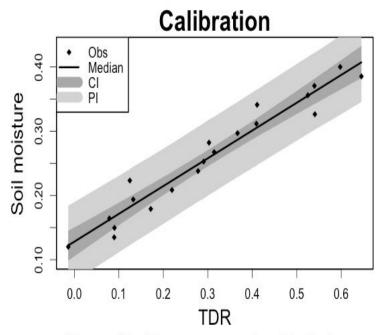




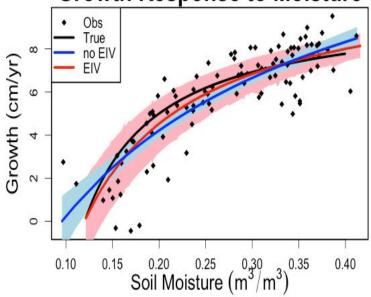


Uncertainty & Variance: Errors in Variables



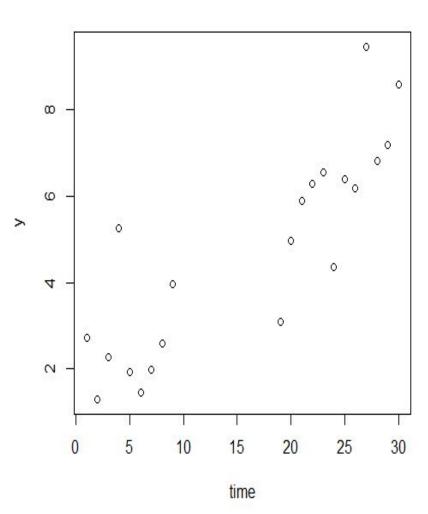




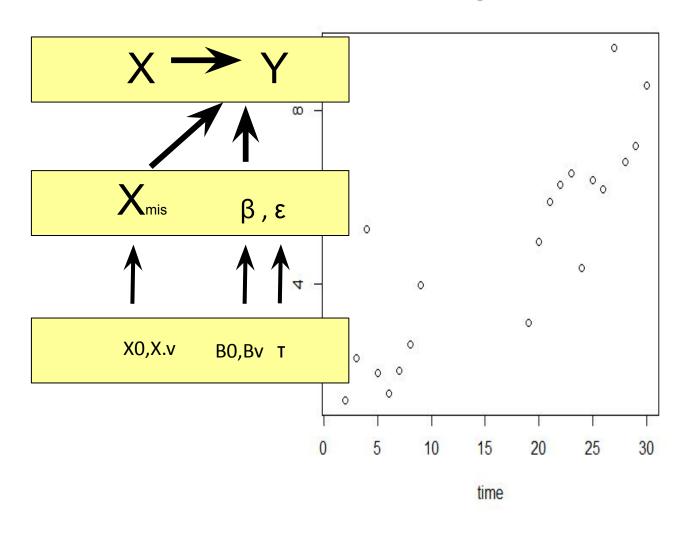


- State not directly observed
 - -Missing data

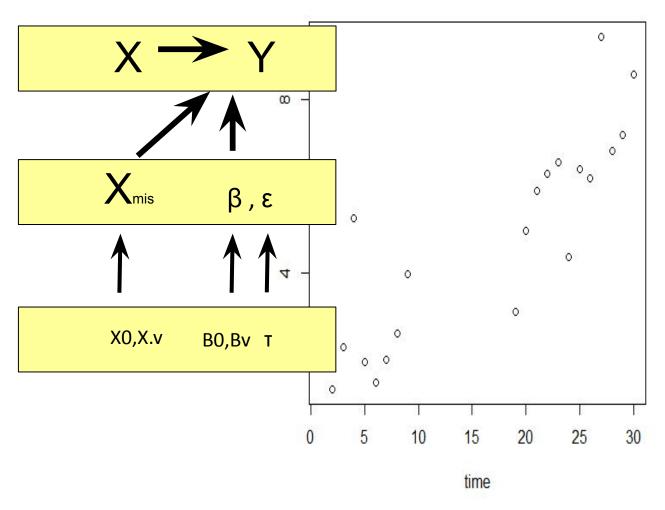
Missing Data



Missing Data

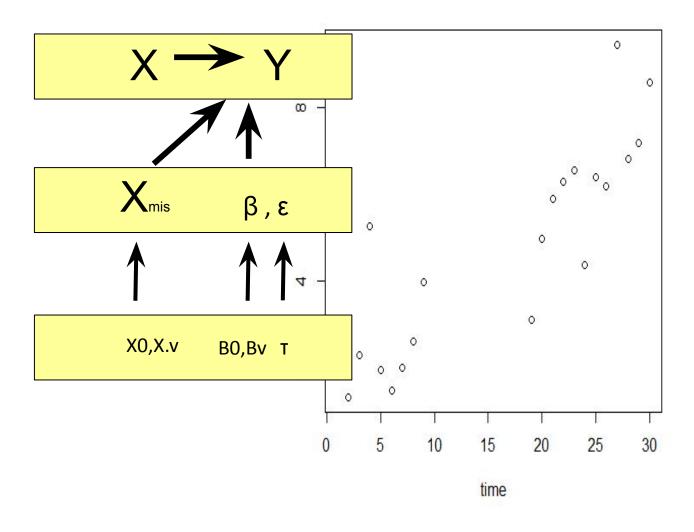


Missing Data



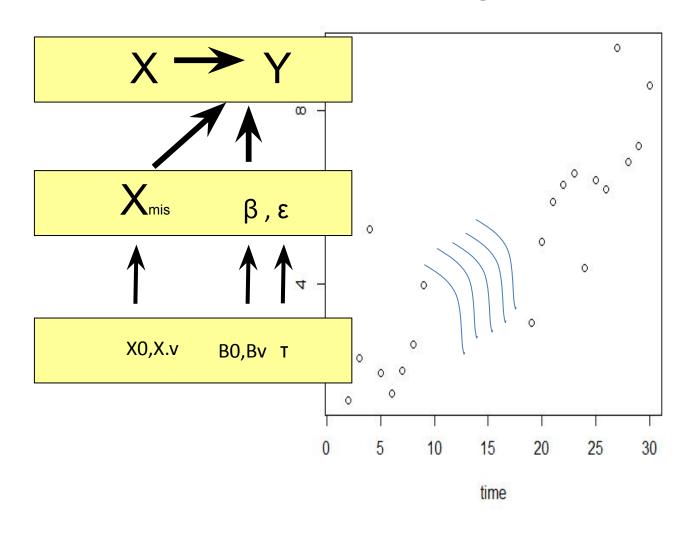
- Update the regression model conditioned on the current values of the missing data
- •Update the missing data based on the current regression model and covariate values

Missing Data



data is <u>missing at</u> <u>random</u>

Missing Data



data is <u>missing at</u> <u>random</u>

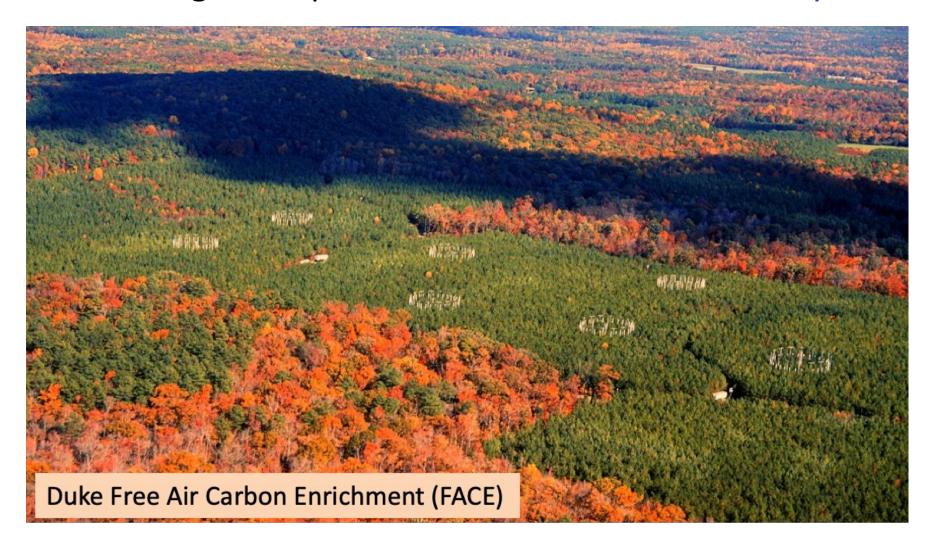
- State not directly observed
 - -Missing data
 - –Proxy measures

- State not directly observed
 - –Missing data
 - –Proxy measures

Will rising atmospheric CO2 increase tree fecundity?

Ignoring variable latency can lead to incorrect or falsely overconfident conclusions (and wonky bad forecasts)

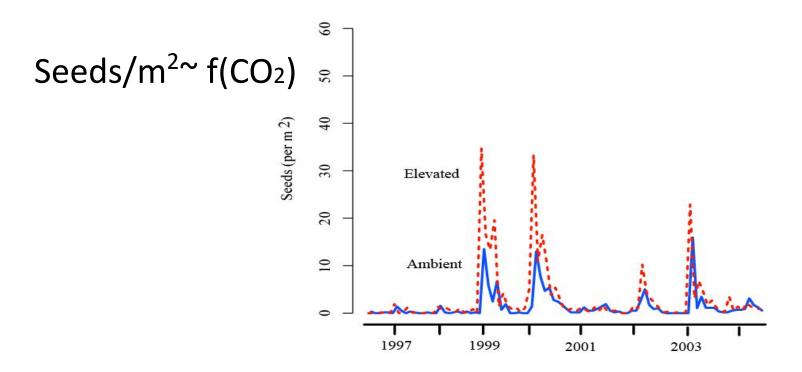
Will rising atmospheric CO₂ increase tree fecundity?



Will rising atmospheric CO₂ increase tree fecundity?

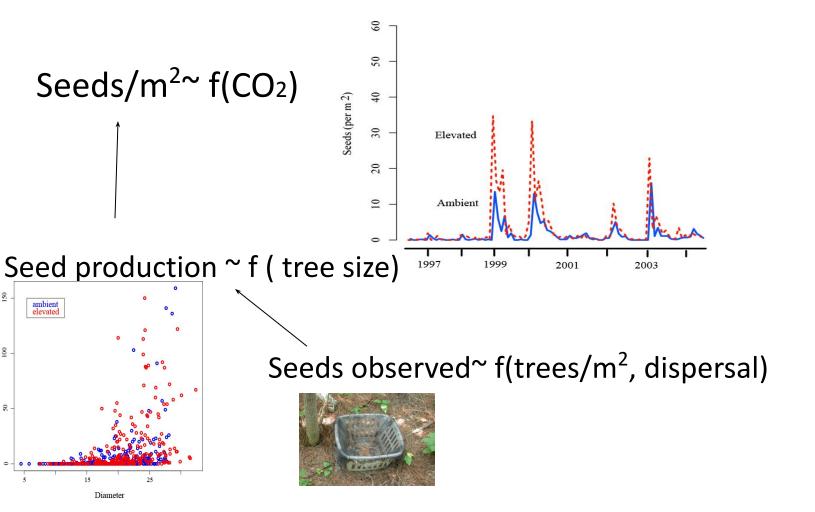
Fecundity ~ f(CO₂)

Will rising atmospheric CO₂ increase tree fecundity?

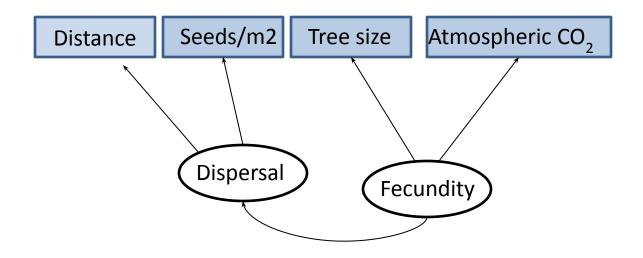


Cones (1998-2003)

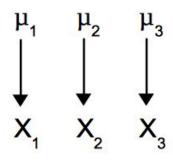
Will rising atmospheric CO₂ increase tree fecundity?



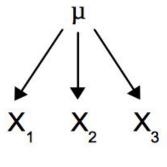
Will rising atmospheric CO₂ increase tree fecundity?



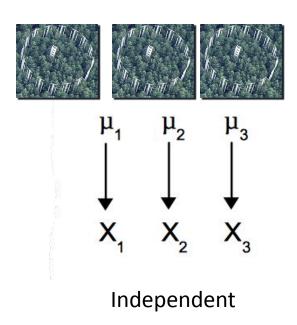
Fecundity \sim f(CO₂ + tree size) & f(Obs)

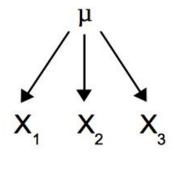


Independent

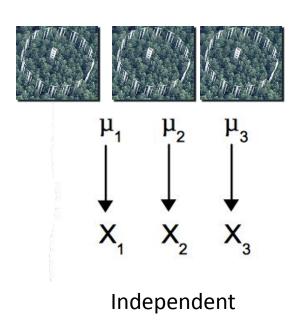


Shared

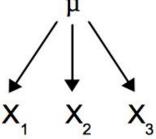




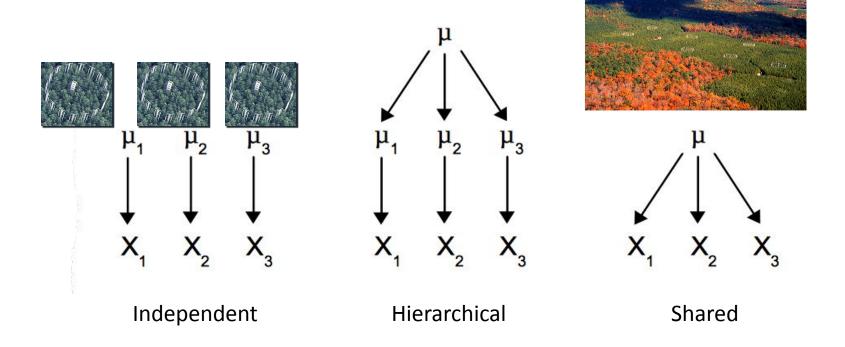
Shared



μ



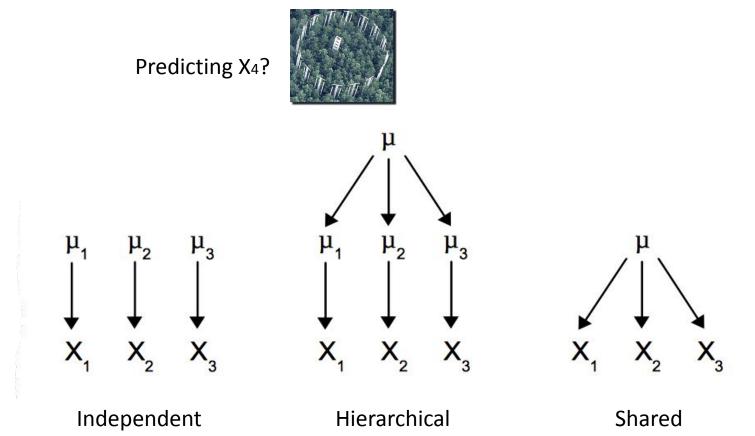
Shared



Model variability in the parameters

Partition variability explicitly into multiple terms

Borrow strength across data sets



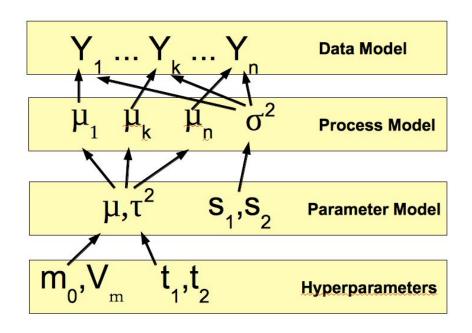
Model variability in the parameters

Partition variability explicitly into multiple terms

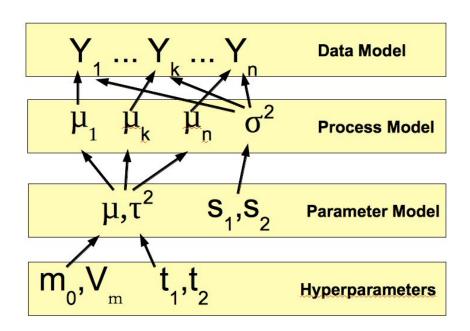
Borrow strength across data sets

Hierarchical with respect to parameters

Pay attention to subscripts!



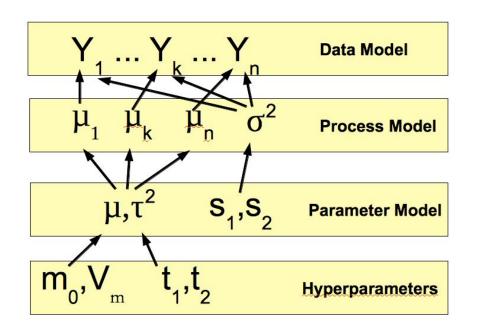
$$Y_k^{\sim} N(u_{k'}\sigma^2)$$



$$Y_k \sim N(u_k, \sigma^2)$$

$$u_k^{\sim} N(u,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$



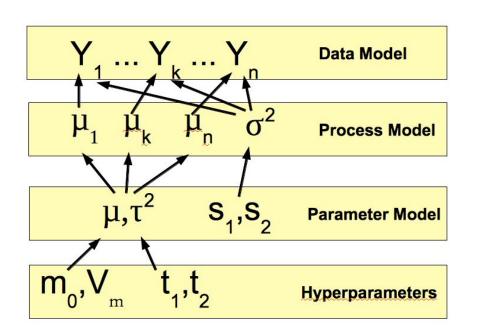
$$Y_k^{\sim} N(u_k, \sigma^2)$$

$$u_k^{\sim} N(u,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$

$$u^{\sim} N(u0,v_m)$$

 $T^2 \sim IG(t1,t2)$



$$Y_k \sim N(u_k, \sigma^2)$$

$$Y_k^{\sim} N(u_g + \alpha_k, \sigma^2)$$

$$u_k^{\sim} N(u,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$

$$u^{N}(u0,v_{m})$$

 $T^{2} \sim IG(t1,t2)$

Random effects attempt to account for the unexplained variance associated with some group (plot, year, etc.) due to all the things that were not measured

May point to scales that need additional explanation

Adding covariates may explain some portion of this variance, but there's always something you didn't measure

$$Y_k \sim N(u_k, \sigma^2)$$

$$Y_k^{\sim} N(u_g + \alpha_k, \sigma^2)$$

$$u_k^{\sim} N(u,T^2)$$

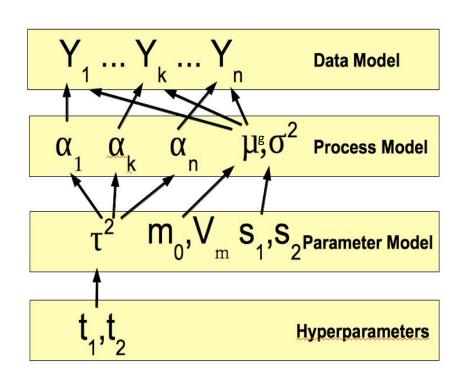
 $\sigma^2 \sim IG(s_1,s_2)$

$$\alpha_k \sim N(0,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$

$$u_g^{\sim} N(u0,v_m)$$

 $T^2 \sim IG(t1,t2)$



$$Y_k^{\sim} N(u_g + \alpha_k, \sigma^2)$$

$$\alpha_k \sim N(0,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$

$$u_g^{\sim} N(u0,v_m)$$

 $T^2 \sim IG(t1,t2)$

$$Y_k \sim N(u_g + \alpha_k, \sigma^2)$$

REs have mean 0

$$\alpha_k \sim N(0,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$

$$u_g^{\sim} N(u0,v_m)$$

 $T^2 \sim IG(t1,t2)$

$$Y_k^{\sim} N(u_g + \alpha_k, \sigma^2)$$

REs have mean 0

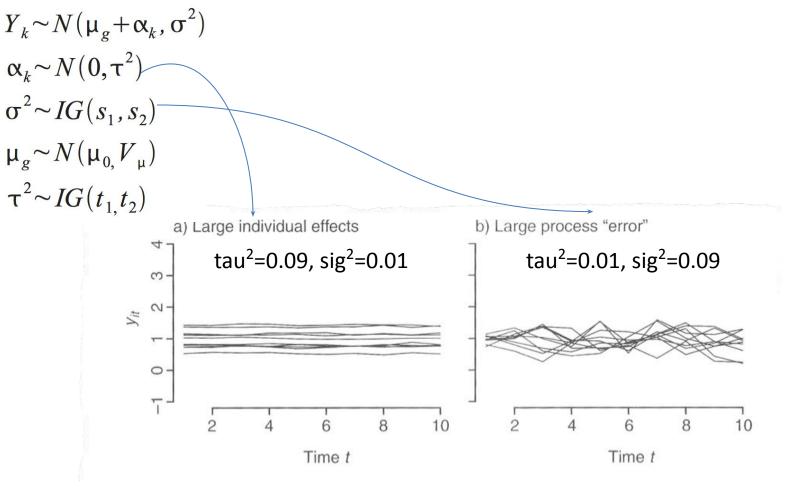
$$\alpha_k \sim N(0,T^2)$$

 $\sigma^2 \sim IG(s_1,s_2)$

REs variance attributes portion of uncertainty to specific source

$$u_g^{\sim} N(u0,v_m)$$

 $T^2 \sim IG(t1,t2)$



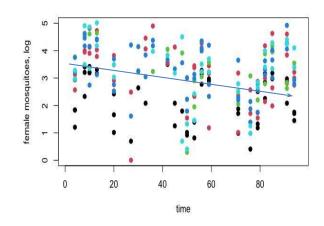
RE's traditionally aspects of the study that would likely change if experiment/observation protocol was replicated

- e.g., Plots, Years
- used to account for lack of independence among reps

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Replication is important for identifiability (i.e., partitioning variance between process error and REs)

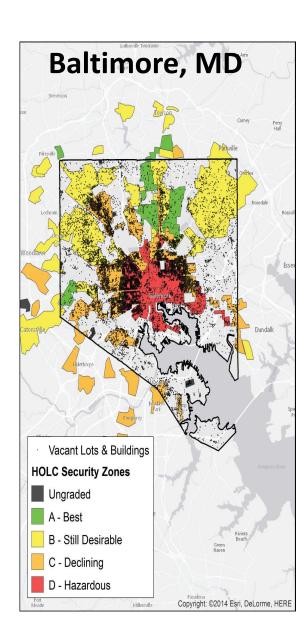


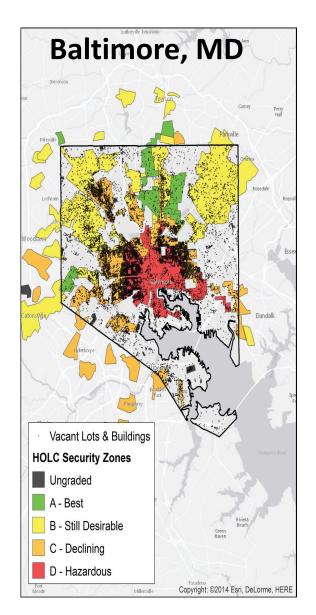
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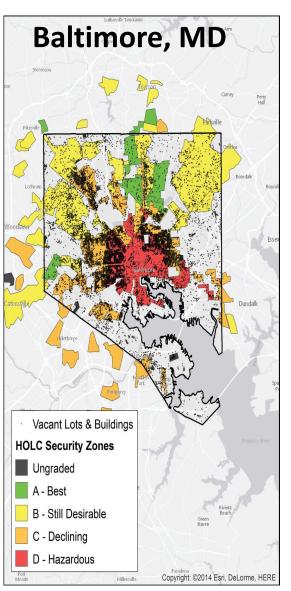
Treatments/covariates of interest are treated as fixed effects (mixed effects model)





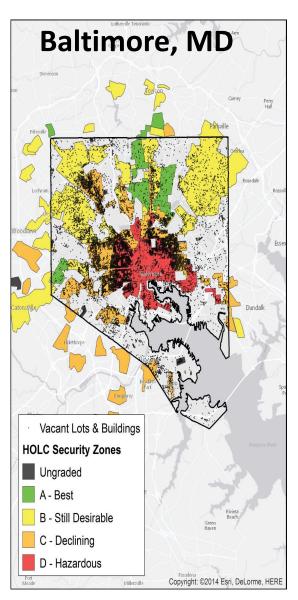




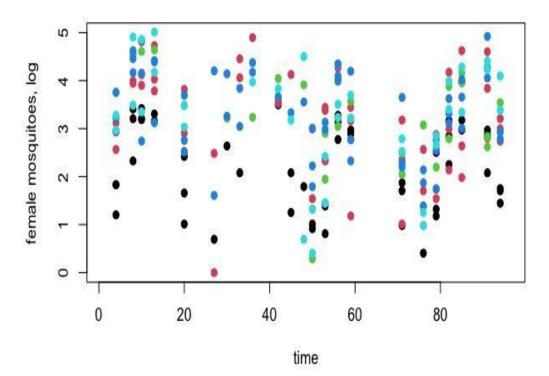












```
Global Mean= "
model\{
mu \sim dnorm(0,0.001)
tau \sim dgamma(0.001,0.001)

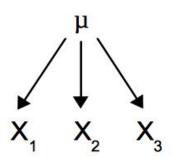
for(i in 1:nblocks){
x_1 \quad x_2
for(t in 1:ndates){
X[i,t] \sim dnorm(mu, tau)
}}
```

```
Global Mean= "
model{
 mu ~ dnorm(0,0.001)
tau ~ dgamma(0.001,0.001)
                                     2
for(i in 1:nblocks){
                                 female mosquitoes, log
 for(t in 1:ndates){
  X[i,t] ~ dnorm(mu, tau)
 }}}
                                    2
                                     0
                                                  20
                                                           40
                                                                    60
                                                                              80
```

time

```
Global Mean= "
model{
mu ~ dnorm(0,0.001)
tau ~ dgamma(0.001,0.001)

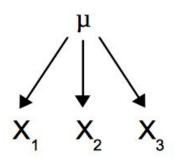
for(i in 1:nblocks){
  for(t in 1:ndates){
    X[i,t] ~ dnorm(mu, tau)
}}}
```



```
Space= "
model{
 mu \sim dnorm(0,0.001)
 tau \sim dgamma(0.001, 0.001)
 tau.sp~dgamma(0.001,0.001)
for(i in 1:nblocks){
 alpha.sp[i]~dnorm(0,tau.sp)
 Emu[i]=mu + alpha.sp[i]
for(t in 1:ndates){
  X[i,t] ~ dnorm(Ex[i], tau)
 }}}
11
```

```
Global Mean= "
model{
mu ~ dnorm(0,0.001)
tau ~ dgamma(0.001,0.001)

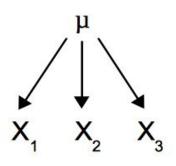
for(i in 1:nblocks){
  for(t in 1:ndates){
    X[i,t] ~ dnorm(mu, tau)
  }}}
```



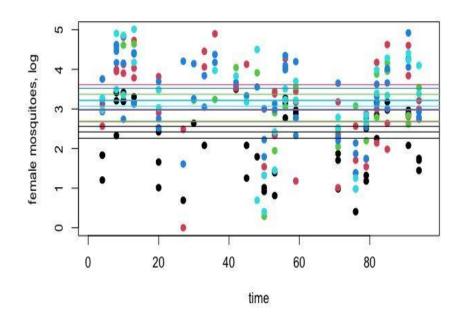
```
Space= "
model{
                            Parameter Model
 mu ~ dnorm(0,0.001) ~
                                 (Priors)
 tau ~ dgamma(0.001,0.001)
 tau.sp~dgamma(0.001,0.001)
for(i in 1:nblocks){
 alpha.sp[i]~dnorm(0,tau.sp)
 Emu[i]=mu + alpha.sp[i]
                            Process Model
for(t in 1:ndates){
  X[i,t] ~ dnorm(Ex[i], tau)
                             Data Model
 }}}
п
```

```
Global Mean= "
model{
mu ~ dnorm(0,0.001)
tau ~ dgamma(0.001,0.001)

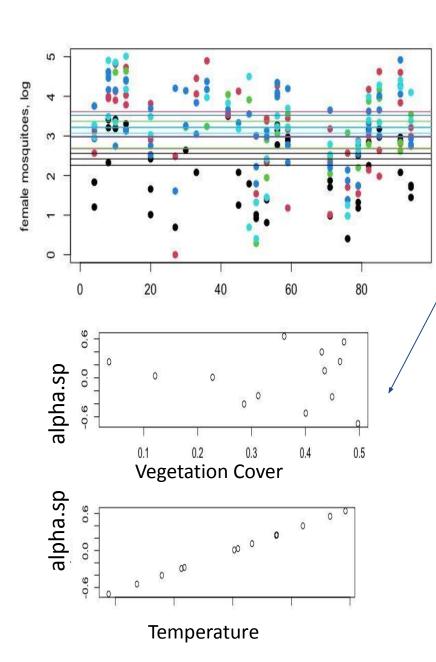
for(i in 1:nblocks){
  for(t in 1:ndates){
    X[i,t] ~ dnorm(mu, tau)
  }}}
```



```
Space= "
model{
                            Parameter Model
 mu ~ dnorm(0,0.001) ~
                                 (Priors)
 tau ~ dgamma(0.001,0.001)
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 Emu[i]=mu + alpha.sp[i]
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for(t in 1:ndates){
  X[i,t] ~ dnorm(Ex[i], tau)
                             Data Model
 }}}
п
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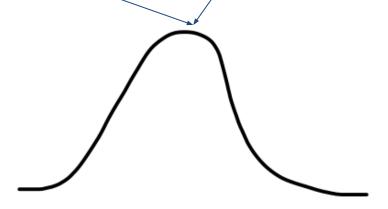
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  X[i,t] ~ dnorm(Ex[i], tau)
 }}}
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Space= "
model{
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 tau ~ dgamma(0.001,0.001)
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for(i in 1:nblocks){
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 Emu[i]=mu + alpha.sp[i]
for(t in 1:ndates){
  X[i,t] ~ dnorm(Ex[i], tau)
 }}}
п
```

Model	mu (95% ci)	tau (sigma)	tau.sp (sigma)	Deviance
Global	2.99 (2.98, 3.13)	0.85 (1.18)	NA	698
Spatial RE	2.97 (2.66, 3.28)	1.04 (0.96)	4.98 (0.20)	654

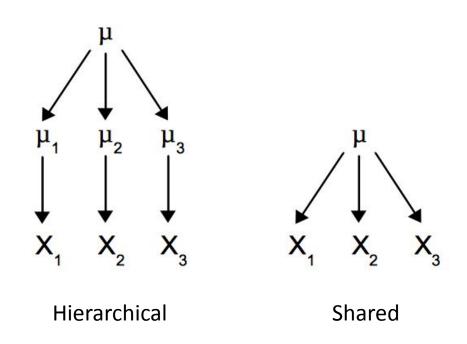
Model	mu (95% ci)	tau (sigma)	tau.sp (sigma)	Deviance
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Spatial RE	2.97 (2.66, 3.28)	1.04 (0.96)	4.98 (0.20)	654



defines a distribution that can be used to infer mean at out-of-sample site. Tighter REs (smaller variance) = more precise prediction

Hierarchical model allows predictions about the unobserved

Out-of-sample predictions integrate over the random effects variance in known sites/years – will include more uncertainty than in-sample estimates.



Ecology is complex. You cannot measure everything.

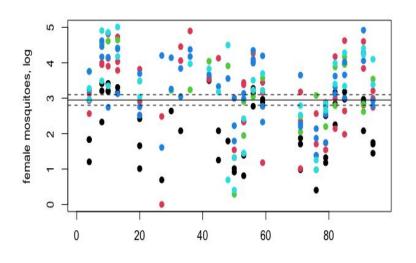
Hierarchical models can help identify important drivers and account for non-independence in sampling scheme

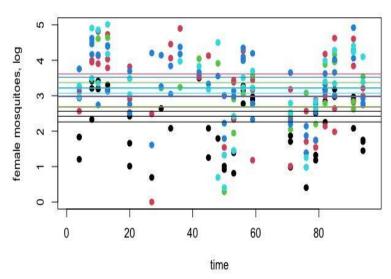
Hierarchical model allows for borrowed strength from datarich to data-poor

- predictions about the unobserved
- integrate over the random effects variance = more real reflection of uncertainty in out-of-sample predictions



Ecological Forecasting is about characterizing uncertainty





What will mosquito abundance be next month?

When and where should city invest in control?