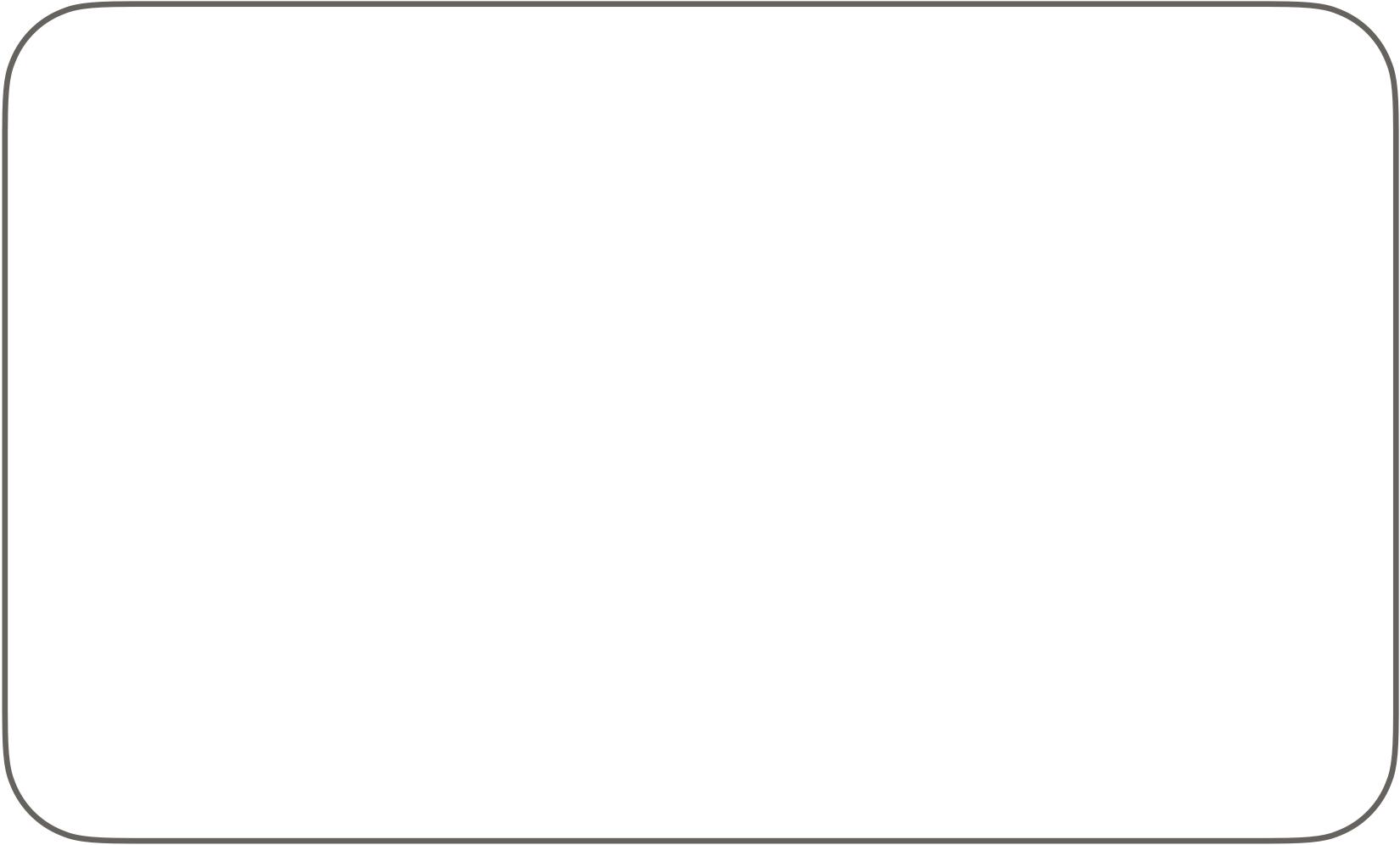




## RANDOM SLOPES MODEL



$$\begin{aligned} y_i &\sim Normal(\mu_i, \, \sigma) \\ \mu_i &= \alpha_0 + \alpha_{block[i]} + (\beta_0 + \beta_{block[i]}) x_i \\ \alpha_k &\sim Normal(0, \, \sigma_\alpha), \text{ for } k \text{ in } \{1, \cdots, N_{blocks}\} \\ \beta_k &\sim Normal(0, \, \sigma_\beta), \text{ for } k \text{ in } \{1, \cdots, N_{blocks}\} \\ \alpha_0, \beta_0 &\sim Normal(0, \, 1) \\ \beta &\sim Normal(0, \, 0.3) \\ \sigma, \sigma_{block} &\sim Exponential(1) \end{aligned}$$



Allows the slope associated with the x predictor to vary across blocks, but maintains a dependency across blocks



## RANDOM SLOPES MODEL

```
y_i \sim Normal(\mu_i, \sigma)
          \mu_i = \alpha_0 + \alpha_{block[i]} + (\beta_0 + \beta_{block[i]})x_i
         \alpha_k \sim Normal(0, \sigma_\alpha), \text{ for } k \text{ in } \{1, \dots, N_{blocks}\}
         \beta_k \sim Normal(0, \sigma_{\beta}), \text{ for } k \text{ in } \{1, \dots, N_{blocks}\}
   \alpha_0, \beta_0 \sim Normal(0, 1)
           \beta \sim Normal(0, 0.3)
\sigma, \sigma_{block} \sim Exponential(1)
```

Allows the slope associated with the x predictor to vary across blocks, but maintains a dependency across blocks

#### MORE COMPLEX STRUCTURING OF THE DEPENDENCY ACROSS COEFFICIENTS

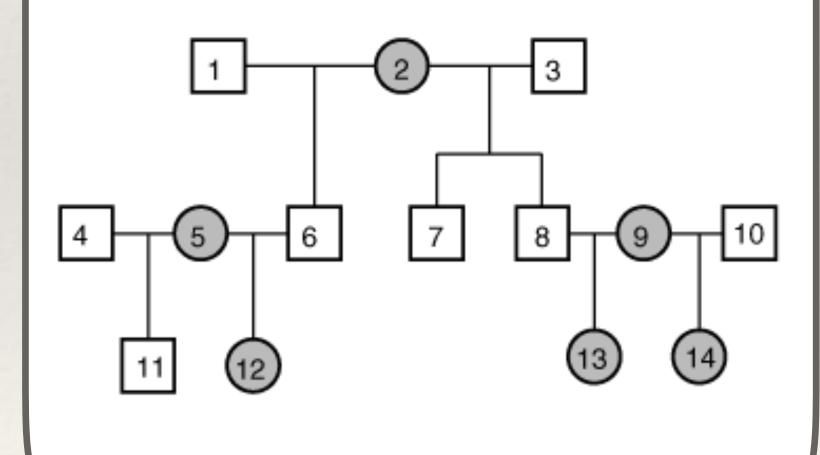
# Phylogenetic models

Evolutionary relatedness



### Animal models

Genetic relatedness



#### Spacial auto-correlation

Spacial proximity

