

Talks, travel, grants

Brachiopods

Mammals

- ▶ Evolution 2014: basic comparison between NA and European mammal survival
- ▶ GSA 2014: current fully Bayesian model of brachiopod survival
 - ▶ lots of positive feedback, ideas

Travel and grants

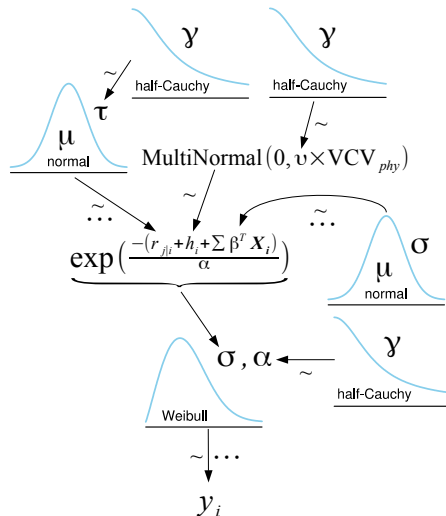
- ▶ AMNH: tooth measures for all notoungulate specimens identified to species level
- ▶ DDIG: applied; travel to Argentina; collaboration with Rick Madden

Survival model development

North American survival

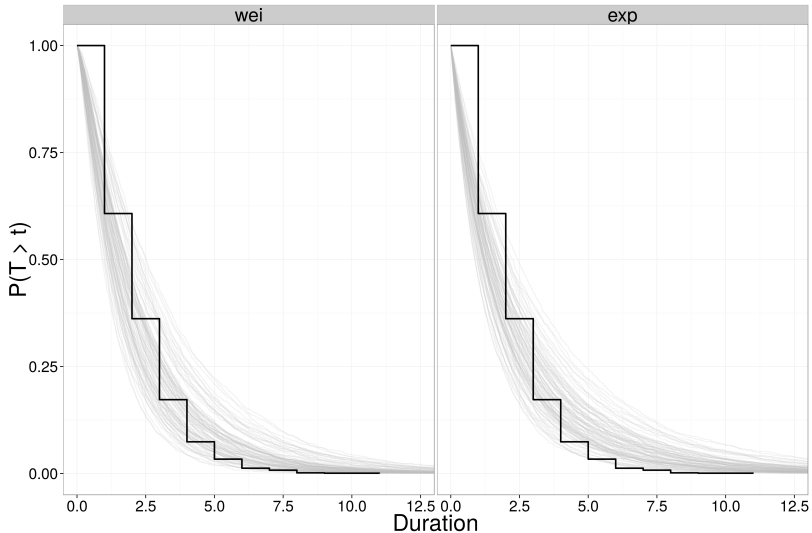
- ▶ species duration as measure of survival
- ▶ traits
 - ▶ organismal: diet, locomotor categories
 - ▶ species: body size, bioprovince occupancy
- ▶ origination cohort
- ▶ phylogeny primarily based on taxonomy
- ▶ duration defined as number of 2My bins from FAD to LAD, inclusive
- ▶ fully Bayesian hierarchical model
- ▶ censoring approach
 - ▶ if still extant, right censored
 - ▶ if not extant and duration of only 1 bin, left censored
- ▶ continuous traits rescaled following Gelman 2009 *Stats. Med.* by $2 * sd$.

Model diagram

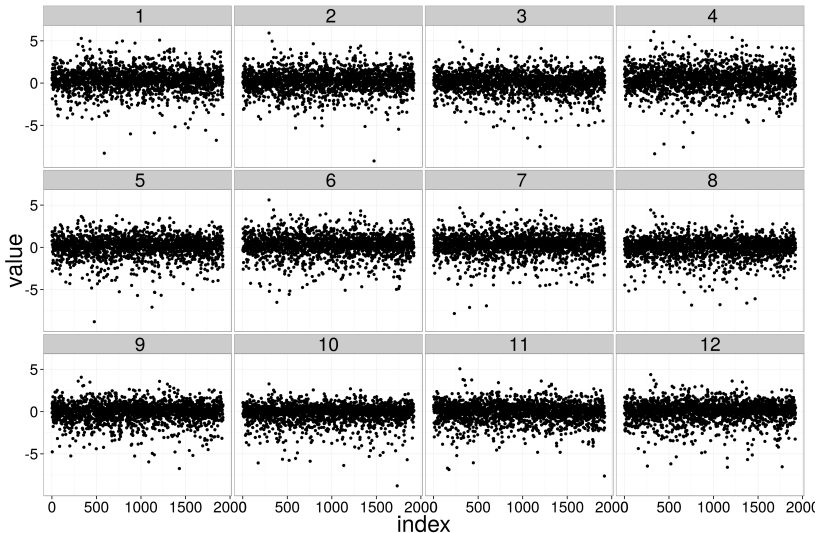


$$\begin{aligned}
 y_i &\sim \text{Weibull}(\sigma, \alpha) \\
 r_{j|i} &\sim \text{Normal}(0, \tau) \\
 \tau &\sim \text{half-Cauchy}(2.5) \\
 h_i &\sim \text{MultiNormal}(0, \Sigma) \\
 \Sigma &= v \times \text{VCV}_{phy} \\
 v &\sim \text{half-Cauchy}(2.5) \\
 \beta &\sim \text{Normal}(0, 10) \\
 \alpha &\sim \text{half-Cauchy}(2.5)
 \end{aligned}$$

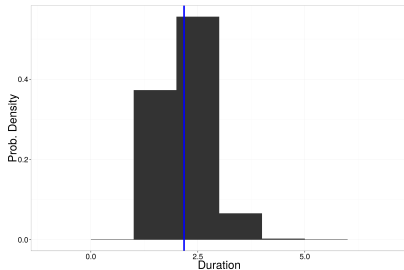
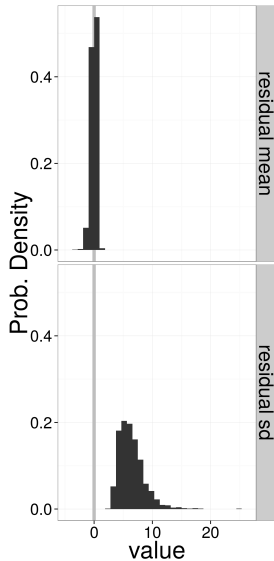
Posterior predictive checks: $S(t)$



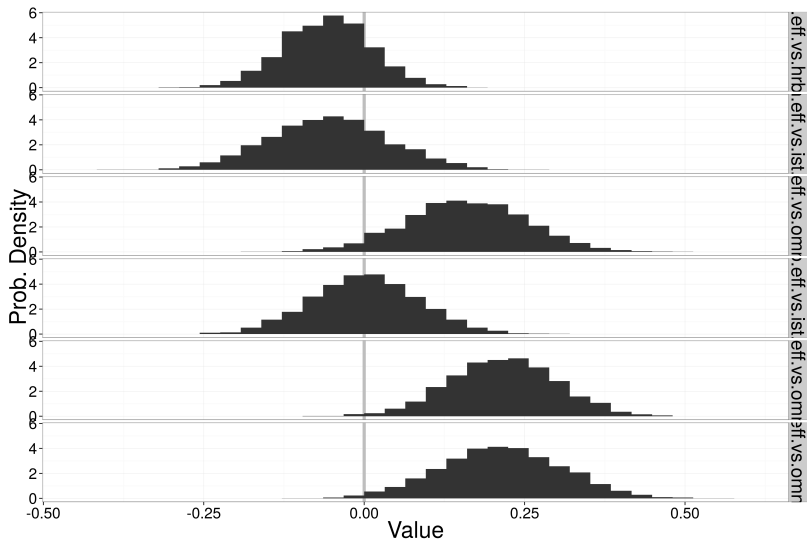
Posterior predictive checks: stdn. residuals



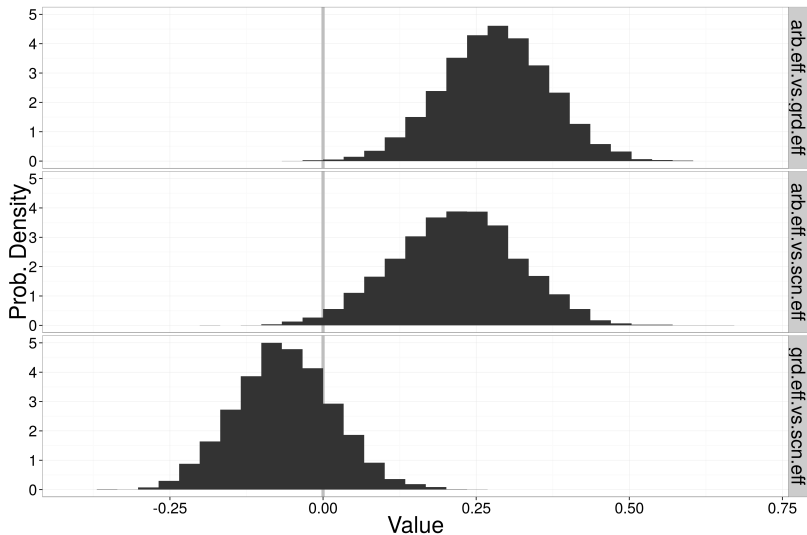
Posterior predictive checks: residuals and mean



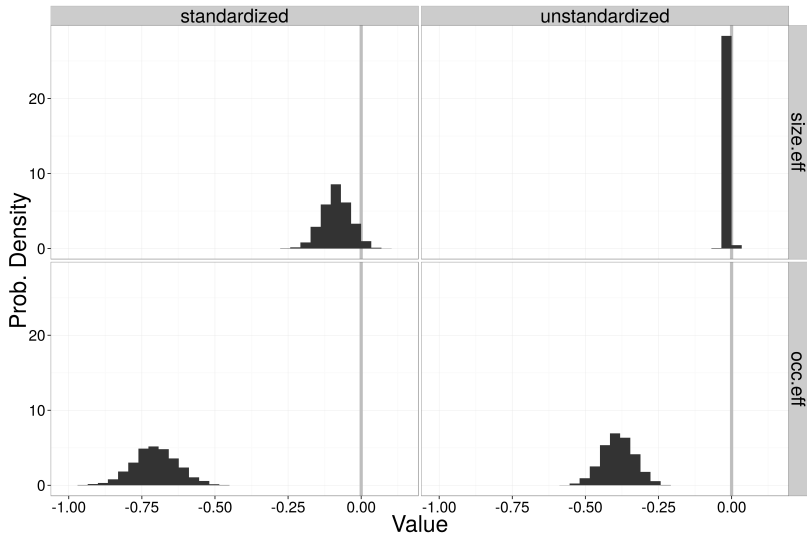
Pairwise differences of β , dietary category



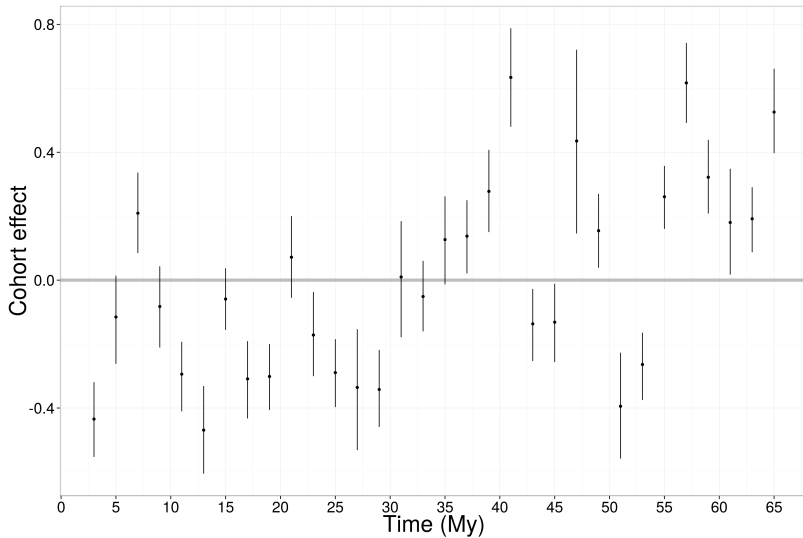
Pairwise differences of β , locomotor category



Other traits

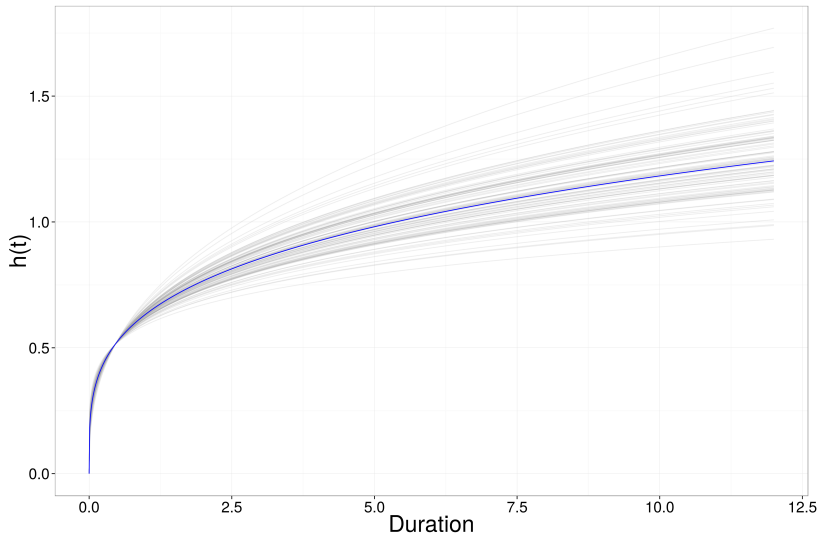


Cohort effect



Phylogenetic heritability *sensu* Lynch '91

Hazard curvature



Meaning and interpretation

- ▶ comparable probabilistic statements of trait, temporal, and historical effects
- ▶ increasing cohort survival risk over Cenozoic ▶
- ▶ $h(t)$ not constant over t , increases
- ▶ model fits. no systematic biases in residuals. noisy.