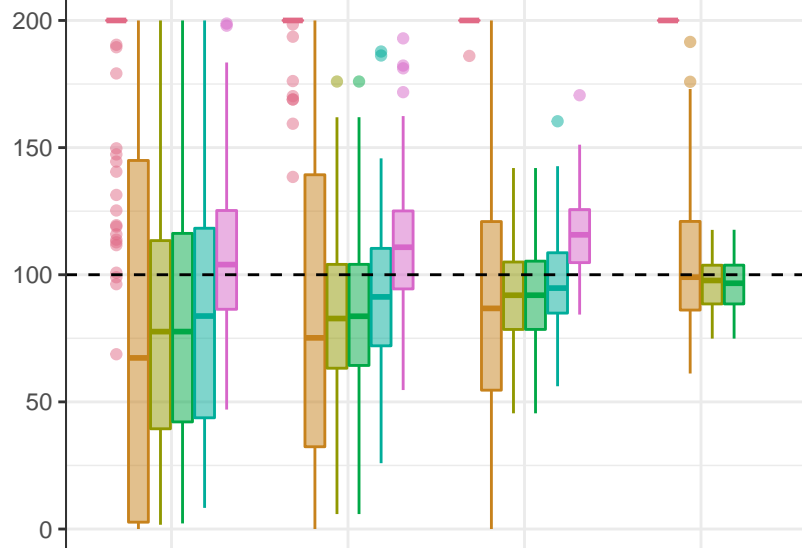
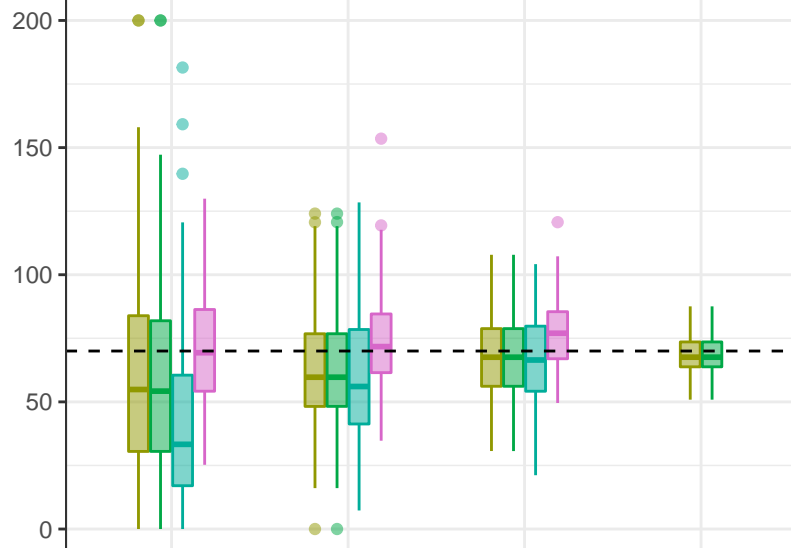
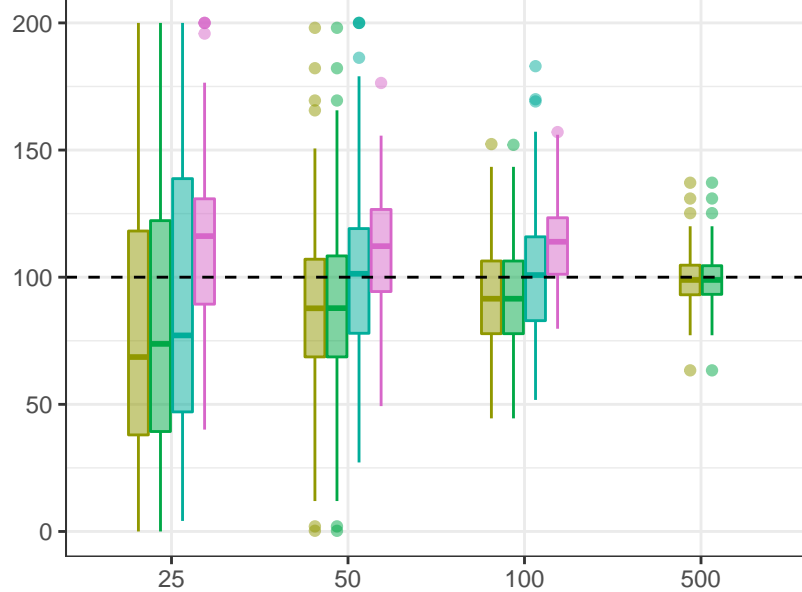
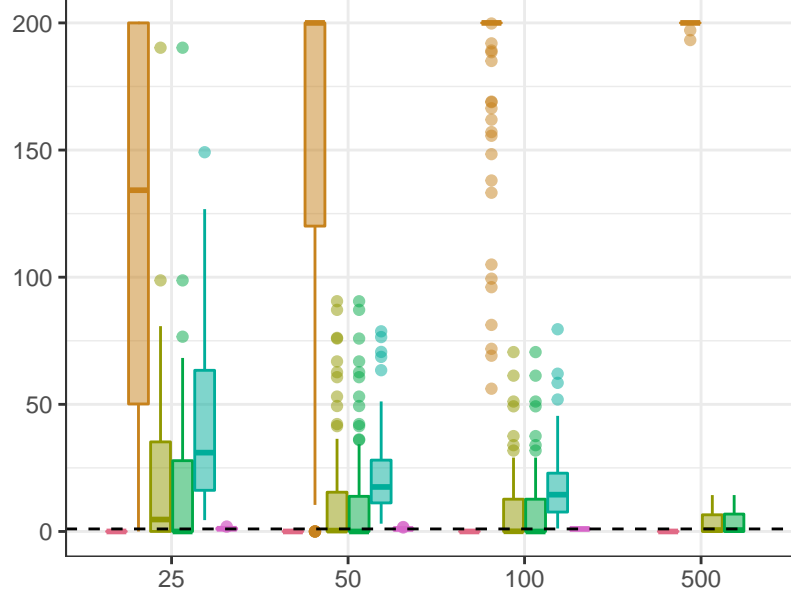


Parameter Estimates

Phylogenetic random intercept $\Sigma_{\text{phy}_{\text{int}}}$ Phylogenetic random intercept-slope correlation $\rho_{\text{phy}_{\text{int-slope}}}$ Phylogenetic random slope $\Sigma_{\text{phy}_{\text{slope}}}$ Residual σ_{ϵ} 

Platform

- glms
- phyloilm
- phyloglm/lme4
- phyloglm/glmTMB
- brms
- MCMCglmm

Number of Species