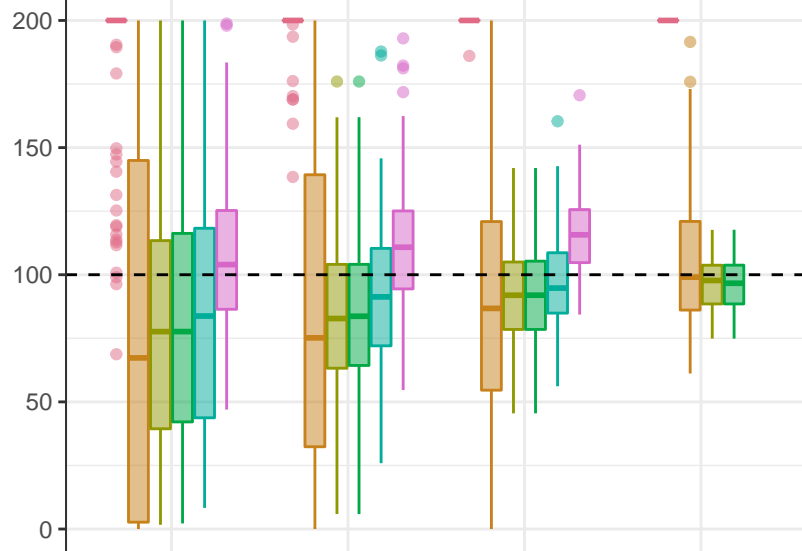
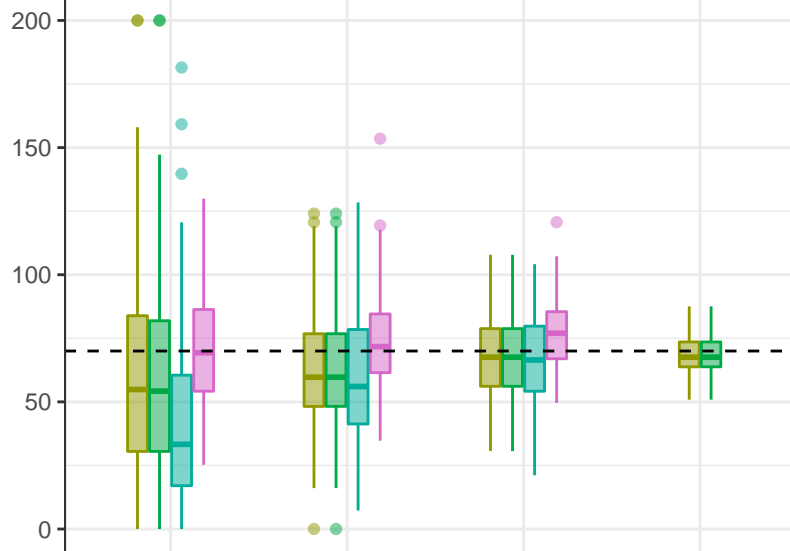
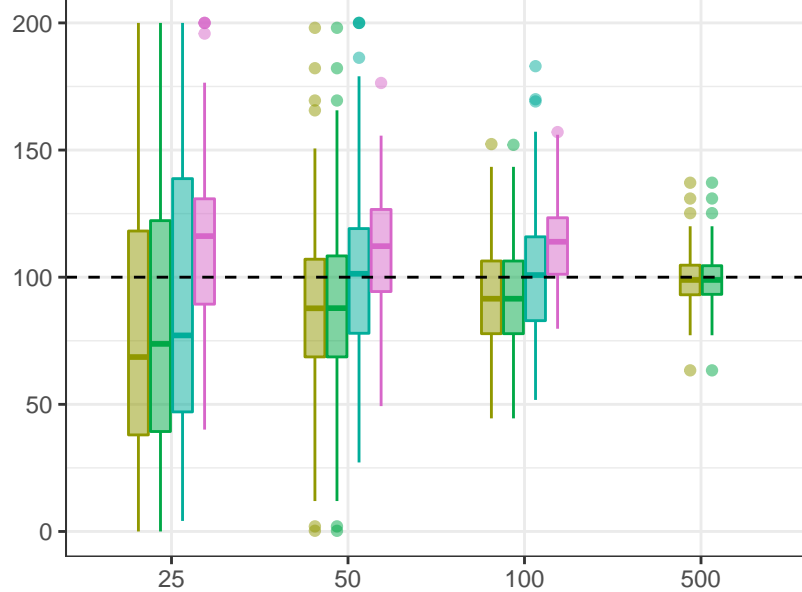
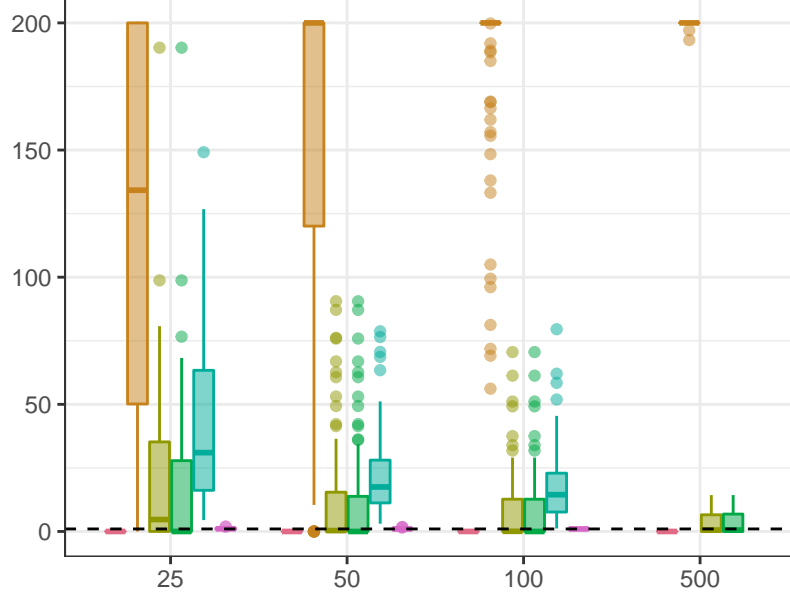


Parameter Estimates

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  $\sigma_{\epsilon}$ 

Platform

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

Number of Species