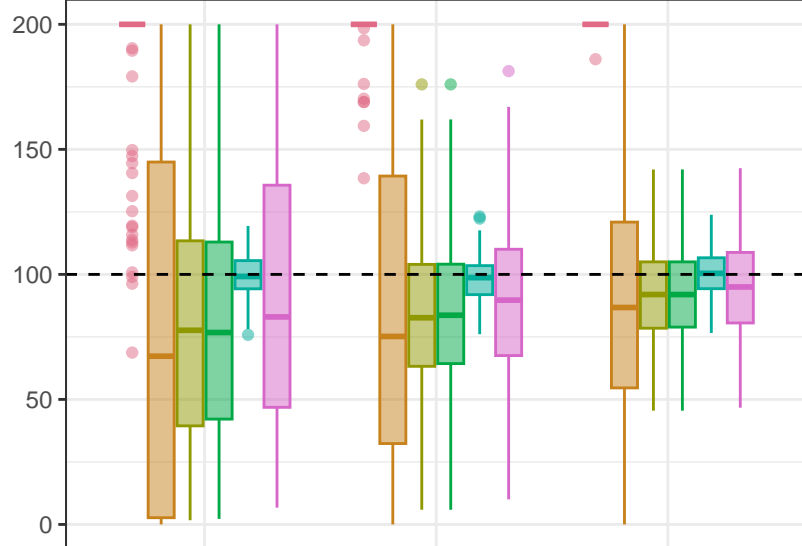
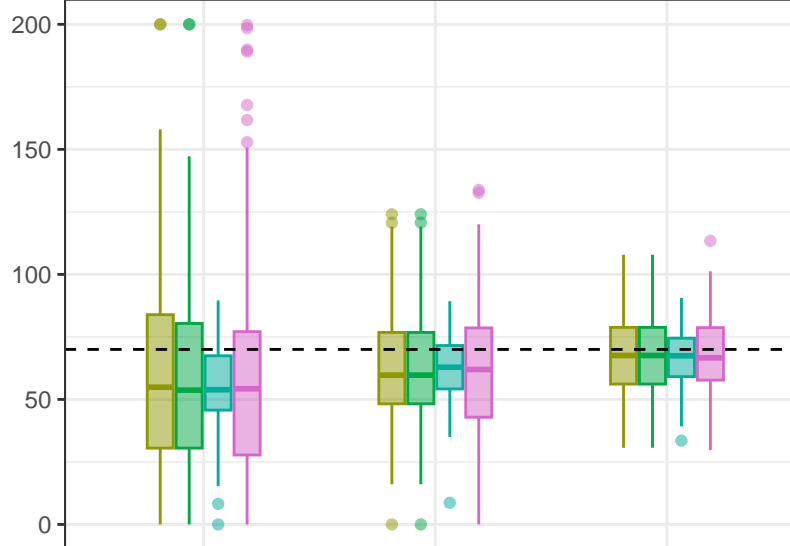
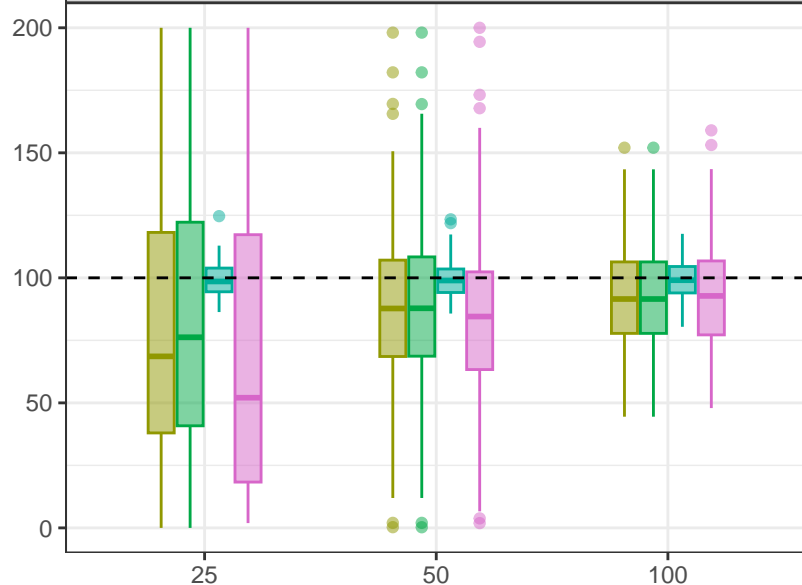
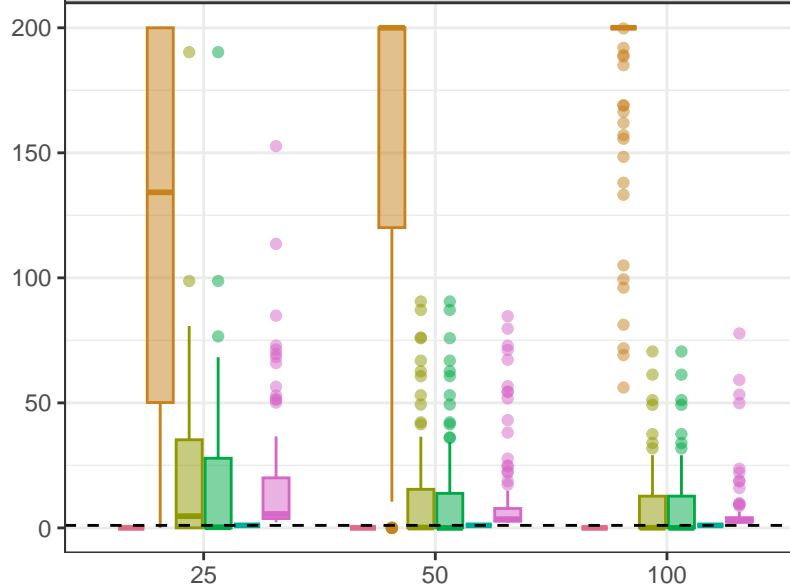


Parameter Estimates

Phylogenetic random intercept  $\sigma^2_{\text{phy}_{\text{int}}}$ Phylogenetic random intercept-slope covariance  $\sigma_{\text{phy}_{\text{int-slope}}}$ Phylogenetic random slope  $\sigma^2_{\text{phy}_{\text{slope}}}$ Residual  $\sigma^2_{\epsilon}$ 

Platform

- gls
- phylolm
- phyloglm/lme4
- phyloglm/glmTMB
- brms
- MCMCglmm

Number of Species