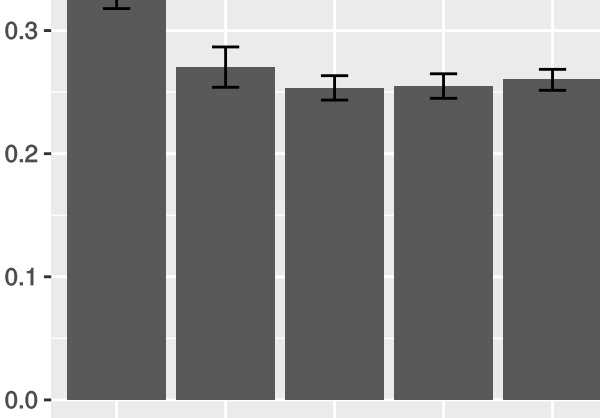
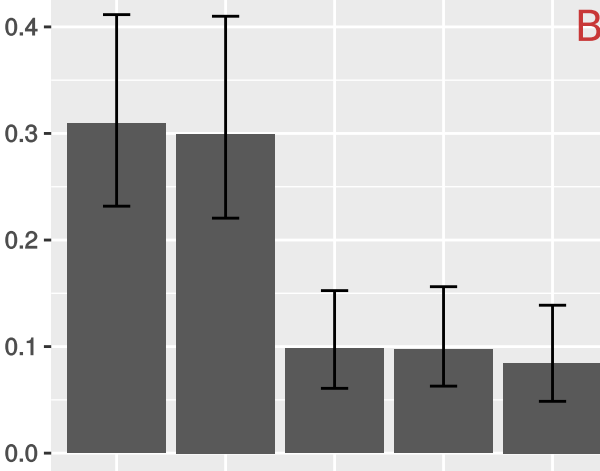


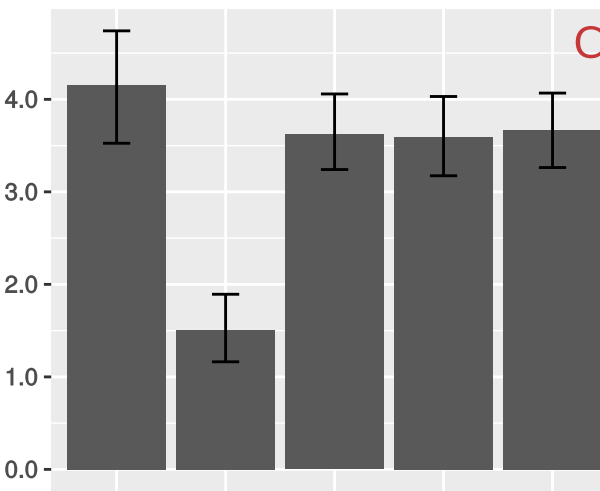
Relative species tree error



Mean pendant edge bias



Mean Robinson-Foulds distance



Bayesian phylogenetic method