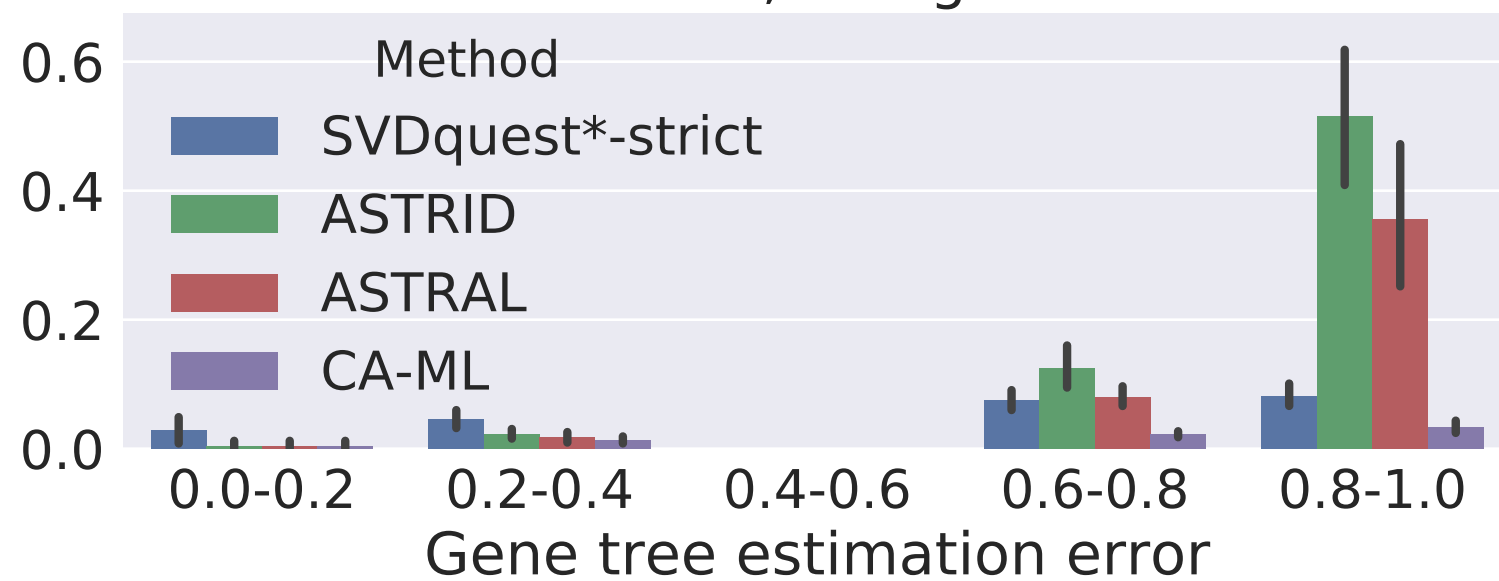
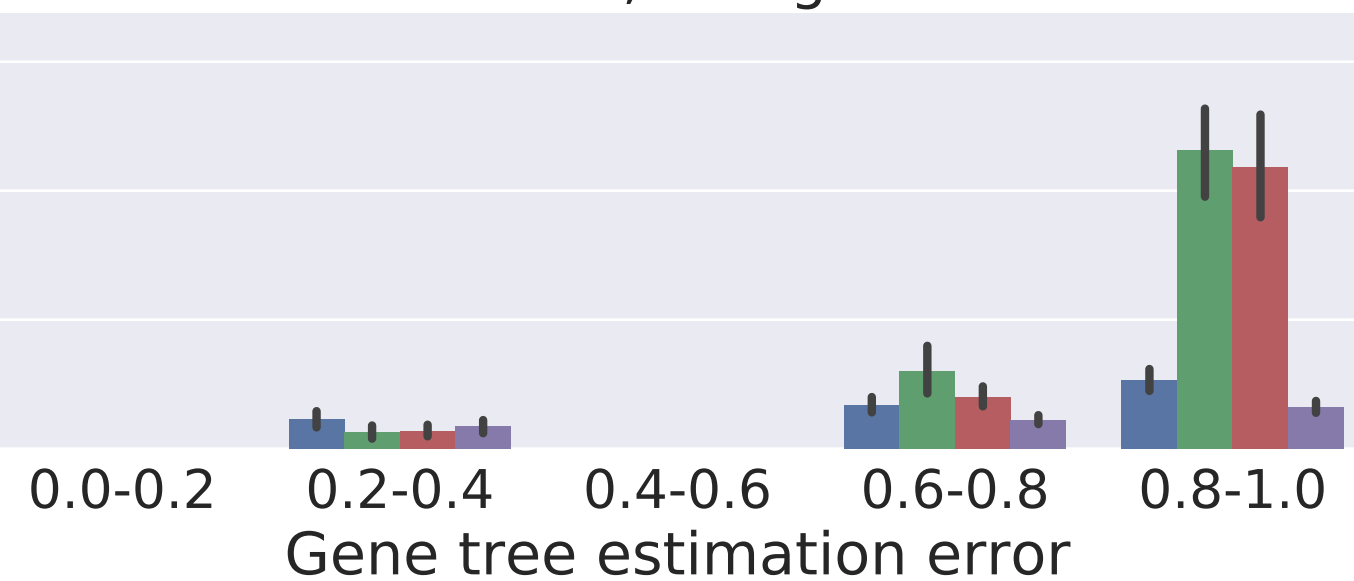


Species tree estimation error

13% AD, 500 genes



33% AD, 500 genes



72% AD, 500 genes

