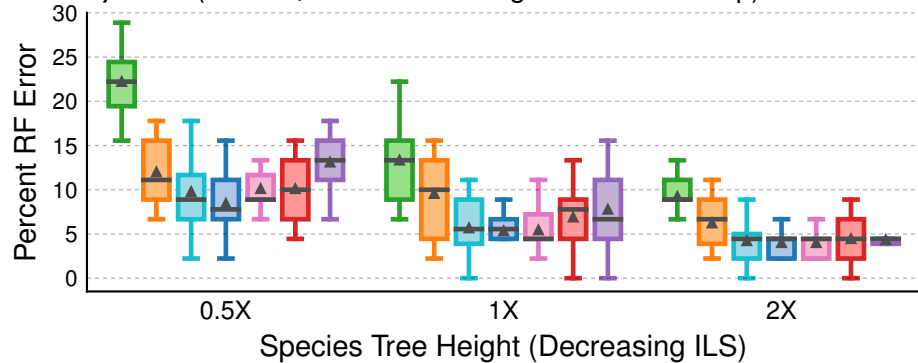
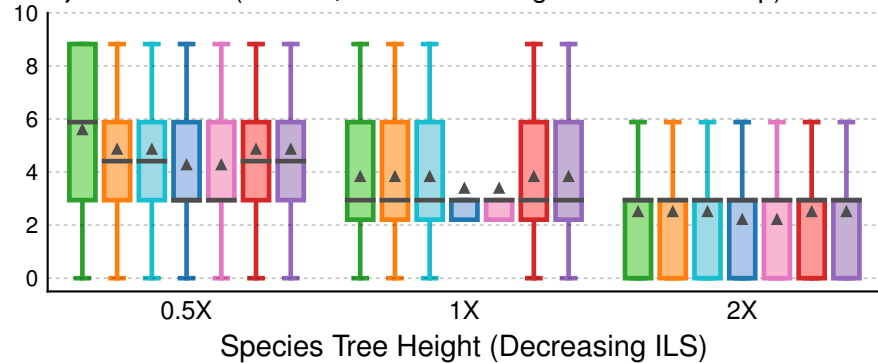


Impact of ILS

a) Avian (48 taxa, 1000 estimated gene trees - 500 bp)



b) Mammalian (37 taxa, 200 estimated gene trees - 500 bp)



wQMC

TQMC-n0

n1

n2

wQFM

FASTRAL

ASTRAL3