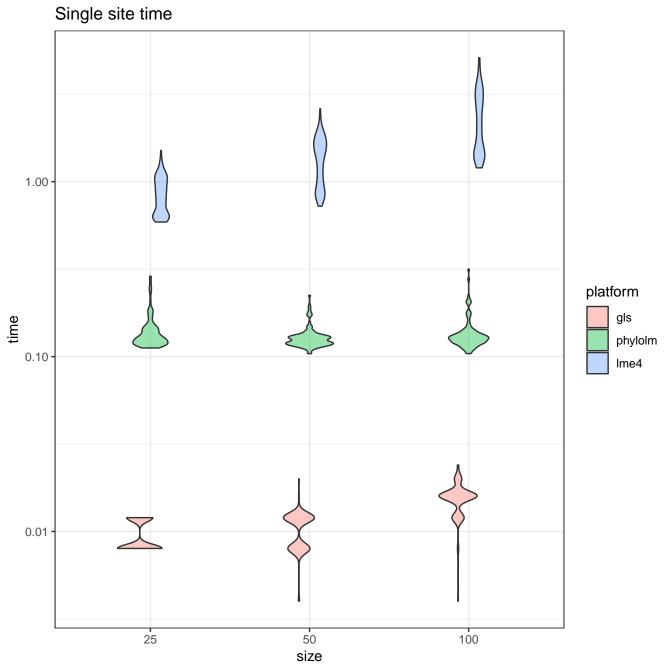
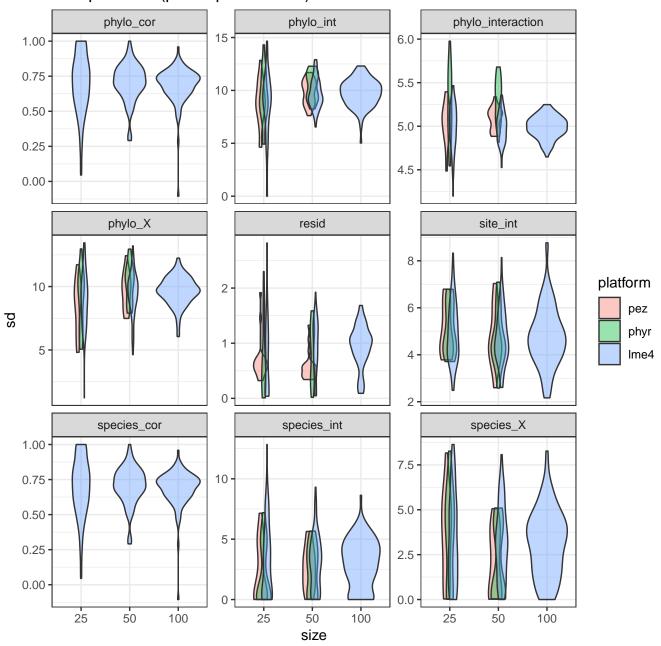
Single site phylo_cor phylo_int 50 -1.0 -40 0.5 -30 0.0 20 --0.5 **-**10 platform -1.00 gls sq phylo_X resid phylolm lme4 30 -20 -15 20 -10 10 -5 -0 0 25 50 25 50 100 100 size



Single Site coverage phylolm lme4 gls 1.0 8.0 platform gls phylolm coverage lme4 fixed_parameter B0_coverage B1_coverage 0.4 25 50 100 25 50 100 25 50 100 size

Multiple Sites (pez is problematic)



TODO: cowplot y-int simulation parameter phylo_cor phylo_int phylo_interaction 15 5.5 1.00 0.75 10 5.0 0.50 5 0.25 4.5 0.00 0 phylo_X resid site_int 8 10-2 platform 6 sd lme4 1 5. 4 0 species_X species_cor species_int 1.00 7.5 10 0.75 5.0 0.50 5 -0.25

0

25

50

size

0.00

25

50

100

2.5

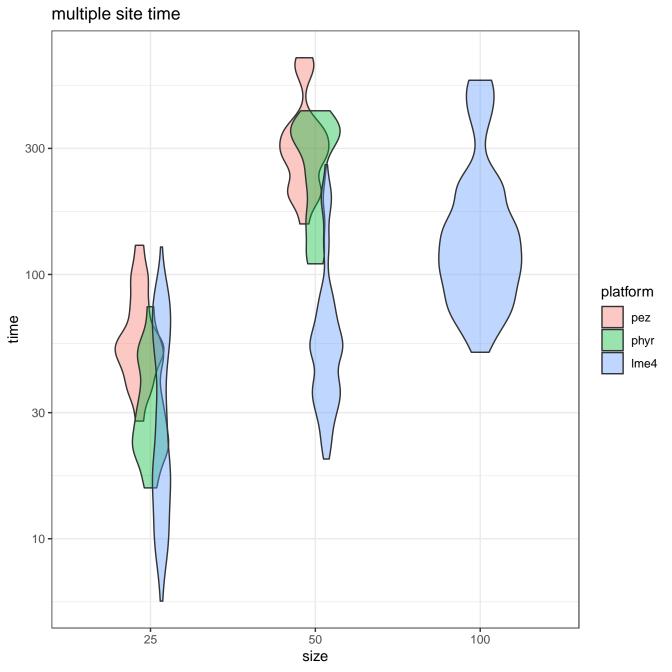
0.0

25

50

100

100



Single Site coverage phyr lme4 pez 0.95 0.90 fixed_parameter B0_coverage B1_coverage coverage platform pez phyr lme4 0.85 -0.80 -25 25 50 25 50 100 100 50 100 size