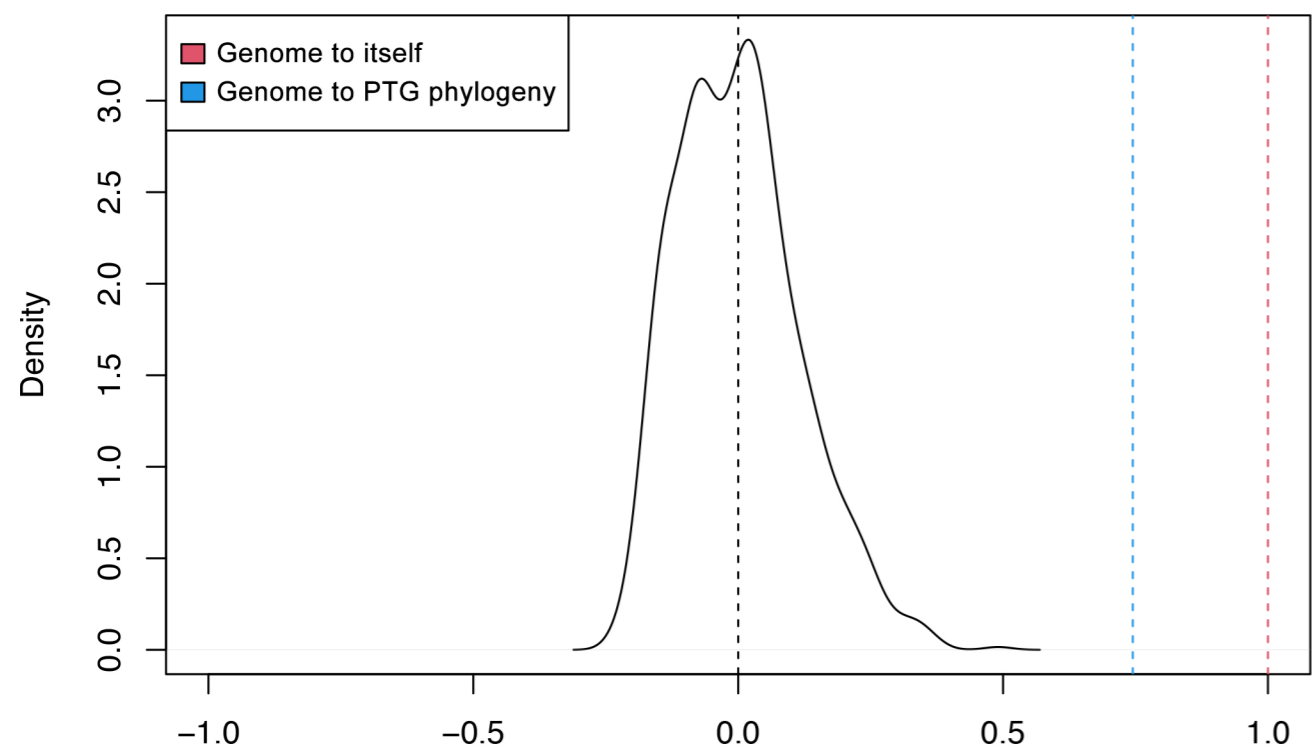
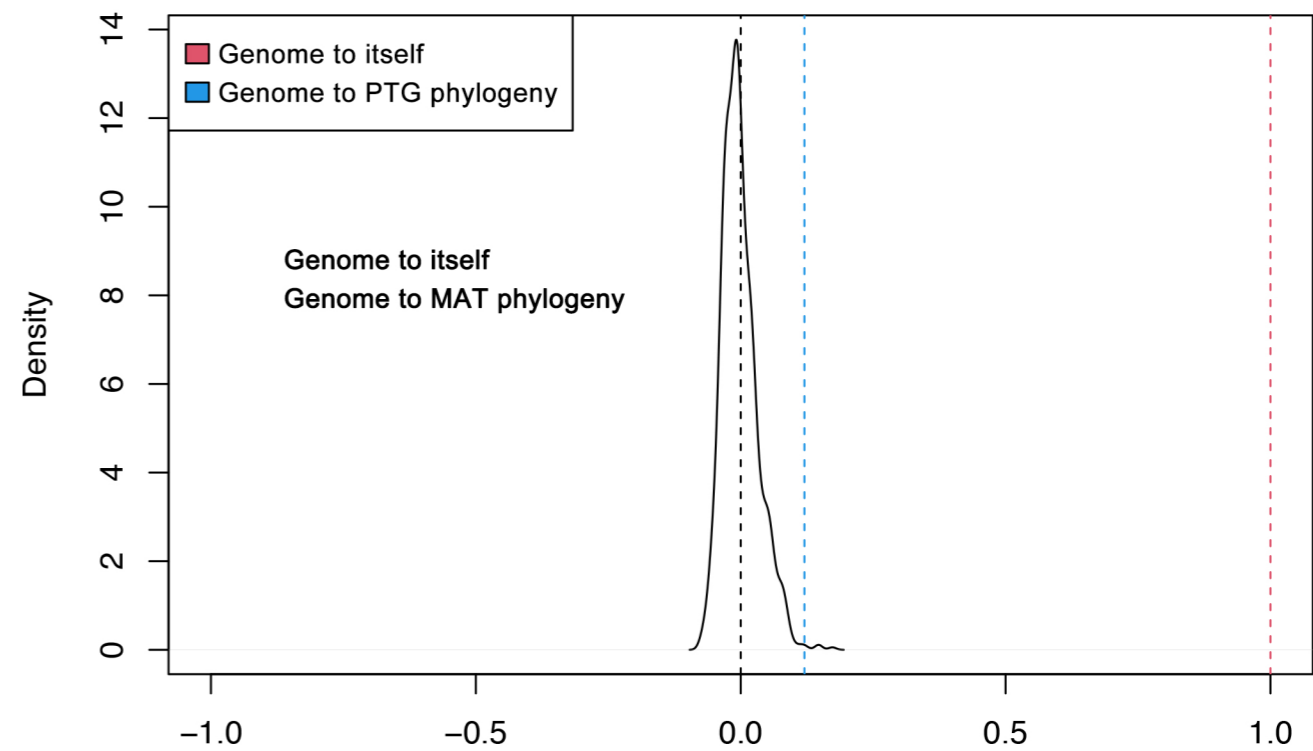


a Phylogeny congruency between Genome and PTG, Baker's gamma distribution



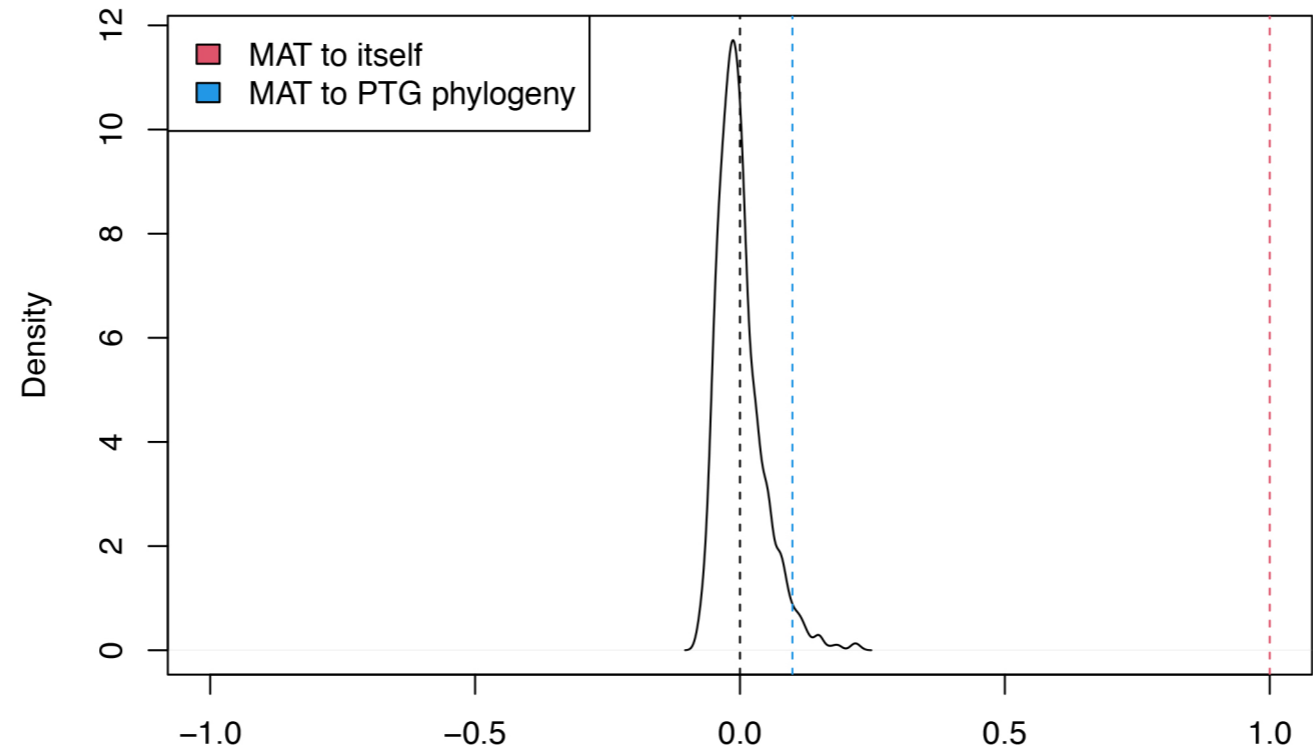
N = 1000 Bandwidth = 0.02619
p value for congruency significance: to itself = 0 ; to PTG phylogeny tree = 0

b Phylogeny congruency between Genome and MAT, Baker's gamma distribution



N = 1000 Bandwidth = 0.007129
p value for congruency significance: to itself = 0 ; to MAT phylogeny tree = 4E-03

c Phylogeny congruency between MAT and PTG, Baker's gamma distribution



N = 1000 Bandwidth = 0.00843
p value for congruency significance: to itself = 0 ; to PTG phylogeny tree = 3.2E-02