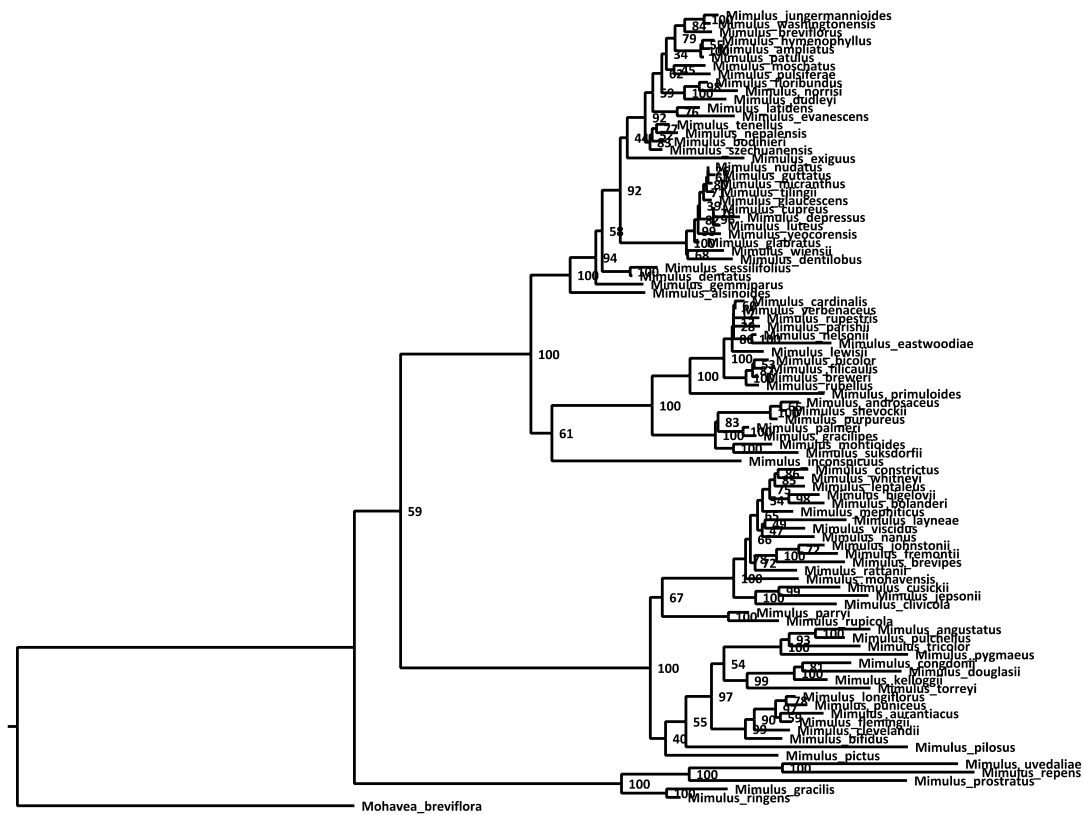


SuppFigure 1. Phylogenetic tree for *Mimulus* using Maximum Likelihood method with all 3 markers we have data for. Node labels= bootstrap percentages





SuppFig3. Bayesian Binary MCMC Distribution Tree: Ancestral Distribution of *Mimulus* on the Bayesian Tree. A= North America (blue), B= South America (Purple), C= Asia (Yellow), D= Oceania (Red).

result of combined:

