Supplementary Table 1: Comparison between phylogeographic spread inferred using parsimony and maximum likelihood on the same tree topologies

	Fréchet tree distance	Number of nodes with different locations	Amount of nodes with different locations
UPGMA	2174.67	3	0.016
NJ	4509.48	4	0.021
Parsimony	10000.33	12	0.063
MLJC	7917.40	10	0.053
MLGTR	8204.06	12	0.063