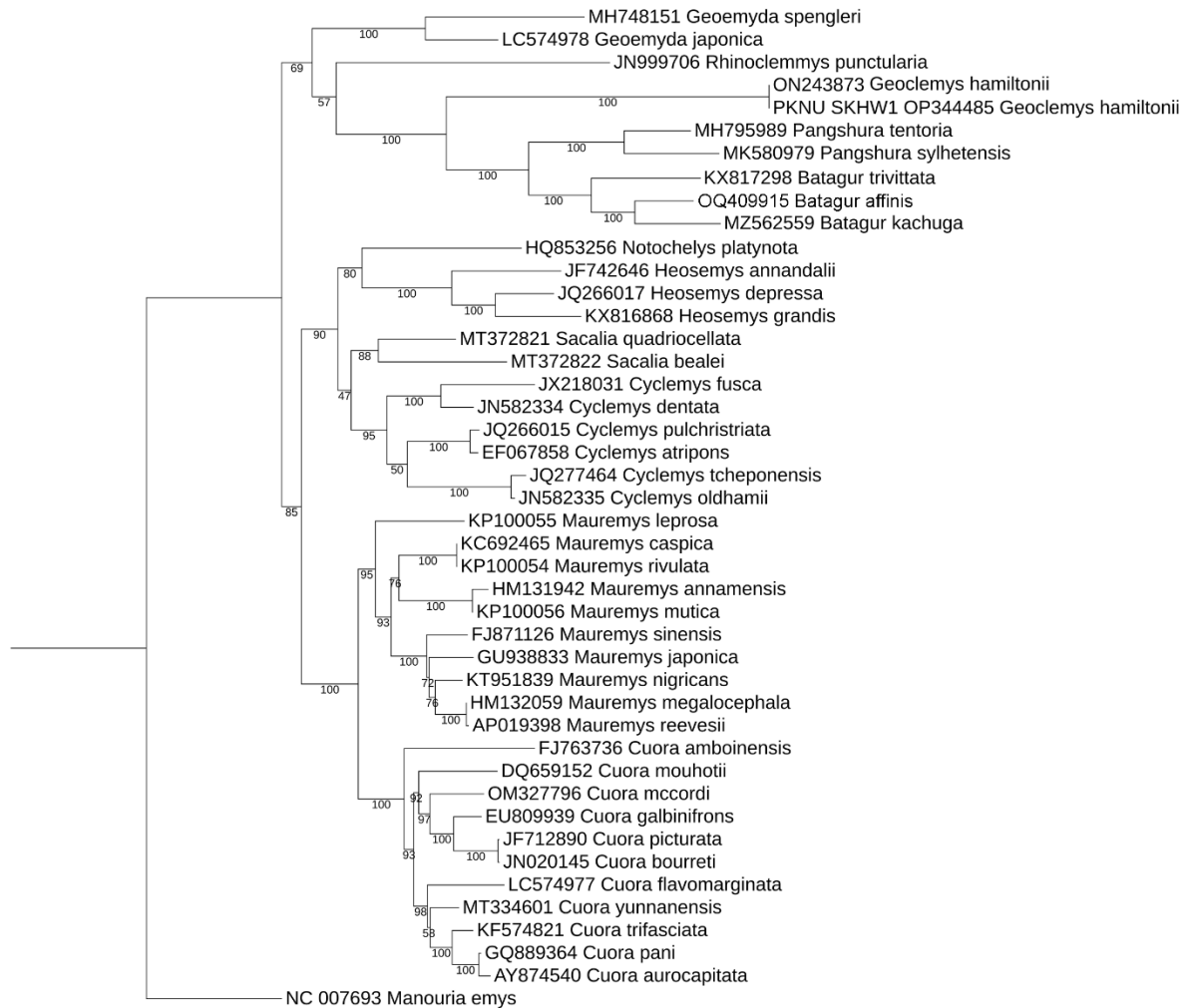


Figure S1. Maximum Likelihood (ML) phylogenetic tree based on the concatenated nucleotide sequences of 13 PCGs showing the phylogenetic position of *Geoclemys hamiltonii* and other Geoemydidae species.



Tree scale: 0.1