

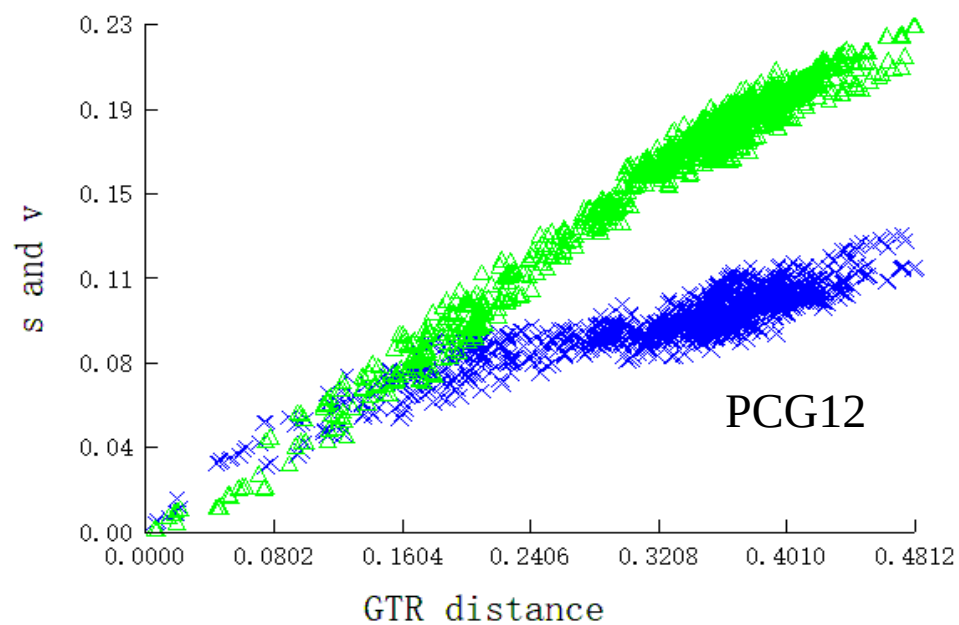
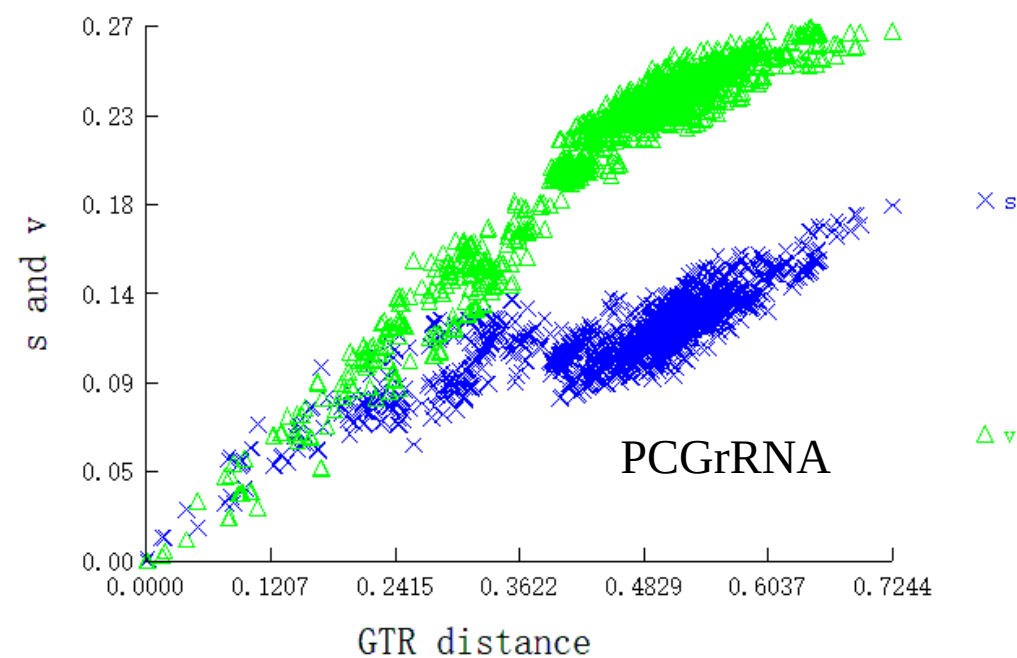
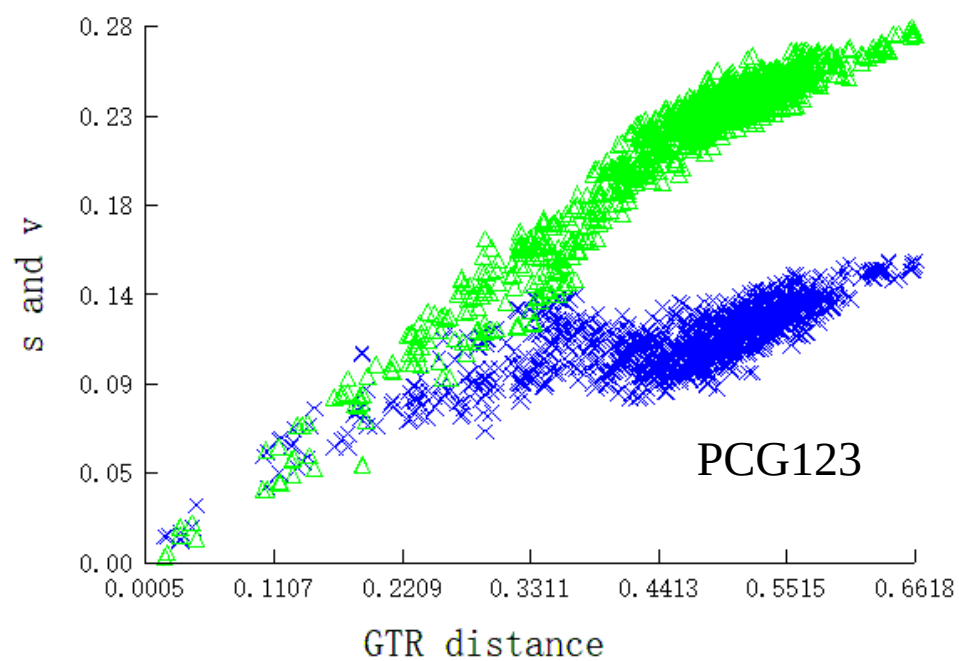


Fig. S4 Saturation analyses of 13 protein-coding genes (PCGs) and 2 tRNA genes in the 51 leech species, with GTR distance as abscissa and Transition (s) and Transversion (v) as ordinate.