

Figure S1. Genotype accumulation curve depicting the efficiency of SSRs in delineating the Cypriot *Vitis* spp. Genotypes.

Estimates of population differentiation

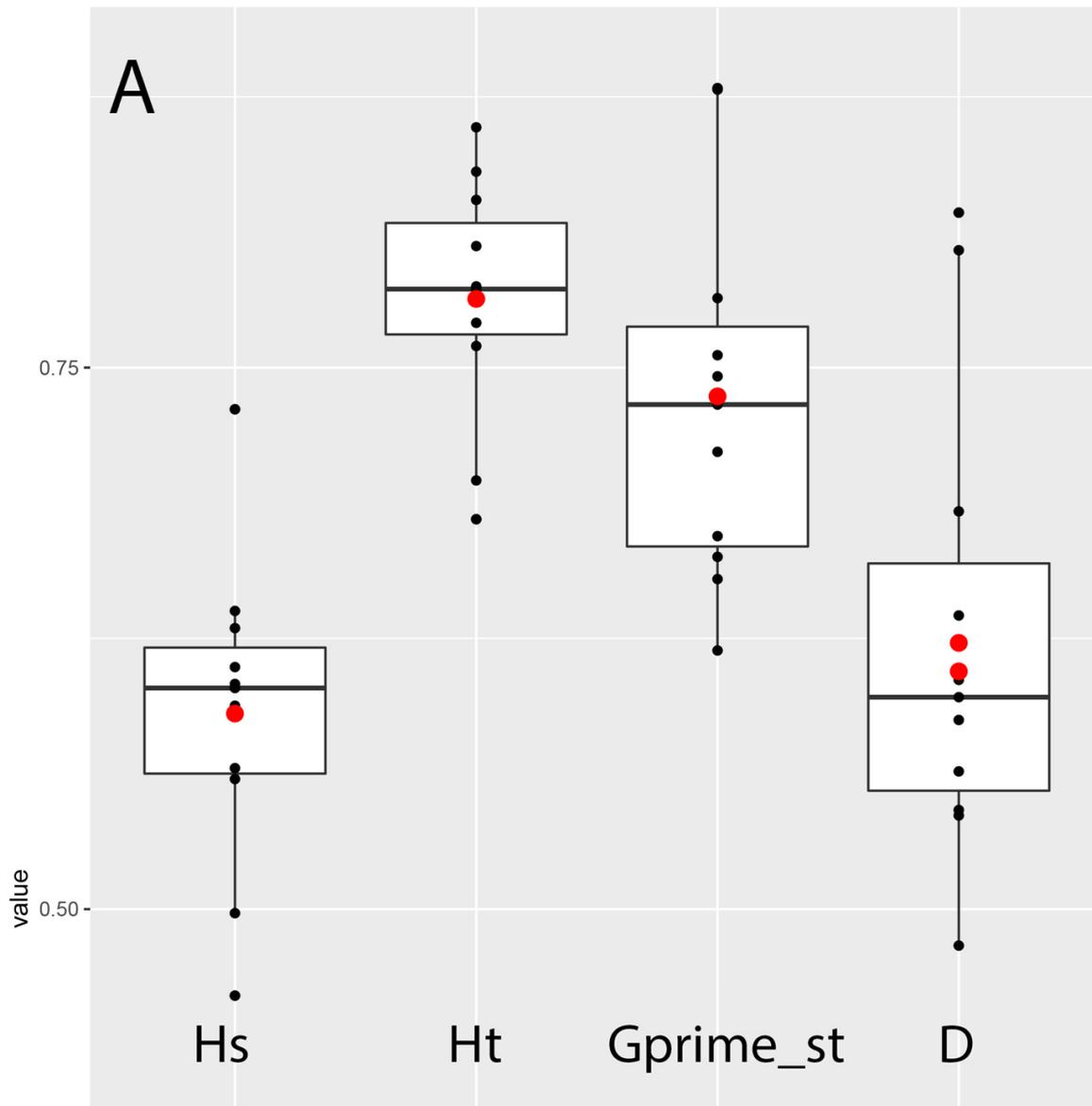


Figure S2. Genetic variability estimates from clone corrected genotypes. Boxplots depict the distribution of genetic indexes across the 11 loci studied. Hs: Heterozygosity with population structure, Ht: Heterozygosity without population structure, Gprime_st: Hedrick index, D: Jost index (all indexes were calculated using the Poppr package).

Tree scale: 0.1

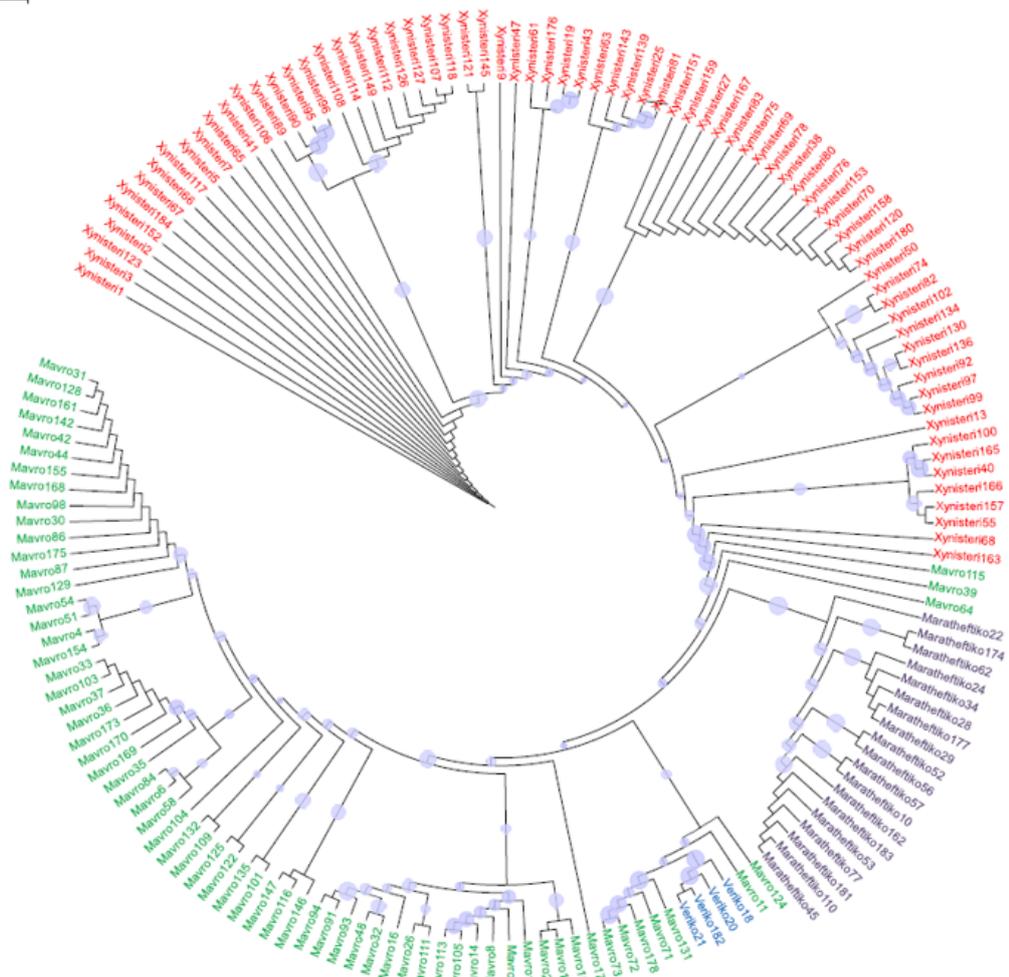


Figure S3. Phylogenetic tree of Cypriot grapevine genotypes. The size of nodes correlates to bootstrap support values.