

Figure S2. Genic frequencies for the 32 populations at the 6 intron markers

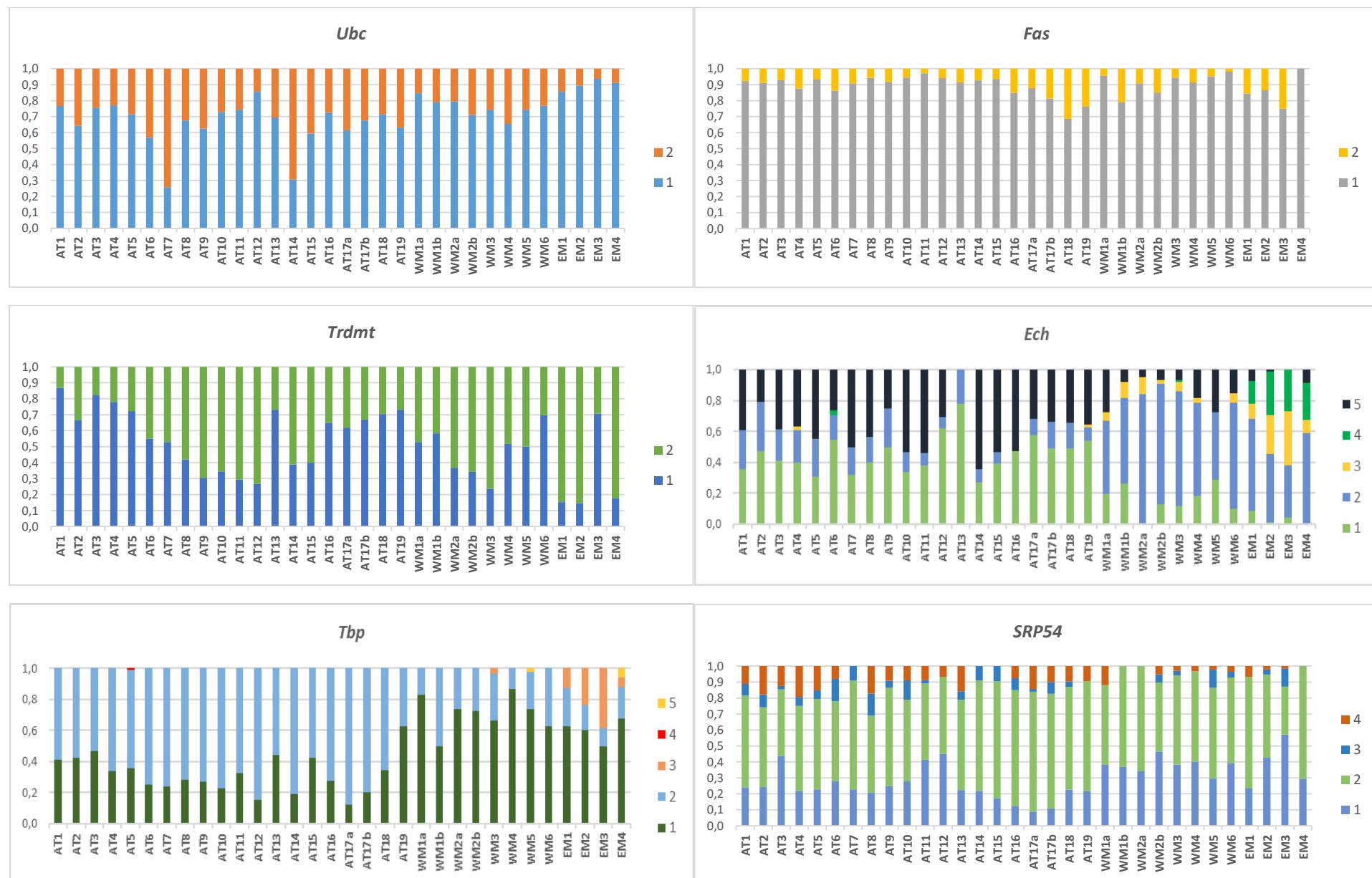


Figure S3. Contribution of alleles at each locus to the clusters of the Bayesian analysis of genetic structure, represented as the absolute value of the difference in allelic frequency between clusters. **(a)** K= 2. **(b)** K=6.

