

Figure S1. Correlation of geographic distance (in kilometers) and genetic distance (pairwise F_{ST}) among 12 populations of *J. regia*, including regression line ($r = 0.14$, $p = 0.22$ at 1000 randomization).

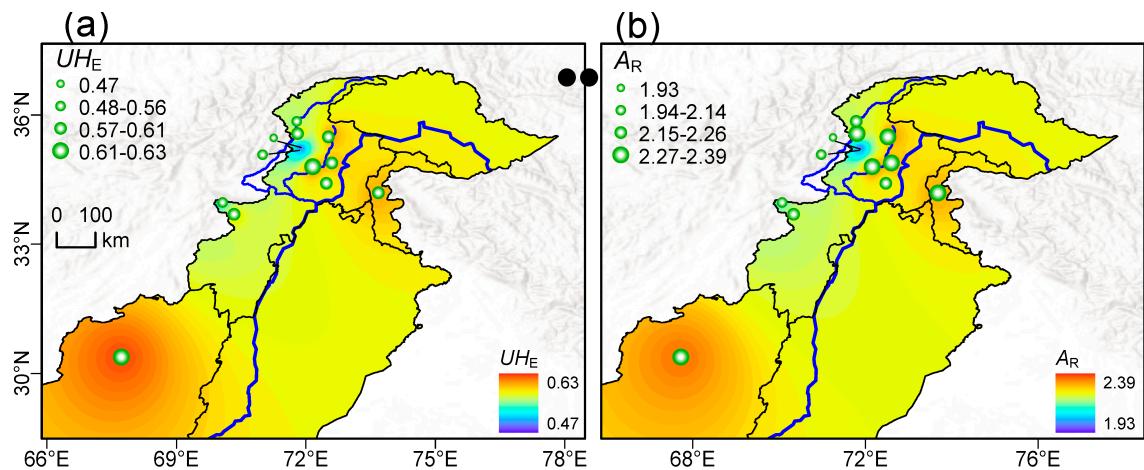


Figure S2. Spatial genetic diversity of 12 populations of Pakistan walnut: (a) Unbiased expected heterozygosity (UH_E), (b) allelic richness (A_R).

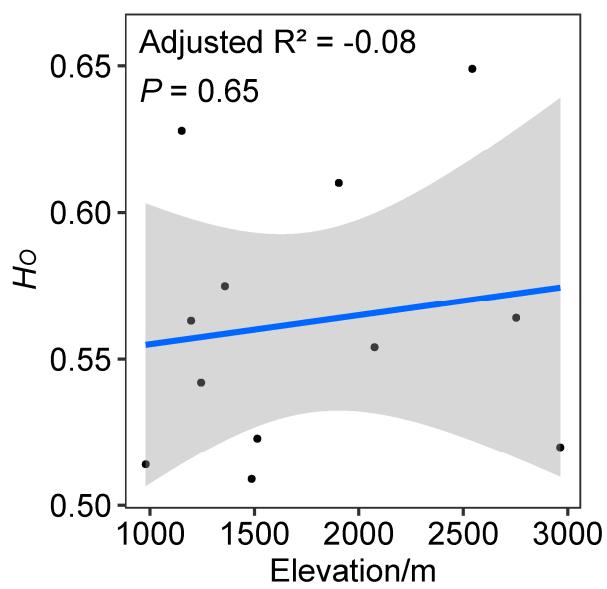


Figure S3. Relationship between observed heterozygosity and elevation for 12 populations of walnut in Pakistan.