Figure S2. Scatterplot expected heterozygosity, unique allele richness and arc surface distance from the most eastern sample. A least-square regression line represents the relationship between the two variables. Significance of the correlation was tested using Spearman’s $r$ (expected heterozygosity: $r=-0.66$, $P<0.0001$; unique allele richness: $r=-0.58$, $P<0.0013$).