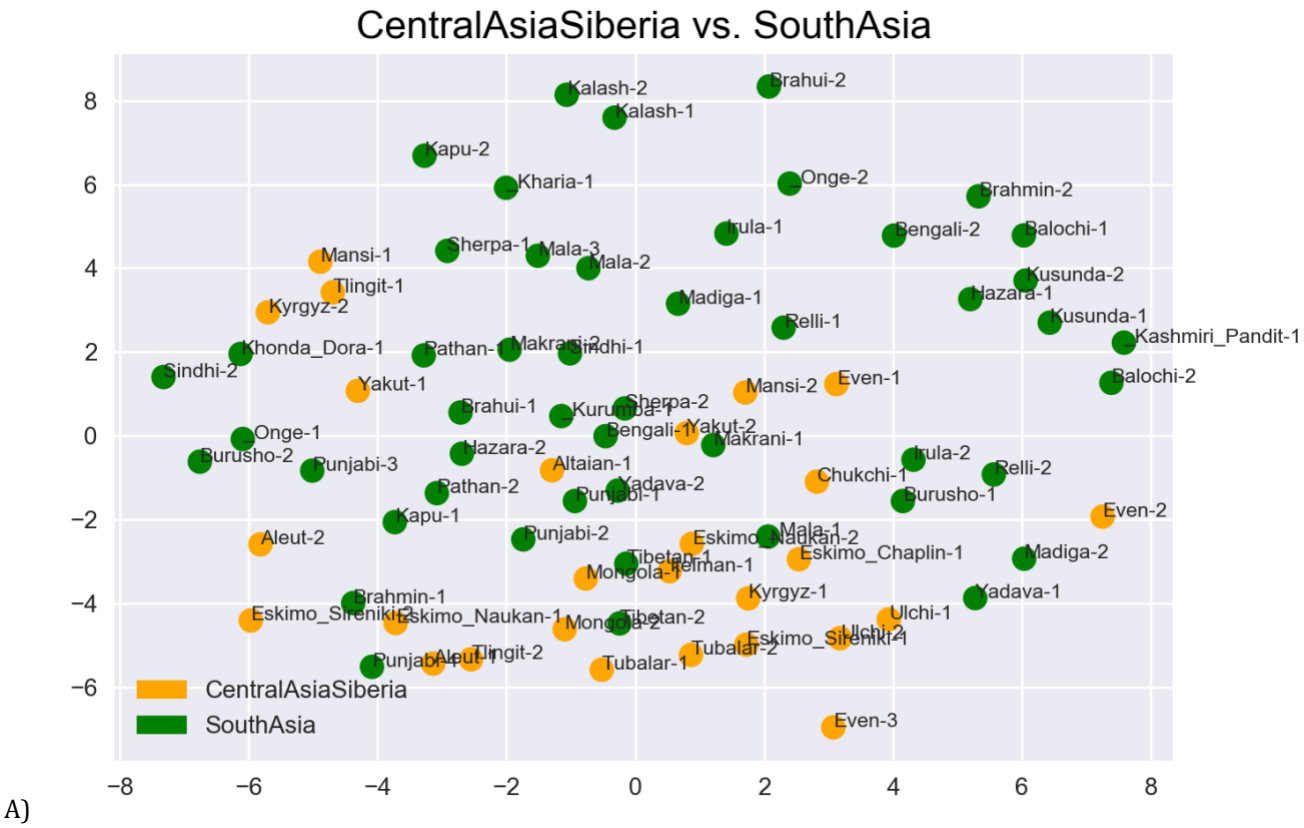
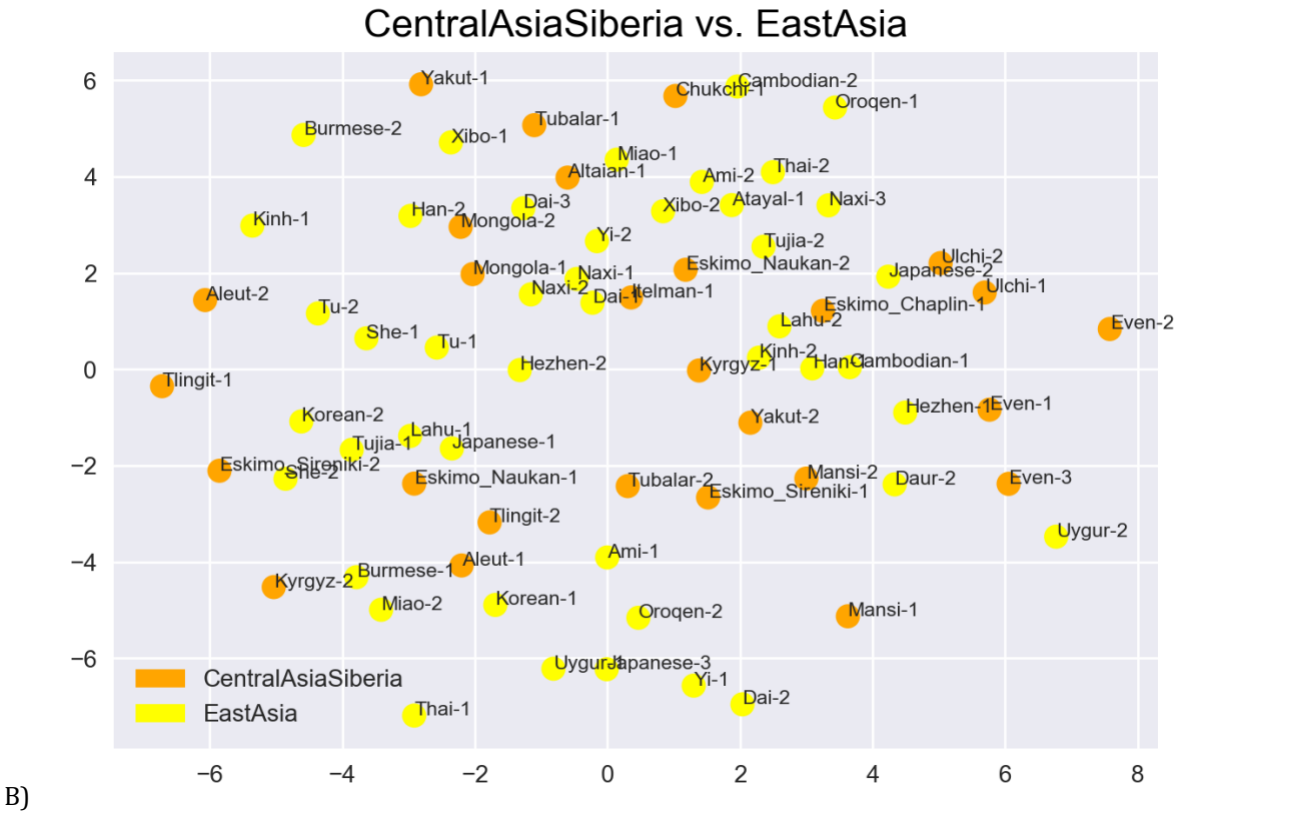


A)



B)

Supplementary Information Figure 5. Multidimensional scaling of 27 individuals from CentralAsiaSiberia and A) 49 individuals from SouthAsia and B) 45 individuals from EastAsia. Data used includes all DNA source types in the SGDP dataset. The metric used for scaling was based on the Euclidean distance between the set of alleles of each individual. Individuals are labeled by an abbreviated version of their sample ID in the SGDP dataset.