

Table S1: Individuals and markers used for analyses.

Analysis	Individuals (sample size)	# of SNPs
Frappe, PCA for genome-wide ancestry; Estimation of locus-specific ancestry	CEU (88); YRI (100); MEX1 (984); MEX2 (46)	482906
Within European PCA	Irish (43); Polish (45); Portuguese (43); Italian (45); MEX1 ^{EUR} (633); MEX2 ^{EUR} (46)	162382
Within Indigenous American PCA	Indigenous Americans (129); MEX1 ^{AMR} (933); MEX2 ^{AMR} (26)	162382
Signature of selection	MEX1 (957)	482906