

Table S2: Mean Rsq values for genome-wide significant loci in the GWAS of allergic rhinitis among Latinos

SNP	Rsq*
rs12973620	0.906
rs6583203	0.722
rs17133587	0.598
rs11680788	0.624

*estimates the squared correlation between imputed and true genotypes. Typically, a cut-off of 0.30 will flag most of the poorly imputed SNPs, but only a small number (<1%) of well imputed SNPs [59].