

Figure S1. The DIND test was performed for CEU (A) and EAS (B). P values were calculated through coalescent simulations that incorporate demographic scenarios. The continuous and hatched blue lines indicate the median and the 95 th percentile, respectively. In the case of EAS, the test was performed using all chromosomes (green dots) and with the exclusion of the single divergent chromosome carrying Hap I (red dots). Line colors refer to the different demographic models: Marth et al. (blue), Voight et al. (green), Gutenkunst et al. (red).

Table S2. Statistical significance of summary statistics calculated using different demographic models. Data refer to the APOBEC3H region we resequenced.

Table S1. Information concerning resequenced HapMap Individuals, APOBEC3H SNPs, and haplotypes.

dind.rtx; fas2ms.rtx: R scripts to compute DIND statistics.

Haplotypes were inferred using PHASE.

Coalescent simulations were performed using both MS and COSI software.