

Figure S1

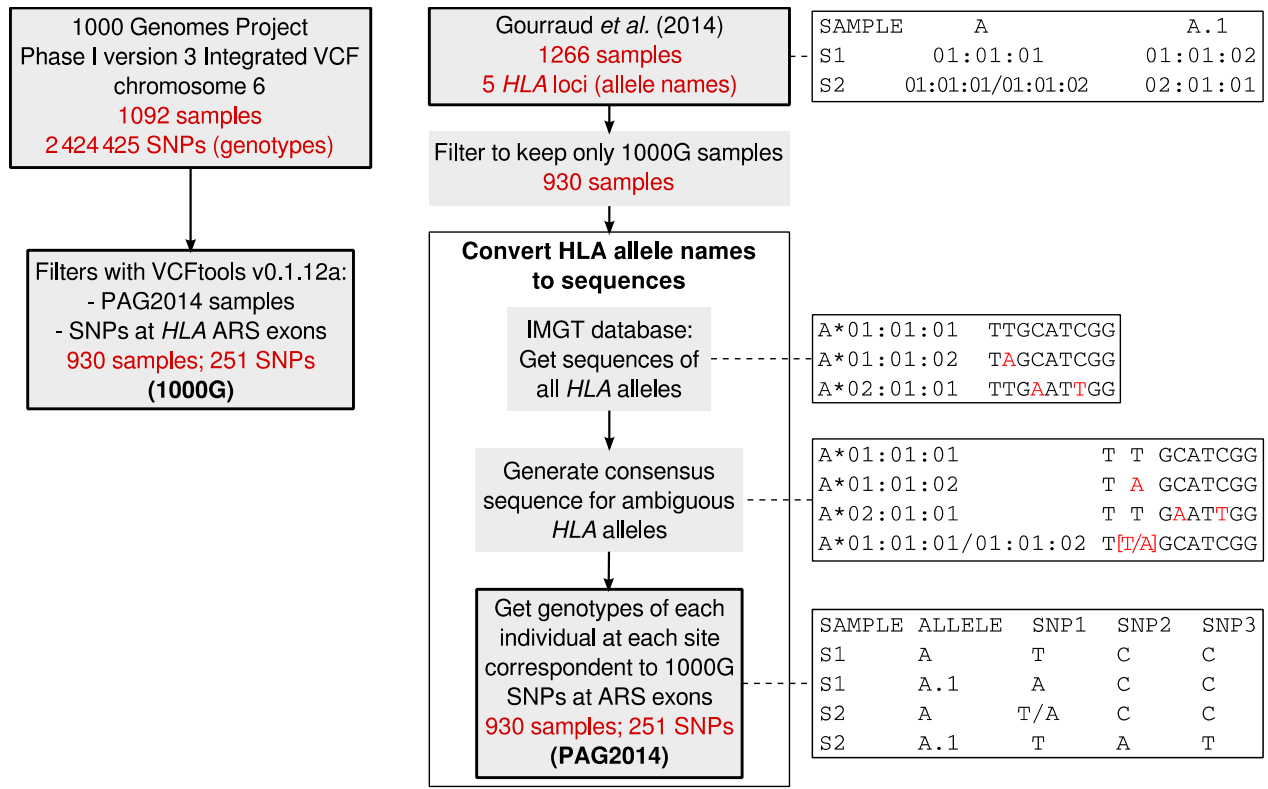


Figure S1 Workflow for preparation of next generation sequencing dataset from the 1000 Genomes Project (1000G) and Sanger sequencing dataset generated by Gourraud *et al.* (2014) (PAG2014) for comparisons of genotypes and allele frequencies (see main text).