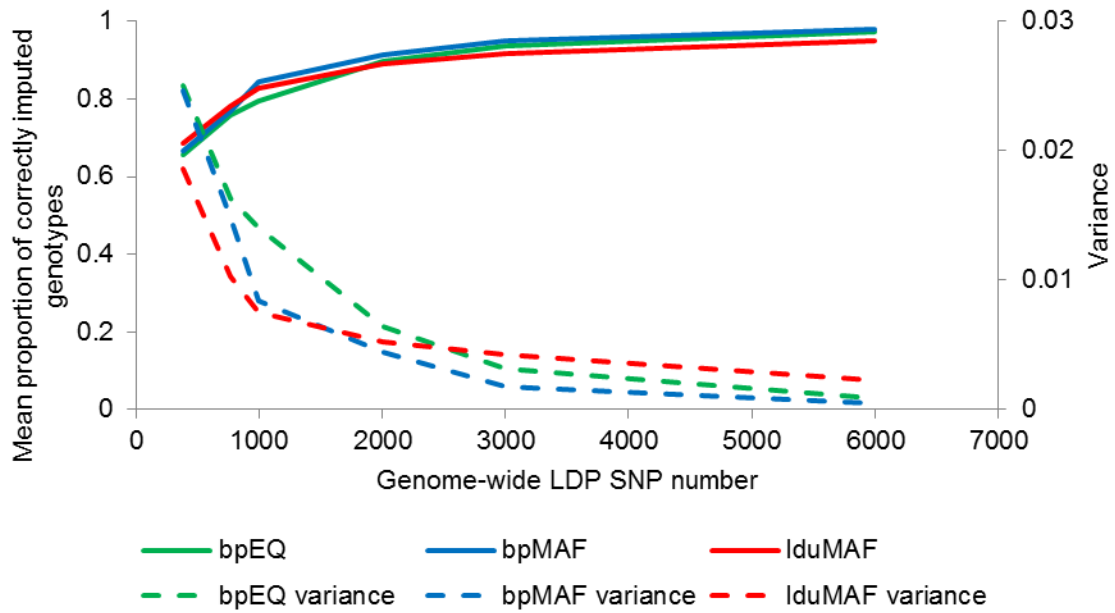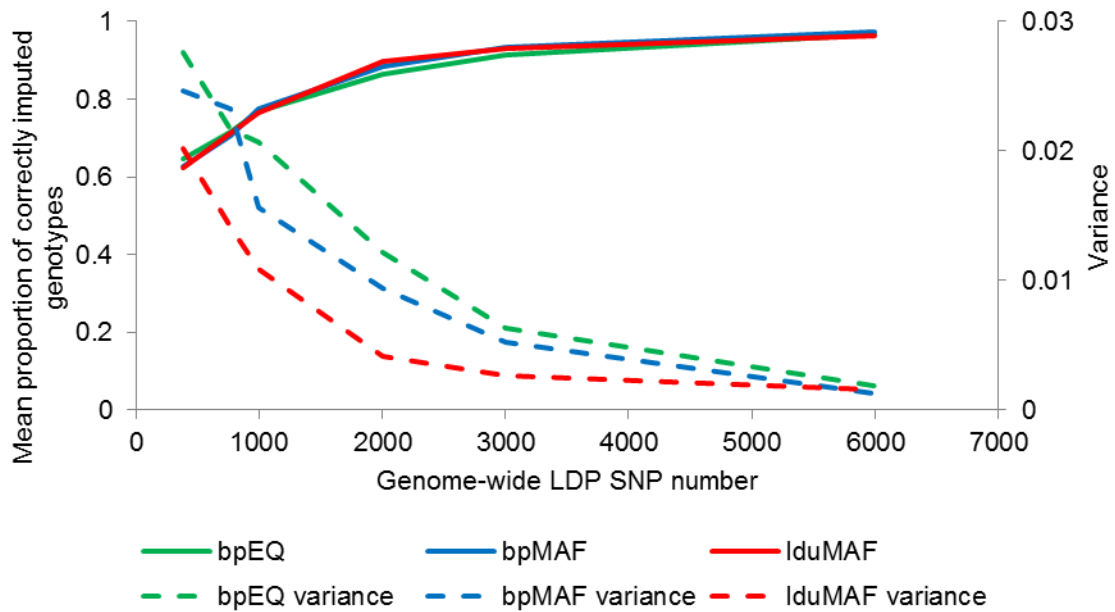**Figure S2** The mean proportion of correctly imputed genotypes and its variance across SNPs
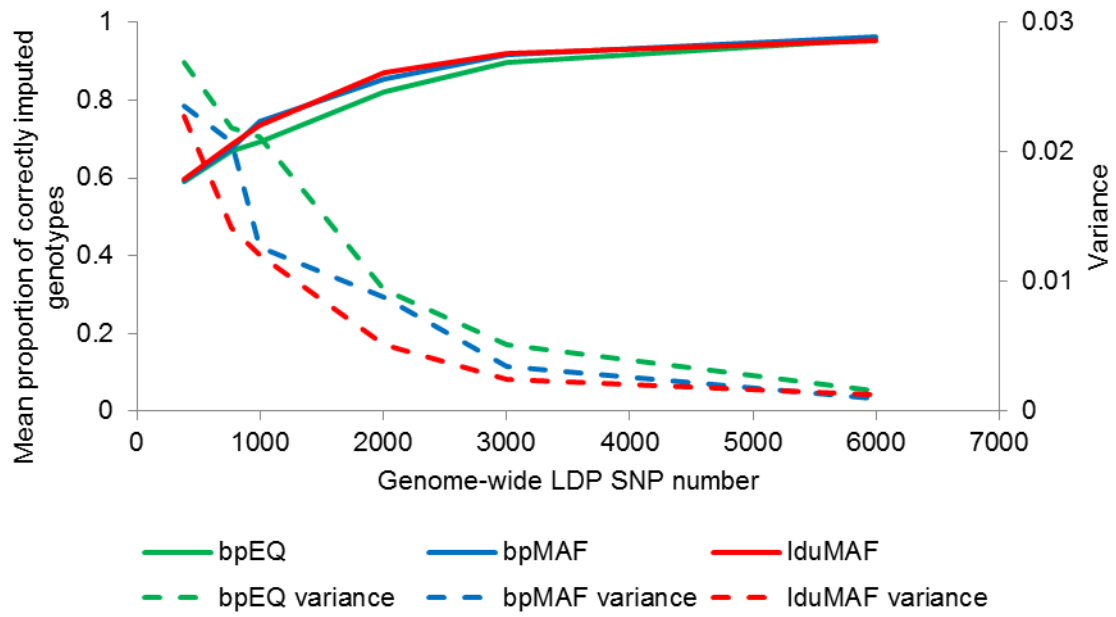
As calculated in the within-population analysis of the UK dataset and plotted against the total number of SNPs that would be on a genome-wide LDP of equivalent density.
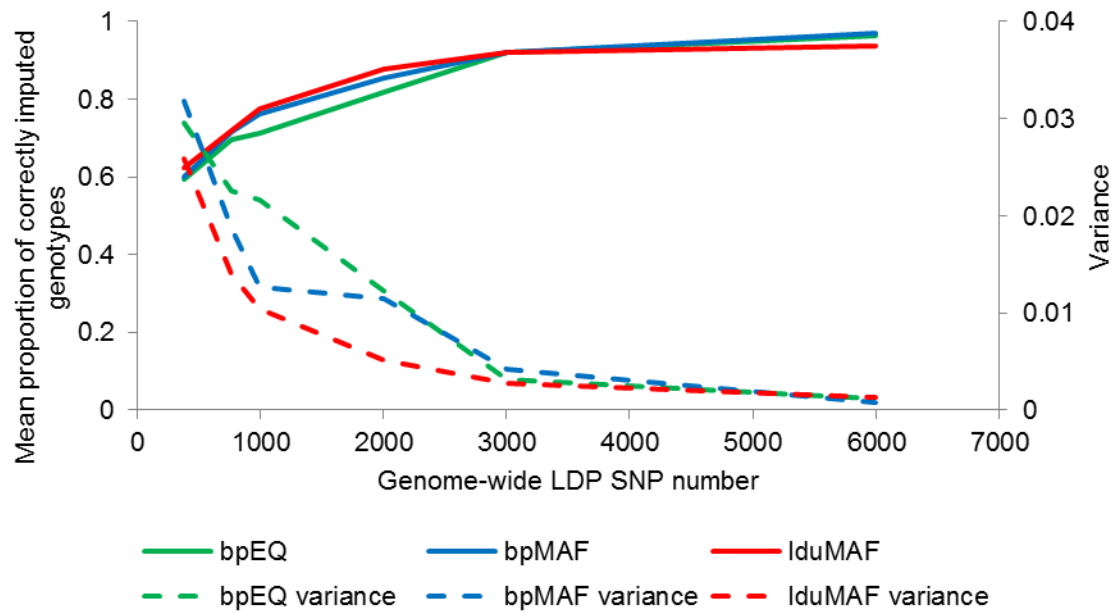

a) ECA1

b) ECA10

c) ECA20

d) ECA26