

Supplementary Table S3. Pearson correlations for 845 haemostatic gene variants with at least 1% MAF in at least 1 population.

	AMR	EAS	EUR	SAS	SOM	YRI
AMR		0.904	0.964	0.957	0.905	0.817
EAS	0.904		0.862	0.899	0.805	0.722
EUR	0.964	0.862		0.955	0.909	0.806
SAS	0.957	0.899	0.955		0.901	0.812
SOM	0.905	0.805	0.909	0.901		0.910
YRI	0.817	0.722	0.806	0.812	0.910	

