

Table S3 Genetic correlation matrix for sulcal lengths

Trait	arsp	Cs	iar	ips	lf	lu	ps	sar	spcd	sts
arsp	1.000	0.241	-1.000	0.113	-0.185	-0.287	-0.372	-0.507	0.328	-0.223
cs	0.241	1.000	-0.284	0.240	-0.272	0.432	0.185	-0.188	-0.322	-0.613
iar	-1.000	-0.284	1.000	-0.329	0.264	0.470	0.509	-0.251	1.000	-0.605
ips	0.113	0.240	-0.329	1.000	0.741	0.275	0.399	1.000	0.793	0.577
lf	-0.185	-0.272	0.264	0.741	1.000	0.443	0.255	-0.245	1.000	0.836
lu	-0.287	0.432	0.470	0.275	0.443	1.000	0.555	-0.719	0.083	1.000
ps	-0.372	0.185	0.509	0.399	0.255	0.555	1.000	0.118	0.571	0.267
sar	-0.507	-0.188	-0.251	1.000	-0.245	-0.719	0.118	1.000	-1.000	-0.337
spcd	0.328	-0.322	1.000	0.793	1.000	0.083	0.571	-1.000	1.000	1.000
sts	-0.223	-0.613	-0.605	0.577	0.836	1.000	0.267	-0.337	1.000	1.000

Averaged across both hemispheres, created from the extended baboon pedigree.