

S4 Table. LDSC based SNP heritability of gestational duration and SPTB in the FinnGen GWAS and in the meta-analysis.

Gestational duration					
population	N_preterm	N_term	N_total	sample_prevalence	H² (se)
FinnGenR6*	2250	22141	24391	0,09224714	0.1263 (0.0593)
Finland	286	488	774	0,369509044	
23andMe	3331	40237	43568	0,076455196	
Meta-analysis	5867	62866	68733	0,085359289	0.1747 (0.0266)

SPTB Case-Control					
population	N_preterm	N_term	N_total	sample_prevalence	H² (se)
FinnGenR6*	4925	49105	54030	0,091153063	0.0181 (0.0248)
Finland	286	488	774	0,369509044	
23andMe	3331	40236	43567	0,076456951	
Meta-analysis	8542	89829	98371	0,086834535	0.0565 (0.0141)