## S2 Table. GREML estimates of SNP heritability and genetic correlation across sexes.

Phenotype	N			Univariate estimates SNP heritability $^1$			Bivariate estimates genetic correlation <sup>1,2</sup>
1	Females	Males	Total	Females	Males	$Average^3$	Females-Males
Height	11,553	8,905	20,458	43.2% (3.0%) ***	45.1% (3.8%) ***	44.0%	0.981 (0.067) ***
BMĬ	11,542	8,907	20,449	22.1% (2.9%) ***	23.8% (3.8%) ***	22.8%	0.794 (0.122) *** †
EduYears	11,653	8,966	20,619	18.1% (2.9%) ***	18.9% (3.7%) ***	18.4%	0.832 (0.162) ***
CurrCigt	11,706	8,980	20,686	22.3% (7.1%) ***	26.7% (9.1%) ***	24.2%	0.543 (0.257) *** †
CurrDrinkFreq	11,312	8,760	20,072	14.1% (4.6%) ***	0.9% (6.0%)	8.3%	1.000 (2.068) *
Self-rated health	10,866	8,318	19,184	8.6% (3.1%) ***	10.8% (4.0%) ***	9.5%	1.000 (0.349) ***
<sup>1</sup> Standard errors between parentheses.					* > 0 at 10	% sign.	† < 1 at 10% sign.
<sup>2</sup> Significance of deviations from one only tested					** > 0  at  5	5% sign.	†† < 1 at 5% sign.
for genetic correlations.					*** > 0 at 1	l% sign.	$\dagger \dagger \dagger < 1 \text{ at } 1\% \text{ sign.}$

for genetic correlations.  $^3$  Sample-size weighted averages of univariate estimates in females and males.

S2 Table 1/1