

An independent validation study of three single nucleotide polymorphisms at the sex hormone-binding globulin locus for testosterone levels identified by genome-wide association studies

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The authors would like to apologise for three mistakes that were in the originally published version of this article.

The mistakes are as follows:

1. Page 2, line 5: The statistics value of rs6259 and rs727428 were reversed.

Mistake: rs6259 and rs727428 were not ($\beta_{STD} = 0.082$, $P = 0.017$; $\beta_{STD} = 0.17$, $P = 0.071$, respectively),

Correctly: rs6259 and rs727428 were not ($\beta_{STD} = 0.17$, $P = 0.071$; $\beta_{STD} = 0.082$, $P = 0.017$, respectively),

2. Page 3, right paragraph line 15: The fragment size was mistake.

Mistake: 275 bp (A-allele)

Correct: 274 bp (A-allele)

3. Table II: Three β_{STD} were incorrect.

Mistake:

SNP (Effect allele)	Trait	Cohort 1		Cohort 2	
		β_{STD} (SE)	P	β_{STD} (SE)	P
rs2075230 (A)	SHBG	0.020 (0.044)	6.5×10^{-6}	0.025 (0.047)	1.3×10^{-7}
	cFT	-0.097 (0.047)	0.039	0.034 (0.050)	0.50
rs6259 (A)	SHBG	0.20 (0.072)	6.4×10^{-3}	0.025 (0.070)	3.0×10^{-4}

Correct:

SNP (Effect allele)	Trait	Cohort 1		Cohort 2	
		β_{STD} (SE)	P	β_{STD} (SE)	P
rs2075230 (A)	SHBG	0.20 (0.044)	6.5×10^{-6}	0.25 (0.047)	1.3×10^{-7}
	cFT	-0.097 (0.047)	0.039	0.034 (0.050)	0.50
rs6259 (A)	SHBG	0.20 (0.072)	6.4×10^{-3}	0.25 (0.070)	3.0×10^{-4}