

Supplementary Table 3. Genomewide Association results for BMI, HIP, WST, and WHR-associated SNPs with p-value $< 5 \times 10^{-8}$ using Meta-Random

Trait	Chr	Position	Minor	Major	MAF	Beta	SE	p(Meta-Random)
BMI	5	171275948	A	ATTT	0.4496	0.2472	0.0483	1.33×10^{-8}
	16	53800954	C	T	0.1258	0.3622	0.0681	1.61×10^{-8}
	16	53803574	A	T	0.1258	0.3558	0.0681	2.11×10^{-8}
	16	53802494	T	C	0.1263	0.3569	0.0683	2.42×10^{-8}
	16	53813367	G	T	0.1252	0.3551	0.0683	2.87×10^{-8}
	16	53803223	A	G	0.1257	0.3521	0.0682	3.07×10^{-8}
	5	171276393	G	A	0.4064	0.2359	0.0494	4.74×10^{-8}
	16	53816752	G	A	0.1242	0.3463	0.0689	4.85×10^{-8}
	16	53816838	A	G	0.1241	0.3459	0.0689	4.99×10^{-8}
WHR	12	111518493	G	A	0.1356	-0.0023	0.0004	5.59×10^{-9}
	12	111414461	G	T	0.1445	-0.0020	0.0004	7.90×10^{-9}
	12	111386127	C	T	0.1420	-0.0020	0.0004	1.21×10^{-8}
	12	111389437	G	C	0.1429	-0.0019	0.0004	1.77×10^{-8}
	12	111367244	T	C	0.1386	-0.0020	0.0004	2.45×10^{-8}
	12	111390454	T	TAGAA	0.1461	-0.0019	0.0004	3.89×10^{-8}

BMI, body mass index; HIP, hip circumference; WST, waist circumference; WHR, waist-hip ratio; SNP, single nucleotide polymorphism; LMM, linear mixed model; MAF, minor allele frequency; SE, standard error.