

**Table S6. Results of the GWAS and the replication study for loci previously studied with European populations.**

SNP	Chr	Position (hg19)	Coded allele	GWAS set			Replication set			Combined			Ref. <sup>†</sup>
				Allele freq.	Effect (s.e.)	P	Allele freq.	Effect (s.e.)	P	Allele freq.	Effect (s.e.)	P	
rs215605	7	32,336,965	T	0.31	-0.23 (0.22)	0.29	0.31	0.057 (0.29)	0.85	0.31	-0.13 (0.18)	0.47	[4]
rs215614	7	32,347,335	G	0.66	0.13 (0.22)	0.54	0.67	0.097 (0.29)	0.74	0.66	0.12 (0.17)	0.49	[4]
rs6474412	8	42,550,498	T	0.85	0.60 (0.30)	0.042	0.86	0.26 (0.4)	0.51	0.85	0.48 (0.24)	0.043	[4]
rs13280604	8	42,559,586	G	0.14	-0.56 (0.31)	0.073	0.13	-0.27 (0.4)	0.49	0.14	-0.45 (0.25)	0.066	[4]
rs1329650	10	93,348,120	T	0.76	0.07 (0.24)	0.77	0.76	-0.65 (0.32)	0.042	0.76	-0.20 (0.19)	0.31	[6]
rs1028936	10	93,349,797	C	0.62	-0.29 (0.22)	0.19	0.61	-0.18 (0.28)	0.51	0.61	-0.25 (0.17)	0.15	[6]
rs588765	15	78,865,425	T	0.12	-0.097 (0.32)	0.76	0.11	0.019 (0.42)	0.96	0.12	-0.055 (0.25)	0.83	[5]
rs16969968	15	78,882,925	G	0.98	-0.56 (0.71)	0.43	0.98	-1.50 (0.86)	0.088	0.98	-0.93 (0.55)	0.091	[5,6]
rs1051730	15	78,894,339	G	0.98	-0.56 (0.71)	0.43	0.98	-1.50 (0.86)	0.088	0.98	-0.93 (0.55)	0.090	[4-6]
rs6495308	15	78,907,656	T	0.22	0.058 (0.25)	0.81	0.22	0.69 (0.32)	0.031	0.22	0.29 (0.20)	0.13	[5]
rs7937	19	41,302,706	T	0.64	0.19 (0.21)	0.37	0.64	0.13 (0.28)	0.65	0.64	0.17 (0.17)	0.32	[4]
rs3733829	19	41,310,571	G	0.39	0.72 (0.21)	0.00065	0.39	0.66 (0.28)	0.018	0.39	0.7 (0.17)	$3.3 \times 10^{-5}$	[6]
rs1801272	19	41,354,533	T	0.012	-11 (2.40)	$1.4 \times 10^{-5}$	0.012	-6.5 (3.70)	0.082	0.012	-9.4 (2.0)	$4.5 \times 10^{-6}$	[4]
rs7260329	19	41,521,638	G	0.56	0.30 (0.21)	0.15	0.56	1.1 (0.27)	$9.4 \times 10^{-5}$	0.56	0.59 (0.17)	0.00041	[4]
rs4105144	19	41,358,624	-	-	-	-	-	-	-	-	-	-	[4]

<sup>†</sup> previous reports provided in the main text.