

ESM Table 9 Heterogeneity in allelic effects between Chinese, East Asians and Europeans

CHR	SNP	Nearby Genes	R/A	OR[95%CI]					P for heterogeneity		Reported Population
				CKB [Up to 7,109 cases and 86,022 controls]	AGEN-T2D [Up to 25,079 cases and 29,611 controls]	Meta-analysis of CKB+AGEN-T2D ^a [up to 32,188 cases and 115,633 controls]	^b DIAGRAMv3 (EU) [up to 12,171 cases and 56,862 controls]	Meta-analysis of CKB+Trans-ethnic GWAS ^c [up to 33,597 cases and 201,655 controls]	CKB vs. DIAGRAMv3	Meta vs. DIAGRAMv3	
1	rs10923931	NOTCH2	T/G	1.16[1.05-1.27]	1.00[0.86-1.16]	1.11[1.02-1.20]	1.10[1.05-1.16]	1.06[1.03-1.10]	3.93x10 ⁻¹	9.17x10 ⁻¹	EU
1	rs340874	PROX1	C/T	1.05[1.01-1.09]	1.08[1.03-1.14]	1.06[1.03-1.09]	1.08[1.04-1.12]	1.06[1.04-1.08]	3.24x10 ⁻¹	5.05x10 ⁻¹	EU
2	rs780094	GCKR	C/T	1.08[1.04-1.12]	1.06[1.01-1.11]	1.07[1.04-1.10]	1.04[1.00-1.08]	1.07[1.04-1.09]	1.41x10 ⁻¹	1.90x10 ⁻¹	EU
2	rs7578597	THADA	T/C	1.27[1.01-1.60]	0.93[0.62-1.40]	1.18[0.96-1.44]	1.14[1.08-1.22]	1.08[1.03-1.13]	4.14x10 ⁻¹	8.18x10 ⁻¹	EU
2	rs243021	BCL11A	A/G	1.07[1.03-1.11]	1.05[1.00-1.10]	1.06[1.03-1.09]	1.09[1.05-1.13]	1.07[1.05-1.09]	5.23x10 ⁻¹	2.98x10 ⁻¹	EU
2	rs7593730	RBMS1	C/T	0.98[0.93-1.02]	1.00[0.94-1.07]	0.99[0.95-1.02]	1.11[1.06-1.15]	1.03[1.00-1.06]	1.22x10 ⁻⁴	7.05x10 ⁻⁵	* EU
2	rs3923113	GRB14	A/C	1.00[0.95-1.05]	1.03[0.95-1.12]	1.01[0.97-1.05]	1.04[1.00-1.08]	1.05[1.02-1.08]	2.51x10 ⁻¹	3.26x10 ⁻¹	SA
2	rs2943641	IRS1	C/T	1.04[0.97-1.11]	1.12[1.03-1.22]	1.07[1.02-1.13]	1.08[1.04-1.12]	1.08[1.05-1.12]	3.55x10 ⁻¹	8.21x10 ⁻¹	EU
3	rs1801282	PPARG	C/G	1.07[0.99-1.16]	1.15[1.01-1.30]	1.09[1.02-1.16]	1.16[1.11-1.22]	1.12[1.08-1.17]	6.74x10 ⁻²	1.14x10 ⁻¹	EU
3	rs6780569	UBE2E2	G/A	1.11[1.06-1.16]	1.17[1.12-1.22]	1.14[1.10-1.18]	1.09[1.02-1.15]	1.11[1.08-1.14]	4.85x10 ⁻¹	1.40x10 ⁻¹	EA
3	rs831571	PSMD6	C/T	1.06[1.02-1.10]	1.09[1.05-1.13]	1.08[1.05-1.11]	1.03[0.99-1.08]	1.05[1.03-1.08]	3.78x10 ⁻¹	1.13x10 ⁻¹	EA
3	rs4607103	ADAMTS9	C/T	1.00[0.97-1.04]	0.99[0.95-1.04]	1.00[0.97-1.03]	1.08[1.04-1.12]	1.03[1.00-1.05]	5.77x10 ⁻³	1.27x10 ⁻³	EU
3	rs11708067	ADCY5	A/G	1.92[1.28-2.88]	1.18[0.80-1.74]	1.49[1.13-1.97]	1.10[1.06-1.15]	1.11[1.07-1.15]	7.53x10 ⁻³	3.75x10 ⁻²	EU
3	rs1470579	IGF2BP2	C/A	1.11[1.07-1.16]	1.15[1.11-1.19]	1.13[1.10-1.16]	1.12[1.08-1.16]	1.12[1.10-1.15]	7.97x10 ⁻¹	6.38x10 ⁻¹	EU
3	rs16861329	ST6GAL1	C/G	1.04[1.00-1.09]	0.92[0.86-0.99]	1.01[0.97-1.04]	1.03[0.97-1.09]	1.07[1.04-1.10]	7.37x10 ⁻¹	5.31x10 ⁻¹	SA
4	rs6815464	MAEA	C/G	1.08[1.04-1.12]	1.13[1.10-1.16]	1.11[1.07-1.13]	1.12[1.01-1.24]	1.05[1.02-1.08]	4.94x10 ⁻¹	8.85x10 ⁻¹	EA
4	rs10010131	WFS1	G/A	1.04[0.94-1.15]	1.00[0.91-1.10]	1.02[0.95-1.09]	1.10[1.06-1.14]	1.09[1.06-1.12]	3.06x10 ⁻¹	5.48x10 ⁻²	EU
5	rs4457053	ZBED3	G/A	1.10[1.02-1.18]	1.00[0.85-1.18]	1.08[1.01-1.16]	1.13[1.08-1.19]	1.10[1.06-1.14]	4.79x10 ⁻¹	2.57x10 ⁻¹	EU
6	rs7754840	CDKAL1	C/G	1.21[1.07-1.26]	1.18[1.14-1.22]	1.20[1.17-1.23]	1.15[1.11-1.19]	1.17[1.15-1.20]	3.06x10 ⁻²	6.39x10 ⁻²	EU
6	rs9470794	ZFAND3	C/T	1.02[0.98-1.05]	1.12[1.08-1.16]	1.07[1.04-1.10]	0.99[0.93-1.05]	1.04[1.01-1.07]	4.49x10 ⁻¹	2.47x10 ⁻²	EA
7	rs2191349	DGKB	T/G	1.05[1.00-1.09]	1.12[1.08-1.17]	1.09[1.06-1.12]	1.07[1.03-1.11]	1.08[1.05-1.10]	4.35x10 ⁻¹	5.15x10 ⁻¹	EU
7	rs864745	JAZF1	T/C	1.04[1.00-1.09]	1.06[1.00-1.12]	1.05[1.01-1.09]	1.12[1.08-1.15]	1.08[1.06-1.11]	1.51x10 ⁻²	1.04x10 ⁻²	EU
7	rs4607517	GCK	A/G	1.01[0.97-1.06]	1.03[0.97-1.09]	1.02[0.98-1.05]	1.05[1.00-1.10]	1.03[1.00-1.05]	2.86x10 ⁻¹	3.24x10 ⁻¹	EU
7	rs6467136	GCC1-PAX4	G/A	1.04[1.00-1.09]	1.11[1.07-1.14]	1.08[1.05-1.11]	0.99[0.95-1.02]	1.02[1.00-1.04]	5.13x10 ⁻²	2.28x10 ⁻⁵	* EA
7	rs972283	KLF14	G/A	1.04[1.00-1.08]	0.99[0.93-1.06]	1.03[0.99-1.06]	1.10[1.06-1.14]	1.05[1.02-1.07]	4.42x10 ⁻²	7.98x10 ⁻³	EU
8	rs896854	TP53INP1	T/C	1.04[1.00-1.08]	1.07[1.02-1.12]	1.05[1.02-1.08]	1.09[1.05-1.13]	1.07[1.04-1.09]	7.94x10 ⁻²	1.33x10 ⁻¹	EU
8	rs13266634	SLC30A8	C/T	1.10[1.06-1.13]	1.10[1.07-1.14]	1.10[1.07-1.13]	1.16[1.11-1.22]	1.12[1.10-1.15]	4.51x10 ⁻²	3.98x10 ⁻²	EU
9	rs7041847	GLIS3	A/G	1.07[1.03-1.10]	1.10[1.07-1.13]	1.09[1.06-1.11]	1.05[1.01-1.09]	1.06[1.04-1.08]	5.36x10 ⁻¹	1.14x10 ⁻¹	EA
9	rs17584499	PTPRD	T/C	1.00[0.95-1.06]	1.09[1.00-1.19]	1.03[0.98-1.08]	1.00[0.95-1.06]	1.01[0.97-1.04]	9.80x10 ⁻¹	5.03x10 ⁻¹	EA
9	rs10811661	CDKN2A/B	T/C	1.22[1.18-1.26]	1.12[1.07-1.16]	1.17[1.14-1.21]	1.18[1.13-1.23]	1.21[1.18-1.24]	2.36x10 ⁻¹	8.50x10 ⁻¹	EU
9	rs13292136	TLE4/CHCHD9	C/T	1.08[1.01-1.14]	0.99[0.92-1.07]	1.04[0.99-1.09]	1.19[1.11-1.27]	1.10[1.07-1.14]	3.04x10 ⁻²	1.84x10 ⁻³	EU
10	rs10906115	CDC123	A/G	1.08[1.05-1.12]	1.08[1.05-1.13]	1.08[1.06-1.11]	1.06[1.02-1.10]	1.08[1.05-1.10]	3.72x10 ⁻¹	3.07x10 ⁻¹	EA
10	rs1802295	VPS26A	T/C	1.03[0.97-1.08]	1.01[0.94-1.09]	1.02[0.98-1.07]	1.02[0.98-1.06]	1.04[1.02-1.07]	8.68x10 ⁻¹	9.68x10 ⁻¹	SA
10	rs1111875	HHEX/IDE	C/T	1.11[1.07-1.15]	1.08[1.04-1.13]	1.10[1.07-1.13]	1.15[1.11-1.18]	1.12[1.09-1.14]	2.41x10 ⁻¹	5.15x10 ⁻²	EU
10	rs7901695	TCF7L2	C/T	1.37[1.25-1.50]	1.18[1.03-1.35]	1.31[1.21-1.41]	1.37[1.32-1.42]	1.32[1.28-1.35]	9.80x10 ⁻¹	2.90x10 ⁻¹	EU
10	rs10886471	GRK5	C/T	1.00[0.96-1.05]	1.06[0.99-1.13]	1.02[0.98-1.06]	0.99[0.95-1.03]	1.01[0.98-1.03]	6.95x10 ⁻¹	2.96x10 ⁻¹	EA
11	rs4752781	DUSP8/INS	T/A	0.99[0.95-1.04]	1.04[0.98-1.10]	1.01[0.97-1.05]	1.04[1.01-1.08]	1.03[1.00-1.05]	7.73x10 ⁻²	1.81x10 ⁻¹	EU
11	rs2237892	KCNQ1	C/T	1.25[1.20-1.30]	1.19[1.14-1.24]	1.22[1.19-1.26]	1.12[1.04-1.21]	1.23[1.20-1.27]	1.17x10 ⁻²	3.88x10 ⁻²	EA
11	rs5215	KCNJ11	C/T	1.07[1.04-1.11]	1.10[1.06-1.14]	1.09[1.06-1.11]	1.08[1.05-1.12]	1.08[1.06-1.10]	7.05x10 ⁻¹	9.17x10 ⁻¹	EU
11	rs1552224	ARAP1	A/C	1.09[1.02-1.16]	1.16[1.05-1.28]	1.11[1.05-1.17]	1.13[1.08-1.19]	1.10[1.06-1.14]	3.18x10 ⁻¹	5.50x10 ⁻¹	EU
11	rs10830963	MTNR1B	G/C	1.02[0.99-1.06]	1.00[0.93-1.08]	1.02[0.99-1.05]	1.11[1.07-1.16]	1.06[1.03-1.08]	1.81x10 ⁻³	6.95x10 ⁻⁴	EU
12	rs1531343	HMG2A	C/G	1.05[0.99-1.11]	1.06[0.99-1.14]	1.05[1.01-1.10]	1.15[1.09-1.22]	1.08[1.05-1.12]	2.02x10 ⁻²	1.40x10 ⁻²	EU
12	rs7961581	TSPAN8/LGR5	C/T	1.04[1.00-1.08]	1.01[0.95-1.06]	1.03[0.99-1.06]	1.08[1.04-1.12]	1.05[1.03-1.07]	1.88x10 ⁻¹	4.65x10 ⁻²	EU
13	rs1359790	SPRY2	G/A	1.06[1.02-1.10]	1.05[1.01-1.10]	1.06[1.03-1.09]	1.10[1.06-1.14]	1.07[1.04-1.09]	2.06x10 ⁻¹	9.87x10 ⁻²	EA
15	rs7403531	RASGRP1	T/C	1.03[0.99-1.07]	1.08[1.02-1.13]	1.05[1.02-1.08]	1.02[0.98-1.06]	1.02[1.00-1.05]	6.23x10 ⁻¹	2.81x10 ⁻¹	EA
15	rs7172432	VPS13C	A/G	1.07[1.03-1.11]	1.11[1.07-1.15]	1.09[1.06-1.11]	1.06[1.02-1.10]	1.07[1.05-1.09]	7.82x10 ⁻¹	2.68x10 ⁻¹	EA
15	rs7178572	HMG20A	G/A	1.07[1.04-1.11]	1.09[1.04-1.14]	1.08[1.05-1.11]	1.08[1.04-1.12]	1.08[1.06-1.11]	8.48x10 ⁻¹	9.90x10 ⁻¹	SA
15	rs11634397	ZFAND6	G/A	1.02[0.96-1.09]	1.00[0.90-1.11]	1.02[0.97-1.07]	1.09[1.05-1.13]	1.06[1.03-1.09]	9.09x10 ⁻²	3.89x10 ⁻²	EU
15	rs2028299	AP3S2	C/A	1.06[1.02-1.11]	1.08[1.02-1.14]	1.07[1.03-1.11]	1.04[1.00-1.09]	1.07[1.04-1.09]	5.81x10 ⁻¹	4.15x10 ⁻¹	SA
15	rs8042680	PRCI	A/C	0.88[0.74-1.05]	1.64[1.16-2.32]	1.00[0.85-1.16]	1.07[1.04-1.11]	1.06[1.04-1.09]	2.59x10 ⁻²	3.50x10 ⁻¹	EU
16	rs9939609	FTO	A/T	1.15[1.09-1.21]	1.13[1.07-1.18]	1.14[1.10-1.18]	1.11[1.07-1.14]	1.11[1.08-1.14]	1.68x10 ⁻¹	2.12x10 ⁻¹	EU
17	rs4523957	SRR	T/G	0.98[0.94-1.02]	1.03[0.97-1.09]	0.99[0.96-1.03]	1.00[0.96-1.04]	1.00[0.98-1.03]	4.74x10 ⁻¹	8.24x10 ⁻¹	EA
17	rs4430796	HNF1B	G/A	1.09[1.05-1.14]	1.12[1.05-1.19]	1.10[1.07-1.14]	1.13[1.07-1.19]	1.10[1.07-1.13]	3.54x10 ⁻¹	4.30x10 ⁻¹	EU
18	rs12970134	MC4R	A/G	1.06[1.02-1.11]	1.07[1.02-1.12]	1.07[1.03-1.10]	1.08[1.04-1.12]	1.07[1.05-1.10]	5.94x10 ⁻¹	6.49x10 ⁻¹	EU
20	rs6017317	HNF4A	G/T	1.05[1.01-1.08]	1.09[1.07-1.12]	1.08[1.06-1.10]	1.06[1.01-1.12]	1.06[1.04-1.08]	6.31x10 ⁻¹	5.75x10 ⁻¹	EA
23	rs5945326	DUSP9	A/G	1.11[1.07-1.15]	NA	NA	NA	NA	NA	NA	EU

Meta: Meta-analysis of AGEN-T2D and CKB study; EU: Europeans; SA: South Asians; EA: East Asians; CKB: China Kadoorie Biobank

^aAGEN-T2D: T2D-GWAS meta-analysis in East Asian ; ^bDIAGRAM: DIAbetes Genetics Replication And Meta-analysis

^cTrans-ethnic GWAS: Trans-ethnic Meta analysis of four ethnic groups (East Asians[AGEN-T2D], Europeans [DIAGRAM], South Asians [SAT2D], and Mexicans [MAT2D])

Reference(PMID): ^a22158537; ^b22885922; ^c24509480; * SNPs showing nominal evidence of heterogeneity ($p < 0.05/110 = 0.00045$)