

9p21.3	rs1011970	9	22062134	T/G	1.03	(0.99-1.08)	0.145	0.999	1.06	(1.05-1.08)	6.4×10-16	4.1E-12	1.07	<i>CDKN2A,CDKN2B</i>	
9q31.2	rs10759243	9	110306115	A/C	1.05	(1.02-1.08)	2.9×10 ⁻⁴	0.658	1.06	(1.05-1.07)	3.4×10-21	1.1E-17	1.06		
9q31.2	rs10816625	9	110837073	A/G	0.91	(0.89-0.94)	1.6×10 ⁻¹⁰	1.1E-06	0.90	(0.89-0.92)	7.7×10-27	2.6E-23	0.90		
9q31.2	rs13294895	9	110837176	T/C	1.03	(0.94-1.13)	0.562	1.000	1.07	(1.05-1.09)	7.1×10-17	2.4E-13	1.07	<i>ASTN2</i>	
9q31.2	rs676256	9	110895353	T/C	1.05	(0.98-1.12)	0.194	0.999	1.10	(1.09-1.12)	2.8×10-53	9.5E-50	1.10		
9q33.1	rs1895062	9	119313486	A/G	1.01	(0.98-1.04)	0.476	1.000	1.04	(1.03-1.06)	1.1×10-13	7.4E-10	1.05		
9q33.3	rs10760444	9	129396434	A/G	0.93	(0.91-0.96)	1.0×10 ⁻⁷	6.7E-04	0.96	(0.95-0.97)	8.3×10-14	5.3E-10	0.96		<i>LMX1B</i>
9q34.2	rs8176636	9	136151579	T/TGGTGCAGGCCG AGGAAAAAATTTGTG GCAATTCCTCA	1.04	(1.01-1.08)	0.014	0.989	1.05	(1.03-1.06)	4.9×10-10	3.2E-06	1.05		<i>ABO</i>
10p15.1	rs2380205	10	5886734	T/C	0.97	(0.93-1.01)	0.135	0.999	0.98	(0.97-0.99)	5.3×10-05	2.6E-01	0.98	<i>ANKRD16</i>	
10p14	rs67958007	10	9088113	T/TG	1.01	(0.96-1.08)	0.638	1.000	1.06	(1.04-1.08)	4.5×10-10	3.1E-06	1.06	<i>DNAJC1</i>	
10p12.31	rs7072776	10	22032942	A/G	1.01	(0.95-1.08)	0.677	1.000	1.06	(1.05-1.08)	2.4×10-19	8.1E-16	1.06		
10p12.31	rs11814448	10	22315843	A/C	0.90	(0.77-1.05)	0.179	0.999	0.84	(0.80-0.87)	4.0×10-18	1.3E-14	0.83	<i>DNAJC1</i>	
10q21.2	rs10995201	10	64299890	A/G	1.08	(0.99-1.18)	0.100	0.998	1.14	(1.12-1.16)	5.1×10-52	1.7E-48	1.14	<i>ZNF365</i>	
10q22.3	rs704010	10	80841148	T/C	1.05	(1.03-1.08)	2.0×10 ⁻⁴	0.575	1.08	(1.06-1.09)	3.4×10-38	1.1E-34	1.08	<i>ZMZ1</i>	
10q23.33	rs140936696	10	95292187	CAA/C	1.02	(0.99-1.05)	0.192	0.999	1.04	(1.03-1.06)	5.4×10-08	3.6E-04	1.05	<i>TCFL2</i>	
10q25.2	rs7904519	10	114773927	A/G	0.98	(0.91-1.05)	0.529	1.000	0.96	(0.94-0.97)	1.8×10-13	1.2E-09	0.96		
10q26.12	rs11199914	10	123093901	T/C	0.97	(0.94-0.99)	0.019	0.992	0.96	(0.95-0.97)	4.2×10-13	2.9E-09	0.96		
10q26.13	rs2981578	10	123340311	T/C	0.83	(0.80-0.85)	6.5×10 ⁻⁴⁶	2.2E-42	0.81	(0.81-0.82)	2.3×10-291	7.6E-288	0.81		<i>FGFR2</i>
10q26.13	rs45631563	10	123349324	A/T	0.95	(0.77-1.17)	0.617	1.000	1.21	(1.18-1.25)	1.1×10-35	3.7E-32	1.22	<i>FGFR2</i>	
11p15	rs6597981	11	803017	A/G	0.98	(0.95-1.01)	0.253	0.999	0.96	(0.95-0.97)	2.5×10-12	1.8E-08	0.96	<i>PIDD1</i>	
11p15.5	rs3817198	11	1909006	T/C	0.94	(0.91-0.98)	0.002	0.922	0.94	(0.93-0.95)	5.5×10-21	1.8E-17	0.94	<i>LSP1</i>	
11q13.1	rs3903072	11	65583066	T/G	0.98	(0.95-1.02)	0.351	1.000	0.96	(0.95-0.97)	5.7×10-12	3.8E-08	0.96	<i>CCND1</i>	
11q13.3	rs75915166	11	69379161	A/C	1.26	(1.00-1.60)	0.054	0.997	1.30	(1.27-1.33)	1.3×10-96	4.3E-93	1.30		
11q22.3	rs11374964	11	108345515	G/GA	1.02	(0.99-1.04)	0.277	0.999	1.00	(0.99-1.01)	6.5×10-01	1.0E+00	1.00		<i>KDEL2C</i>
11q24.3	rs11820646	11	129461171	T/C	0.97	(0.95-1.00)	0.025	0.994	0.96	(0.95-0.97)	3.3×10-15	2.2E-11	0.95		
12p13.1	rs12422552	12	14413931	C/G	1.08	(1.05-1.11)	6.0×10 ⁻⁷	4.0E-03	1.06	(1.05-1.07)	2.4×10-20	7.9E-17	1.06	<i>NTN4</i>	
12p11.22	rs7297051	12	28174817	T/C	0.89	(0.86-0.91)	3.9×10 ⁻¹⁴	2.6E-10	0.89	(0.87-0.90)	5.5×10-72	1.8E-68	0.89		
12q21.31	rs202049448	12	85009437	T/C	0.99	(0.96-1.03)	0.711	1.000	1.03	(1.02-1.05)	8.5×10-07	5.6E-03	1.04	<i>TBX3</i>	
12q22	rs17356907	12	96027759	A/G	1.05	(1.02-1.08)	0.002	0.935	1.09	(1.07-1.10)	1.3×10-40	4.3E-37	1.09		
12q24.21	rs1292011	12	115836522	A/G	1.10	(1.07-1.13)	1.4×10 ⁻¹⁰	9.4E-07	1.09	(1.08-1.10)	1.0×10-47	3.4E-44	1.09	<i>PAX9</i>	
12q24.31	rs206966	12	120832146	T/C	1.02	(0.96-1.09)	0.496	1.000	1.05	(1.03-1.07)	5.0×10-08	3.4E-04	1.05		
13q22.1	rs6562760	13	73957681	A/G	0.98	(0.81-1.20)	0.881	1.000	0.96	(0.94-0.97)	1.3×10-09	8.9E-06	0.96	<i>RAD51B</i>	
14q13.3	rs2236007	14	37132769	A/G	0.93	(0.91-0.96)	6.4×10 ⁻⁶	0.041	0.93	(0.92-0.94)	1.1×10-25	3.6E-22	0.93		
14q24.1	rs2588809	14	68660428	T/C	1.08	(1.00-1.17)	0.058	0.997	1.07	(1.05-1.08)	1.3×10-14	8.6E-11	1.06	<i>RAD51B</i>	
14q24.1	rs999737	14	69034682	T/C	1.12	(0.86-1.46)	0.403	1.000	0.91	(0.90-0.92)	1.2×10-38	4.0E-35	0.91	<i>CCDC88C</i>	
14q32.11	rs941764	14	91841069	A/G	0.96	(0.93-1.00)	0.050	0.997	0.96	(0.94-0.97)	1.7×10-13	1.2E-09	0.95	<i>RIN3</i>	
14q32.12	rs11627032	14	93104072	T/C	1.02	(0.99-1.05)	0.307	1.000	1.04	(1.03-1.06)	2.0×10-10	1.3E-06	1.05		
14q32.33	rs10623258	14	105212261	CTT/C	1.01	(0.98-1.04)	0.565	1.000	1.03	(1.02-1.04)	7.9×10-08	5.4E-04	1.04	<i>ADSSL1</i>	
16p13.3	rs11076805	16	4106788	A/C	0.99	(0.94-1.04)	0.630	1.000	0.97	(0.96-0.98)	3.2×10-05	0.177	0.97	<i>ADCY9</i>	
16q12.2	rs17817449	16	53813367	T/G	1.07	(1.03-1.11)	6.2×10 ⁻⁴	0.805	1.06	(1.05-1.07)	5.4×10-24	1.8E-20	1.06	<i>FTO</i>	
16q12.2	rs11075995	16	53855291	A/T	1.05	(1.02-1.08)	9.2×10 ⁻⁴	0.861	1.04	(1.03-1.06)	3.2×10-11	2.2E-07	1.04	<i>FTO</i>	
16q12.2	rs28539243	16	54682064	A/G	1.03	(1.01-1.06)	0.018	0.992	1.05	(1.04-1.06)	1.1×10-15	7.2E-12	1.05	<i>AMFR</i>	
16q13	rs2432539	16	56420987	A/G	1.05	(1.02-1.08)	0.001	0.880	1.04	(1.03-1.05)	2.7×10-10	1.7E-06	1.04		
16q23.2	rs13329835	16	80650805	A/G	0.93	(0.88-0.99)	0.017	0.991	0.92	(0.91-0.94)	2.9×10-28	9.6E-25	0.92	<i>CDYL2</i>	
16q24.2	rs4496150	16	87085237	A/C	0.96	(0.94-0.99)	0.007	0.980	0.96	(0.95-0.97)	1.9×10-10	1.3E-06	0.96	<i>ATAD5</i>	
17q11.2	rs146699004	17	29230520	G/GGT	0.96	(0.91-1.01)	0.095	0.998	0.96	(0.95-0.97)	5.6×10-10	3.8E-06	0.96		
17q22	rs2787486	17	53209774	A/C	1.08	(1.05-1.12)	4.7×10 ⁻⁷	3.2E-03	1.08	(1.07-1.09)	2.1×10-34	7.0E-31	1.08		

17q25.3	rs745570	17	77781725	A/G	1.03	(1.00-1.06)	0.026	0.994	1.04	(1.03-1.05)	3.3×10 ⁻¹¹	2.1E-07	1.04	
18q11.2	rs527616	18	24337424	C/G	0.97	(0.94-1.00)	0.022	0.993	0.95	(0.94-0.96)	7.7×10 ⁻¹⁶	5.5E-12	0.95	
18q11.2	rs1436904	18	24570667	T/G	1.03	(1.01-1.06)	0.015	0.990	1.05	(1.04-1.06)	7.8×10 ⁻¹⁶	5.2E-12	1.05	<i>CHST9</i>
18q12.1	rs36194942	18	25401204	A/AT	1.01	(0.98-1.04)	0.468	1.000	1.03	(1.02-1.04)	2.7×10 ⁻⁰⁶	0.017	1.03	<i>CDH2</i>
18q12.1	rs117618124	18	29977689	T/C	0.93	(0.69-1.24)	0.604	1.000	1.11	(1.08-1.15)	1.1×10 ⁻¹¹	7.3E-08	1.11	<i>GAREM1</i>
18q12.3	rs6507583	18	42399590	A/G	1.06	(1.02-1.10)	0.006	0.974	1.08	(1.06-1.11)	9.7×10 ⁻¹⁴	6.6E-10	1.09	<i>SETBP1</i>
19p13.2	rs322144	19	11423703	C/G	1.01	(0.98-1.05)	0.488	1.000	1.02	(1.01-1.04)	2.1×10 ⁻⁰⁴	5.8E-01	1.03	<i>TSPAN16</i>
19p13.12	rs2594714	19	13954571	A/G	0.96	(0.93-0.99)	0.009	0.984	0.96	(0.95-0.97)	3.8×10 ⁻¹⁰	2.5E-06	0.96	
19p13.11	rs4808801	19	18571141	A/G	1.06	(1.03-1.09)	1.5×10 ⁻⁴	0.495	1.07	(1.06-1.08)	2.3×10 ⁻³¹	7.7E-28	1.07	<i>ELL</i>
19p13.11	rs2965183	19	19545696	A/G	1.02	(0.99-1.05)	0.180	0.999	1.04	(1.03-1.05)	9.8×10 ⁻¹²	6.7E-08	1.05	<i>GATAD2A, MIR640</i>
19q12	rs113701136	19	30277729	T/C	1.02	(0.99-1.06)	0.178	0.999	1.02	(1.01-1.03)	5.6×10 ⁻⁰⁴	0.785	1.02	<i>CCNE1</i>
19q13.31	rs3760982	19	44286513	A/G	0.99	(0.96-1.03)	0.746	1.000	1.05	(1.03-1.06)	1.6×10 ⁻¹⁴	1.1E-10	1.05	<i>KCCN4, LYPD5</i>
19q13.22	rs71338792	19	46183031	A/AT	0.96	(0.92-0.99)	0.022	0.993	0.95	(0.94-0.97)	2.7×10 ⁻¹⁰	1.8E-06	0.95	<i>GIPR</i>
20q11.22	rs2284378	20	32588095	T/C	0.99	(0.95-1.02)	0.392	1.000	1.01	(1.00-1.02)	8.8×10 ⁻⁰²	0.998	1.01	<i>RALY</i>
20q13.13	rs6122906	20	48945911	A/G	0.97	(0.94-1.00)	0.029	0.995	0.95	(0.94-0.97)	3.7×10 ⁻¹¹	2.4E-07	0.95	
21q21.1	rs2823093	21	16520832	A/G	0.94	(0.87-1.00)	0.056	0.997	0.94	(0.92-0.95)	1.7×10 ⁻²¹	5.6E-18	0.94	<i>NRIP1</i>
22q13.1	rs738321	22	38568833	C/G	1.03	(1.00-1.06)	0.087	0.998	1.05	(1.03-1.06)	7.2×10 ⁻¹⁴	4.8E-10	1.05	<i>PLA2G6</i>
22q13.1	rs6001930	22	40876234	T/C	0.95	(0.92-0.97)	2.0×10 ⁻⁴	0.580	0.90	(0.89-0.92)	5.4×10 ⁻³⁴	1.8E-30	0.89	<i>MKL1</i>
22q13.31	rs28512361	22	46283297	A/G	1.04	(0.95-1.14)	0.351	1.000	1.06	(1.04-1.08)	1.4×10 ⁻⁰⁸	9.4E-05	1.06	

^aOR literature: The ORs for risk variants identified in Asian populations were extracted from Zheng W et al. Nat Gen 2009, Cai Q et al. Hum Mol Genet. 2011, Long J et al. PLoS Genet. 2012, Cai Q et al. Nat Genet. 2014, and Han MR et al. Hum Mol Genet. 2016

; those identified in European populations were extracted from Michailidou K et al. Nature. 2017

^bFPRP: The false positive report probability