

Ancestry	Approach	CHR	SNP	BP	A1	A2	FRQ	INFO	Beta	SE	Pval
African											
	Baseline	2	rs17367970	14976634	A	G	0.82	0.97	0.21	0.04	1.88x10 ⁻⁶
		3	rs2639227	81209252	C	T	0.35	0.97	-0.18	0.04	6.73x10 ⁻⁷
		6	rs9463613	50575789	C	G	0.95	0.84	-0.40	0.08	1.70x10 ⁻⁶
		7	rs5018071	122362272	A	G	0.71	0.99	0.18	0.04	5.97x10 ⁻⁷
	Averaged	2	rs13394665	* 64244935	A	C	0.73	0.97	0.15	0.03	1.27x10 ⁻⁶
		2	rs1069256	* 173881973	C	T	0.75	1.02	0.15	0.03	1.36x10 ⁻⁶
		3	rs2639228	81208798	A	G	0.68	0.99	0.15	0.03	6.64x10 ⁻⁷
		6	rs7775991	47008467	C	T	0.73	1.02	0.14	0.03	4.78x10 ⁻⁶
		6	rs280312	50712854	G	T	0.95	1.02	-0.29	0.06	1.90x10 ⁻⁶
		7	rs17168027	* 134099287	A	G	0.89	0.89	0.21	0.05	4.38x10 ⁻⁶
		10	rs10825165	55354348	A	G	0.11	0.92	-0.22	0.05	3.11x10 ⁻⁶
		15	rs9806467	48965568	C	T	0.83	1.11	-0.16	0.03	3.81x10 ⁻⁶
	Repeated Measures	1	rs1188356	30983669	G	T	0.79	1.02	0.16	0.03	7.33x10 ⁻⁷
		1	rs1413227	97477486	C	T	0.92	1.02	-0.22	0.05	3.19x10 ⁻⁶
		2	rs13394665	* 64244935	A	C	0.73	0.97	0.15	0.03	6.70x10 ⁻⁷
		2	rs1069256	* 173881973	C	T	0.75	1.02	0.17	0.03	6.00x10 ⁻⁷
		3	rs9874245	81211967	A	G	0.85	1.03	0.20	0.04	1.63x10 ⁻⁷
		4	rs1380155	128947268	A	G	0.95	1.91	0.45	0.10	4.22x10 ⁻⁶
		7	rs17168027	* 134099287	A	G	0.89	0.89	0.22	0.05	1.98x10 ⁻⁶
		10	rs11003972	55421987	C	G	0.85	0.98	0.19	0.04	2.09x10 ⁻⁶
		15	rs1026672	64280411	C	T	0.81	0.98	0.16	0.03	4.10x10 ⁻⁶
		15	rs8029458	89405378	G	T	0.80	0.93	-0.16	0.04	3.67x10 ⁻⁶
		15	rs11630508	99138869	A	G	0.15	0.87	0.20	0.04	1.39x10 ⁻⁶
European											
	Baseline	1	rs1543251	105839118	G	T	0.86	1.02	0.20	0.04	1.33x10 ⁻⁷
		2	rs2602569	168998331	C	T	0.08	0.98	0.22	0.05	4.11x10 ⁻⁶
		4	rs4399935	156175906	A	C	0.65	0.97	0.13	0.03	4.71x10 ⁻⁶
		7	rs7457833	* 33521559	A	G	0.90	1.01	-0.20	0.04	3.26x10 ⁻⁶
		8	rs1460396	2913482	A	G	0.19	0.89	0.16	0.03	4.90x10 ⁻⁶
		10	rs6583751	92828756	C	G	0.61	0.96	0.13	0.03	2.48x10 ⁻⁶
		13	rs1838113	18394504	C	T	0.07	0.81	-0.26	0.06	4.11x10 ⁻⁶
		13	rs4884095	77898753	A	T	0.78	0.98	-0.15	0.03	9.09x10 ⁻⁷
	Averaged	1	rs12564617	105832151	A	G	0.14	1.01	-0.15	0.03	1.71x10 ⁻⁶

	3	rs17694541	*	135885739	C	T	0.08	1.06	-0.18	0.04	4.91x10 ⁻⁶
	7	rs7457833	*	33521559	A	G	0.90	1.01	-0.18	0.04	2.12x10 ⁻⁶
	9	rs10739282	*	111488202	A	C	0.41	0.99	0.11	0.02	8.26x10 ⁻⁷
	10	rs12146350		22883117	C	G	0.40	0.97	-0.11	0.02	3.48x10 ⁻⁶
Repeated Measures	1	rs897632		2816023	A	G	0.23	1.00	-0.13	0.03	2.51x10 ⁻⁶
	1	rs12563710		105831868	A	C	0.86	1.01	0.16	0.03	1.30x10 ⁻⁶
	3	rs17694541	*	135885739	C	T	0.08	1.06	-0.19	0.04	3.51x10 ⁻⁶
	7	rs4720629		5441935	C	T	0.91	1.01	0.19	0.04	2.12x10 ⁻⁶
	7	rs7457833	*	33521559	A	G	0.90	1.01	-0.19	0.04	7.93x10 ⁻⁷
	9	rs10739282	*	111488202	A	C	0.41	0.99	0.11	0.02	8.40x10 ⁻⁷
	10	rs7916922		22859264	C	T	0.37	0.92	-0.11	0.02	3.93x10 ⁻⁶
	10	rs11013046		22880589	C	T	0.41	1.00	-0.11	0.02	6.04x10 ⁻⁷
	10	rs12267148		92850261	A	G	0.67	0.93	0.12	0.03	1.52x10 ⁻⁶
Chinese											
Baseline	10	rs12769343		29247812	A	G	0.33	0.92	-0.24	0.05	2.48x10 ⁻⁶
Averaged	11	rs2461672	*	80525681	C	T	0.79	0.90	0.25	0.05	1.42x10 ⁻⁶
	10	rs4529793	*	65960699	A	G	0.09	1.02	0.33	0.07	4.28x10 ⁻⁶
Repeated Measures	2	rs12990176		150385322	G	T	0.06	0.96	-0.32	0.07	3.53x10 ⁻⁶
	10	rs4529793	*	65960699	A	G	0.09	1.02	0.34	0.07	2.71x10 ⁻⁷
	11	rs2461672	*	80525681	C	T	0.79	0.90	0.24	0.05	3.89x10 ⁻⁶
	12	rs17115484		74370892	C	T	0.10	0.97	-0.29	0.06	7.05x10 ⁻⁷
Hispanic											
Baseline	2	rs17023193		83192390	A	T	0.12	1.03	-0.28	0.06	3.94x10 ⁻⁷
	3	rs1127233	*	126109714	G	T	0.23	1.00	0.24	0.04	3.85x10 ⁻⁸
	4	rs2903246		11750932	A	T	0.38	1.03	-0.18	0.04	3.21x10 ⁻⁶
	4	rs4695747	*	175015856	A	G	0.93	0.99	0.39	0.07	1.25x10 ⁻⁷
	5	rs806600		172847545	A	G	0.49	0.99	-0.17	0.04	2.93x10 ⁻⁶
	7	rs909092		141759938	A	G	0.90	0.98	0.31	0.06	7.03x10 ⁻⁷
Averaged	4	rs4695747	*	175015856	A	G	0.93	0.99	0.28	0.06	4.78x10 ⁻⁶
	5	rs7728557	*	74329715	A	G	0.32	1.03	0.15	0.03	1.61x10 ⁻⁶
	14	rs8016794		52070396	A	G	0.83	1.06	0.18	0.04	4.49x10 ⁻⁶
Repeated Measures	2	SNP_A-1966287		191577187	C	T	0.11	1.03	0.24	0.05	1.71x10 ⁻⁶
	3	rs1127233	*	126109714	G	T	0.23	1.00	0.18	0.04	2.06x10 ⁻⁶
	4	rs4695747	*	175015856	A	G	0.93	0.99	0.31	0.06	9.25x10 ⁻⁷
	5	rs7728557	*	74329715	A	G	0.32	1.03	0.16	0.03	1.96x10 ⁻⁶

5	rs244897	167851370	C	T	0.11	0.92	0.24	0.05	4.23x10 ⁻⁶
6	rs3805944	43153440	G	T	0.91	1.03	-0.24	0.05	3.11x10 ⁻⁶
7	rs123778	157062905	C	G	0.85	1.31	-0.24	0.05	2.43x10 ⁻⁶
14	rs7148635	66979360	A	G	0.59	1.02	-0.15	0.03	4.25x10 ⁻⁶
14	rs17106996	77702071	A	G	0.92	0.95	-0.27	0.06	1.03x10 ⁻⁶
16	rs9652625	51663724	C	G	0.68	0.89	-0.17	0.04	2.73x10 ⁻⁶

*indicates SNP that reached genome-wide suggestive p-value in multiple approaches

MAF>5%, INFO > 0.80, LD R² > 0.80

CHR: chromosome, SNP: single nucleotide polymorphism, BP: base pair, A1/A2: coded and non-coded allele, FRQ: coded allele frequency, INFO: imputation information score, SE: standard error, MAF: minor allele frequency, LD: linkage disequilibrium