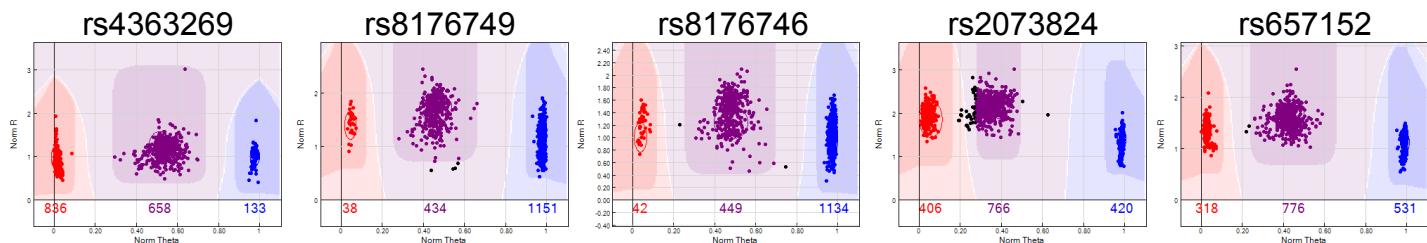


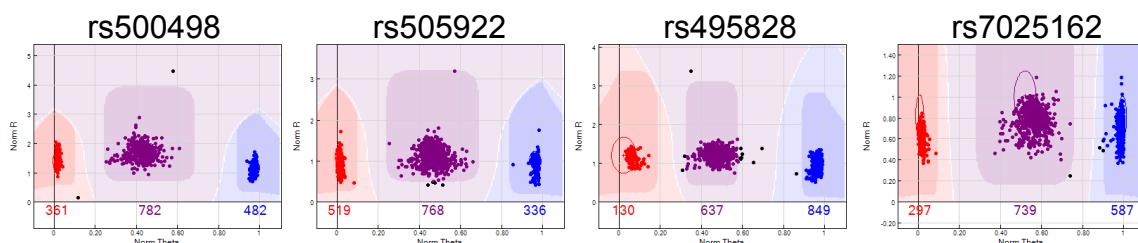
Figure S1

ABO in 9q32

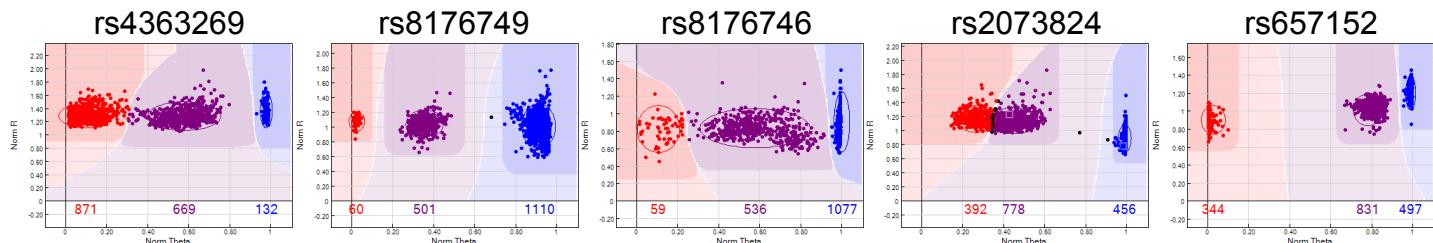
Infinium assay



Infinium assay



Golden Gate assay



Golden Gate assay

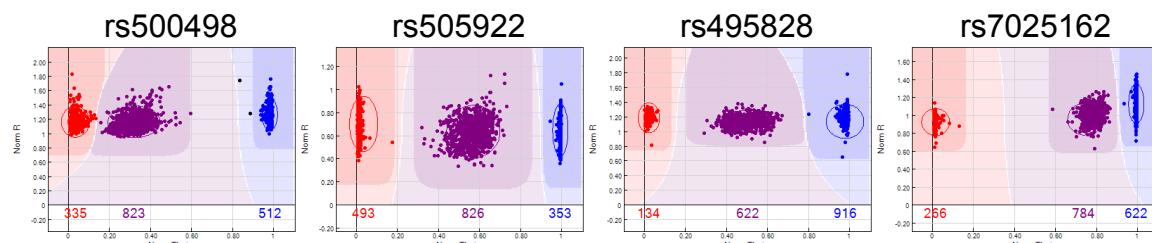
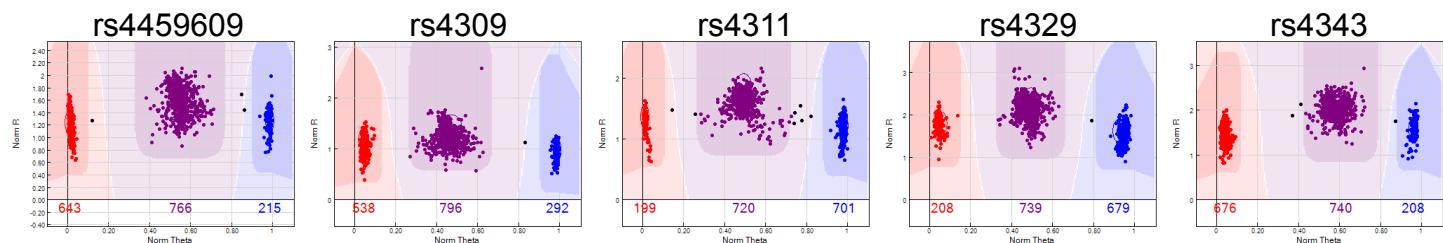


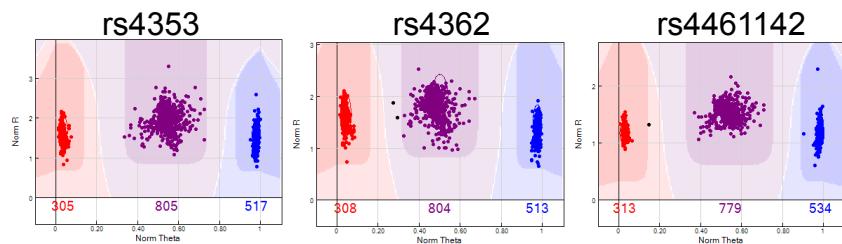
Figure S2

ACE1 in 17q23.2

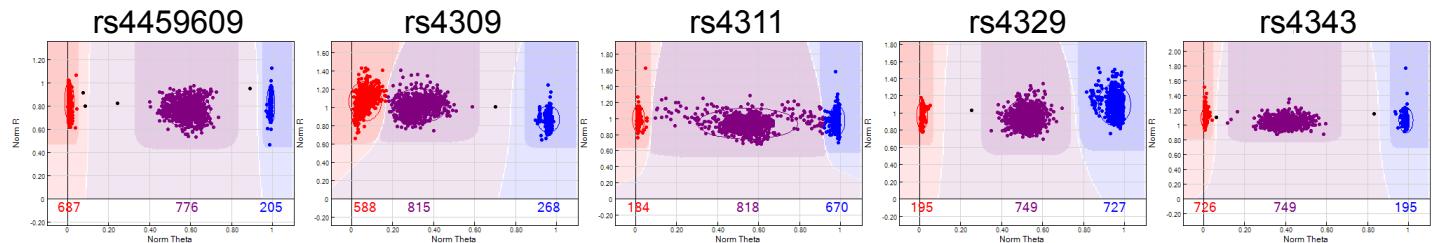
Infinium assay



Infinium assay



Golden Gate assay



Golden Gate assay

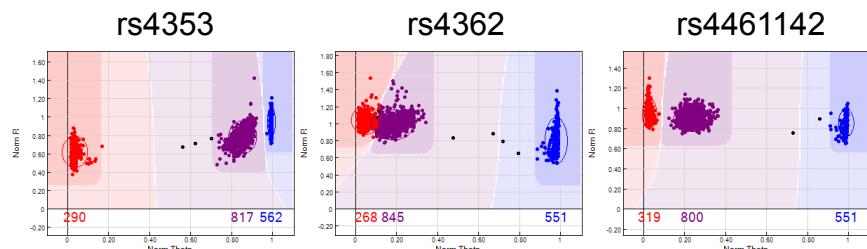


Figure S3

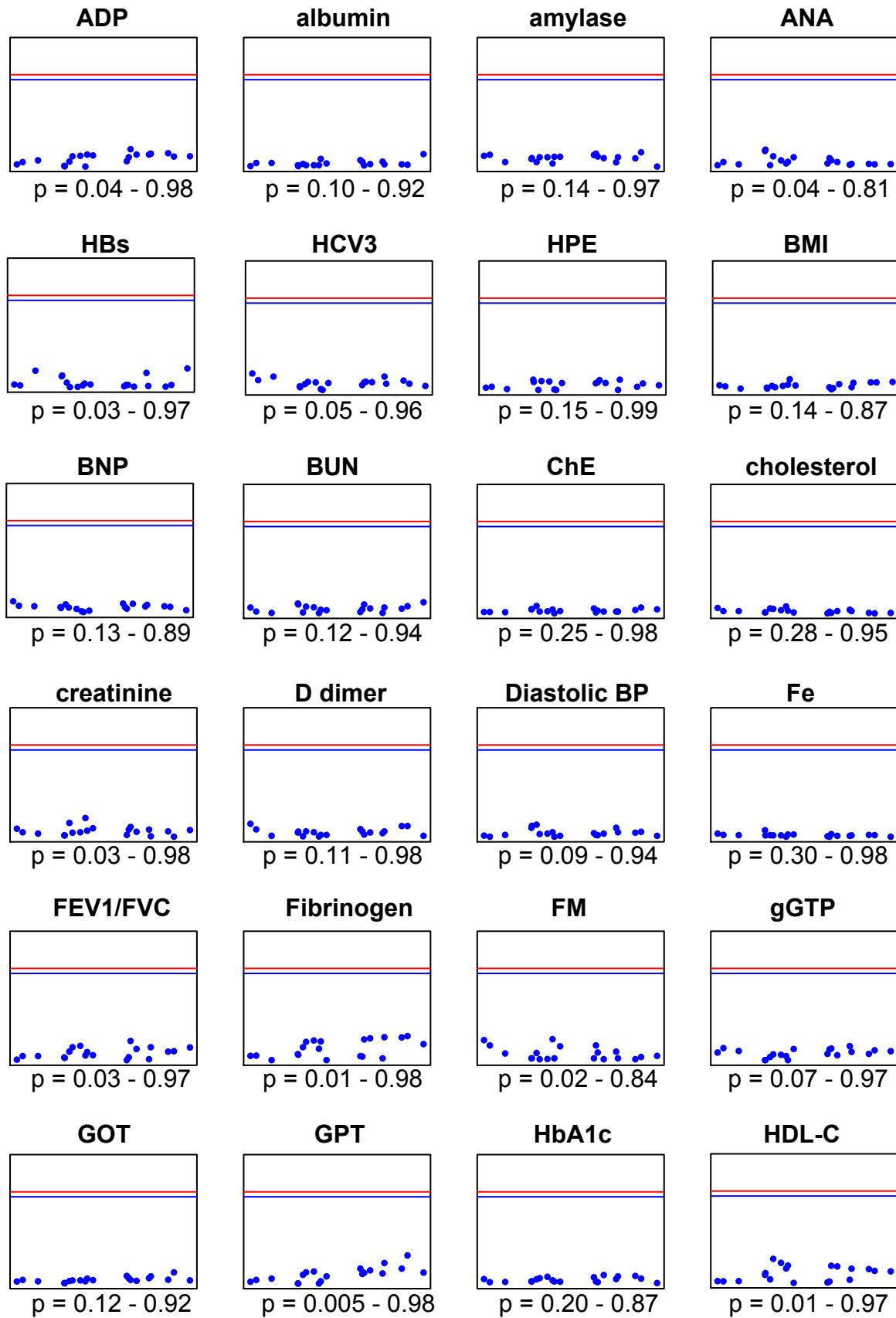


Figure S3 (continued)

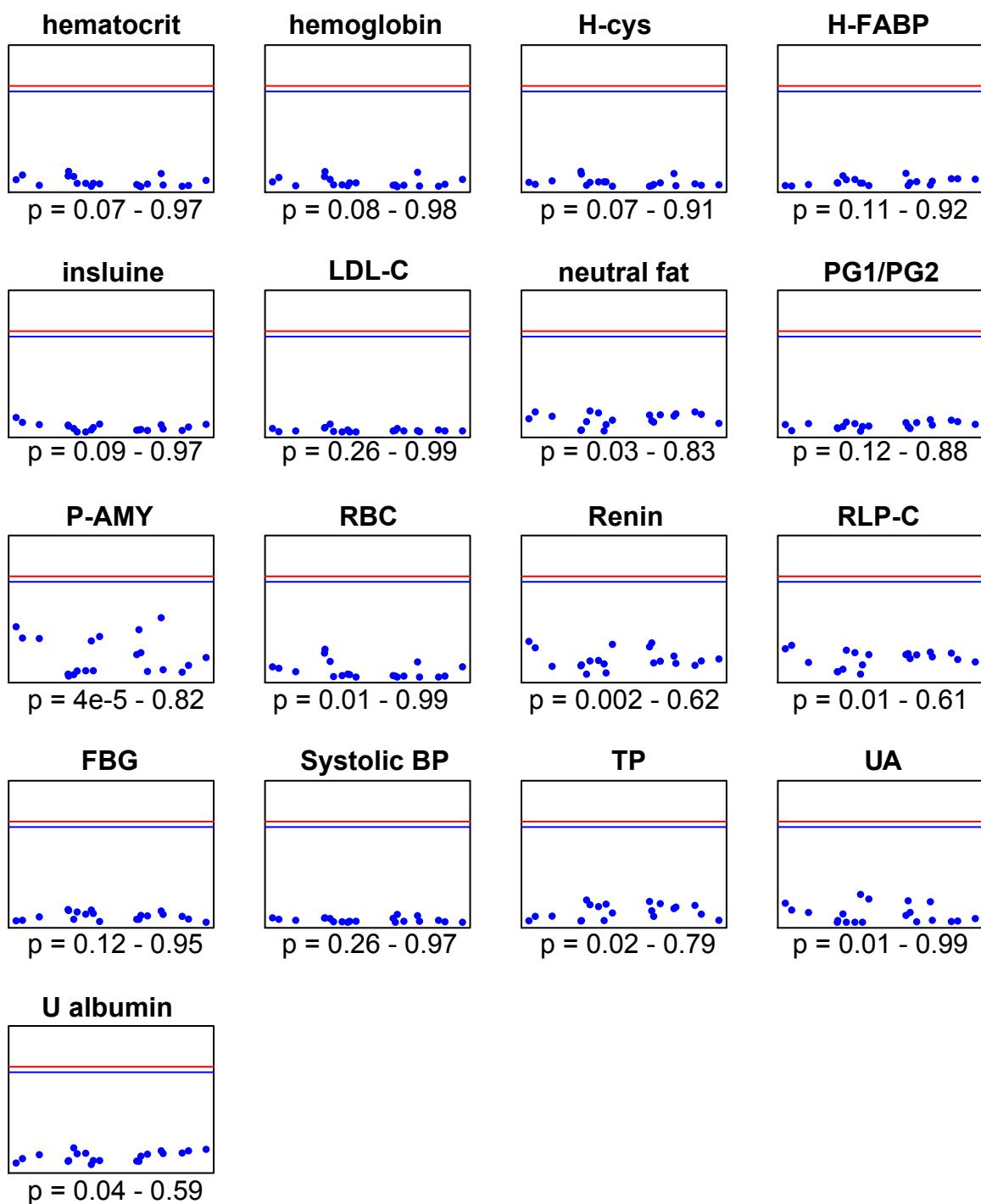
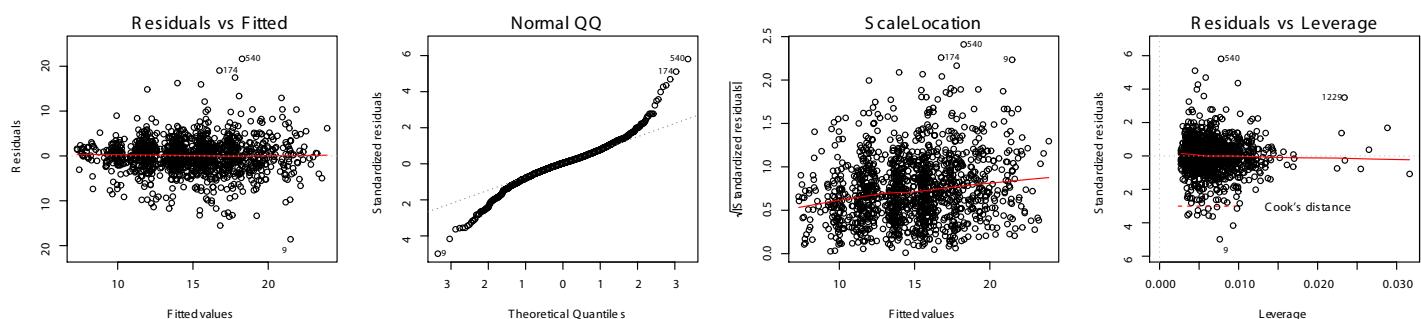
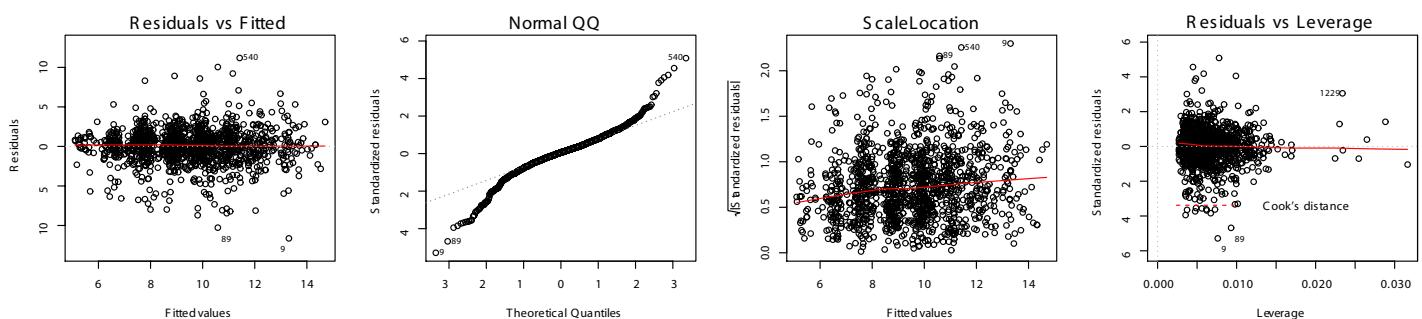


Figure S4

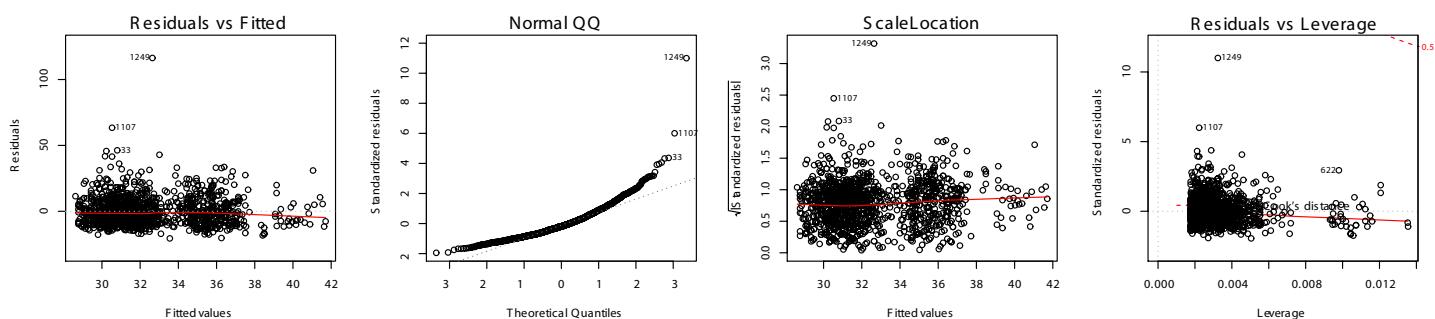
Untransformed ACE



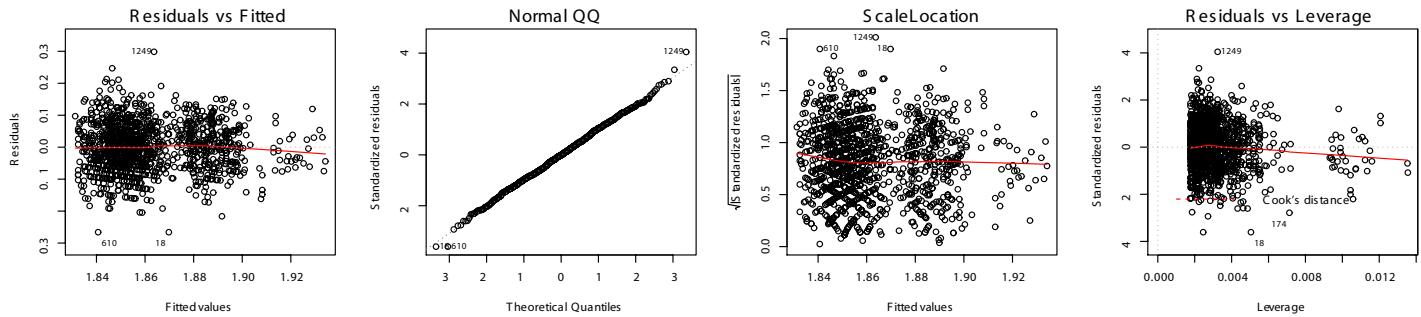
Box-Cox transformed ACE



Untransformed P-LIP



Box-Cox transformed P-LIP



Methods S1

Imputation and test: We first carried out pre-phasing [1] using SHAPEIT [2] based on 1000 Genome Project for chromosome 9, then applied IMPUTE2 [3] to ABO locus to complete imputation for ABO locus. Subsequently we used SNPTEST [4] to examine quantitative traits association with loci after imputation, in which adjustment by age and sex was incorporated. The above procedures were conducted separately for the first and second stage datasets.

[1] Howie B, Fuchsberger C, Stephens M, Marchini J & Abecasis GR. Fast and accurate genotype imputation in genome-wide association studies through pre-phasing. *Nat Genet* 44(8): 955-959 (2012).

[2] Delaneau O, Marchini J & Zagury JF A linear complexity phasing method for thousands of genomes. *Nat. Methods* 9, 179-181 (2011).

[3] Howie BN, Donnelly P & Marchini J. A flexible and accurate genotype imputation method for the next generation of genome-wide association studies. *PLoS Genet* 5(6): e1000529 (2009).

[4] Marchini J, Howie B, Myers S, McVean G & Donnelly P. A new multipoint method for genome-wide association studies via imputation of genotypes. *Nat*

Genet 39: 906-913 (2007).

Regression diagnosis: We carried out regression diagnosis for each marker being significant in GWAS on both ACE and P-LIP for 1st and 2nd stage samples, utilizing four diagnosis plots implemented in R (fitted values vs residuals, normal QQ plot for standardized residuals, fitted values vs $\sqrt{\text{standardized residuals}}$, and leverage vs standardized residuals). We performed the diagnosis plots both for untransformed and for optimally Box-Cox transformed traits ($\lambda = 0.8$ for ACE and $\lambda = -0.4$ for P-LIP).

Table S1. Associations of imputed genotypes in the ABOlocus with P-LIP in Takahata population.

index	id*	rsid	chr	pos	Allele	average maximum posterior call	info	Genotypes				MAF	beta	SE	P-value	
								AA	BB	Null						
1	---	rs7562368	NA	136123019	C	T	0.99972	0.058588	1251.6	0.351	0	0.00014018	ND	ND	ND	
2	---	rs11240409	NA	136123092	T	C	0.99546	0.98526	907.73	314.42	29.845	0.008	0.14941	0.34474	0.055764	6.3238e-10
3	---	rs148031614	NA	136123172	T	C	0.99928	0.25135	1251.1	0.9	0	0	0.00035942	ND	ND	ND
4	---	rs9328546	NA	136123267	C	G	0.93562	0.84338	838.89	376.56	36.519	0.034	0.17956	0.06006	0.058599	0.30539
5	---	rs114204311	NA	136123277	G	A	0.99987	0.083469	1251.8	0.166	0	0	0	ND	ND	ND
6	---	rs192446092	NA	136123600	A	C	0.9999	0.01904	1251.9	0.131	0	0	0	ND	ND	ND
7	---	rs18380736	NA	136123679	C	T	0.99982	0.10392	1251.8	0.221	0	0	0	ND	ND	ND
8	---	rs6597618	NA	136123740	T	C	0.99414	0.98988	459.62	583.73	208.63	0.01	0.39976	0.20029	0.040161	6.1246e-07
9	rs4363269	rs4363269	NA	136123840	A	G	1	1	634	517	101	0	0.28714	-0.0030706	0.044089	0.94448
10	---	rs186098467	NA	136123994	T	C	0.9999	0.0029988	1252	0.003	0	0	0	ND	ND	ND
11	---	rs7025839	NA	136124190	G	A	0.99694	0.99335	703.13	474.76	74.108	0.005	0.24879	0.033506	0.046373	0.46998
12	---	rs72775410	NA	136124314	G	A	0.96768	0.3896	1204.1	47.501	0.434	0.002	0.019317	-0.02986	0.20035	0.88152
13	---	rs73562371	NA	136124326	T	C	0.99955	0.72665	1250.2	1.76	0	0	0.00070288	-0.96353	0.85558	0.26009
14	---	rs9411473	NA	136124399	G	A	0.99562	0.98996	702.96	47.493	74.095	0.021	0.24885	0.03319	0.046467	0.47506
15	---	rs9411370	NA	136124432	A	G	0.99538	0.98938	702.97	47.494	74.071	0.015	0.24884	0.033183	0.046484	0.47531
16	---	rs78755596	NA	136124590	T	A	0.99747	0.075064	1248.8	3.171	0	0	0.0012664	ND	ND	ND
17	---	rs11537044	NA	136124602	T	G	0.99982	0.017178	1251.8	0.225	0	0	0	ND	ND	ND
18	---	rs12338026	NA	136124857	C	G	0.9993	0.58542	1250.4	1.624	0	0	0.00064856	ND	ND	ND
19	---	rs7047076	NA	136125329	A	T	0.989	0.97695	702.56	47.581	73.599	0.031	0.24881	0.032534	0.046852	0.48743
20	---	rs193302481	NA	136125592	C	T	1	0.0009992	1252	0.002	0	0	0	ND	ND	ND
21	---	rs9411474	NA	136125716	C	G	0.98636	0.97321	702.47	47.616	73.432	0.031	0.24882	0.032289	0.046956	0.49168
22	---	rs61527302	NA	136125994	G	A	0.98778	0.97299	813.68	388.96	49.337	0.007	0.19474	0.077463	0.050608	0.12585
23	---	rs191151099	NA	136126026	T	A	1	1	125	0	0	0	0	ND	ND	ND
24	---	rs10901251	NA	136126129	A	C	0.97626	0.96443	456.28	577.46	218.27	0.044	0.40497	0.20263	0.040336	5.0714e-07
25	---	rs148763789	NA	136126132	C	T	0.9999	0.0058266	1252	0.012	0	0	0	ND	ND	ND
26	---	rs186858949	NA	136126153	G	T	1	0.0019992	1252	0.002	0	0	0	ND	ND	ND
27	---	rs76320709	NA	136126394	C	T	0.989	0.49772	1231.6	20.321	0.068	0.002	0.0081697	-0.39605	0.26639	0.13709
28	---	rs34266699	NA	136126495	C	T	0.98406	0.96903	702.47	47.64	73.112	0.022	0.24866	0.032268	0.047075	0.49305
29	---	rs151316644	NA	136126625	G	A	0.99948	0.039007	1251.3	0.653	0	0	0.0026078	ND	ND	ND
30	---	rs3205136	NA	136126631	C	A	0.98673	0.96021	895.8	322.73	33.451	0.018	0.15561	0.33809	0.055421	1.0577e-09
31	---	rs7849280	NA	136126636	A	G	0.94202	0.87242	783.14	416.66	52.162	0.033	0.20807	0.028529	0.053719	0.59537
32	---	rs116958345	NA	136127122	A	C	0.99974	0.0183185	1251.7	0.3	0	0	0.0013179	ND	ND	ND
33	---	rs9411475	NA	136127268	T	C	0.95823	0.92124	649.8	507.3	94.676	0.044	0.27823	0.018941	0.046111	0.68124
34	---	rs110805674	NA	136127346	C	T	0.9999	0.0019952	1252	0.012	0	0	0	ND	ND	ND
35	---	rs12216891	NA	136127366	C	T	0.97965	0.94627	834.13	376.17	41.661	0.04	0.18351	0.3858	0.0503041	3.4977e-13
36	chr9:136127382:I	NA	136127382	A	AGCCCTC	0.99991	0.015851	1251.9	0.115	0	0	0	ND	ND	ND	
37	---	rs13291798	NA	136127481	A	G	0.99927	0.25957	1242.2	9.51	0	0.002	0.0038982	-0.49236	0.24544	0.27327
38	---	rs18619566	NA	136127634	C	A	1	1	1252	0	0	0	0	ND	ND	ND
39	---	rs9411476	NA	136127801	G	A	0.98081	0.95873	741.55	443.51	66.925	0.017	0.23058	0.0303088	0.047903	0.49055
40	---	rs34085694	NA	136127805	C	G	0.91957	0.24806	1241.4	10.634	0.003	0.004	0.0042492	-0.49702	0.04063	0.21933
41	---	rs1692653	NA	136127841	T	C	0.99988	0.00838	895.06	57.97	54.92	0.009	0.02355	-0.17926	0.044681	8.305e-05
42	---	rs115634200	NA	136127859	A	G	0.98949	0.18283	1250.1	1.89	0	0	0.00375479	ND	ND	ND
43	---	rs14657022	NA	136127736	C	T	0.99748	0.11506	248.8	314.5	0.002	0	0.0012598	ND	ND	ND
44	---	rs115413527	NA	136127737	G	A	0.98845	0.26106	1236.9	15.001	0	0	0.0062688	-0.14746	0.38746	0.70352
45	---	rs62574563	NA	136127764	C	T	0.96103	0.91922	675.26	48.643	92.006	0.071	0.26704	-0.12629	0.046073	0.006123
46	---	rs62574564	NA	136127811	G	A	0.96476	0.92562	683.52	482.47	85.92	0.057	0.26132	-0.14733	0.046479	0.001525
47	---	rs73660468	NA	136127907	G	A	0.99996	0.012682	1251.9	0.064	0	0	0	ND	ND	ND
48	---	rs189931088	NA	136127934	A	C	0.99995	0.033943	1251.5	0.058	0	0	0	ND	ND	ND
49	---	rs43039247	NA	136127956	A	C	0.99135	0.23454	1241.2	10.838	0.001	0.001	0.0043251	-0.63652	0.39745	0.24362
50	---	rs10901251	NA	136128000	G	C	0.99201	0.97662	838.99	369.68	43.306	0.017	0.18223	0.37794	0.05184	2.3166e-13
51	---	rs141318395	NA	136128175	G	A	0.99976	0.008487	1251.7	0.299	0	0	0.0011941	ND	ND	ND
52	---	rs62574565	NA	136128259	G	A	0.98307	0.96009	706.68	47.13	78.136	0.054	0.24897	-0.13633	0.046705	0.003511
53	---	rs58081338	NA	136128329	T	C	0.98293	0.96091	705.84	46.79	78.203	0.057	0.24933	-0.1371	0.046684	0.0033167
54	---	rs62574567	NA	136128421	C	G	0.98235	0.95978	706.69	46.716	78.094	0.05	0.24895	-0.13692	0.046709	0.0033756
55	---	rs60484807	NA	136128467	A	G	0.98558	0.97444	425.78	60.68	216.5	0.04	0.16462	0.12501	0.004749	0.0021568
56	---	rs78575390	NA	136128546	A	G	0.97543	0.94996	706.49	51.04	630.16	0.056	0.29281	0.176	0.044343	7.2195e-05
57	---	rs77693339	NA	136128558	T	C	0.99453	0.98877	868.78	348.12	35.085	0.016	0.16705	0.339978	0.053789	1.0667e-13
58	---	rs12554580	NA	136128603	T	C	0.9858	0.97488	425.82	60.68	215.04	0.043	0.16369	-0.12047	0.20744	0.0021543
59	---	rs125543436	NA	136128663	A	G	0.96931	0.93122	719.42	46.02	72.445	0.053	0.24159	-0.13583	0.047909	0.0048513
60	---	rs12554591	NA	136128664	T	C	0.96954	0.93141	705.68	45.28	73.77	0.049	0.24122	-0.13513	0.047933	0.0048133
61	---	rs112404501	NA	136128731	G	A	0.99583	0.95956	1037.23	26.21	0.003	0	0.0020839	ND	ND	ND
62	---	rs12554339	NA	136128737	A	C	0.98236	0.95916	705.65	46.74	78.678	0.034	0.2496	-0.13505	0.046741	0.003861
63	---	rs10901251	NA	136128772	T	C	0.98353	0.96237	706.03	46.75	78.166	0.047	0.24924	-0.13677	0.046658	0.0033761
64	---	rs10751502	NA	136129079	A	G	0.98545	0.97436	425.26	60.23	217.47	0.042				

continued 1

120	---	rs8176731	NA	136132350	T	C	0.99741	0.99567	428.81	606.82	216.35	0.015	0.41515	0.11926	0.040332	0.0031066
121	---	rs2073823	NA	136132516	G	A	0.99758	0.99453	860.98	352.1	38.922	0	0.1717	0.3813	0.052409	3.4373e-13
122	---	rs8176730	NA	136132525	T	C	0.99761	0.99463	860.99	352.07	38.936	0	0.1717	0.3813	0.052406	3.44e-13
123	---	rs8176728	NA	136132561	G	C	0.99785	0.99567	718.21	456.02	77.765	0.005	0.24423	-0.13836	0.045954	0.026059
124	---	rs8176727	NA	136132570	G	A	0.99823	0.99639	718.31	456.29	77.395	0.008	0.24405	-0.13809	0.045985	0.026742
125	---	rs112655244	NA	136132608	C	T	0.99795	0.9959	718.35	455.96	77.685	0.007	0.24414	-0.13841	0.045967	0.026033
126	---	rs8176725	NA	136132617	G	A	0.99526	0.99115	721.82	446.91	83.272	0	0.24499	0.3069	0.045456	1.4632e-11
127	rs2073824	rs2073824	NA	136132633	A	G	1	1	327	609	316	0	0.49561	0.12405	0.039195	0.0015511
128	---	rs2073825	NA	136132707	A	T	0.99883	0.99749	718.55	456.18	77.27	0.001	0.2439	-0.13807	0.045986	0.0026785
129	---	rs8176722	NA	136132754	C	A	0.99867	0.99626	860.62	351.83	39.546	0.001	0.1721	0.37927	0.025228	4.0392e-13
130	---	rs8176721	NA	136132852	G	A	0.99934	0.062922	1251.2	0.82	0	0.001	0.00032748	ND	ND	ND
131	---	rs8176720	NA	136132873	T	C	0.99964	0.9993	428.06	606.21	217.73	0	0.416	0.11869	0.04023	0.0031746
132	---	chr9:136132908:1	NA	136132908	T	TC	0.99875	0.99767	387.31	595.14	269.04	0.002	0.45297	0.23404	0.039133	2.221e-09
133	---	rs75179845	NA	136132954	T	C	0.99743	0.99301	862.38	350.51	39.138	0.003	0.17124	0.38078	0.05249	4.0388e-13
134	---	rs8176718	NA	136132957	C	T	0.99986	0.99962	718.94	456.01	77.051	0	0.24365	-0.13811	0.045994	0.0026754
135	---	rs8176717	NA	136133034	G	T	0.99933	0.99837	719.63	455.38	76.989	0	0.24335	-0.13799	0.046009	0.0027072
136	---	rs8176716	NA	136133065	G	T	0.99863	0.15653	1250.3	0	0	0	0.00068371	ND	ND	ND
137	---	rs8176715	NA	136133148	T	C	0.99802	0.97381	83.13	476.46	926.39	0.003	0.25661	0.15219	0.045798	0.00089066
138	---	rs191976170	NA	136133149	G	A	0.99921	0.24201	1251.1	0.82	0	0.001	0.00032748	ND	ND	ND
139	---	rs8176714	NA	136133178	G	A	0.99356	0.98601	648.13	505.84	98.016	0.001	0.2803	-0.16059	0.04443	0.00030096
140	---	rs115166192	NA	136133218	C	T	0.99965	0.0086304	1251.6	0.441	0	0	0.00017612	ND	ND	ND
141	---	rs182985414	NA	136133346	A	C	1	1	1252	0	0	0	0	ND	ND	ND
142	---	chr9:136133380:D	NA	136133380	CGGG	C	0.99867	0.99649	76.608	455.3	720.09	0.004	0.24302	0.13713	0.046111	0.0029396
143	---	rs45610939	NA	136133396	A	G	0.9792	0.8157	1146.1	105.73	204	0.003	0.042387	-0.23914	0.10983	0.0076077
144	rs512770	rs512770	NA	136133506	A	G	1	77	456	719	0	0	0.24361	0.1381	0.045998	0.0026784
145	---	rs4962040	NA	136133531	A	G	0.99764	0.99606	629.73	513.26	109.01	0.008	0.29204	-0.16	0.043586	0.000116
146	---	chr9:136133582:D	NA	136133582	AG	A	0.99258	0.98658	91.997	483.41	676.59	0.002	0.26564	0.1278	0.044937	0.0044552
147	---	rs641959	NA	136133699	C	A	0.99758	0.99575	89.25	468.68	694.07	0.001	0.25846	0.13101	0.044934	0.0035502
148	---	rs641943	NA	136133714	G	A	0.99778	0.99624	89.165	468.59	694.24	0.001	0.25836	0.13091	0.044938	0.0035781
149	---	rs514708	NA	136133743	T	C	0.99778	0.99625	89.165	468.59	694.24	0	0.25836	0.13091	0.044938	0.0035785
150	---	rs517414	NA	136134034	A	G	0.9996	0.99919	80.042	459.41	712.54	0	0.2474	0.13739	0.045696	0.0026413
151	---	rs9411372	NA	136134068	G	A	0.97122	0.96464	730.07	444.16	77.763	0.012	0.23949	0.036719	0.047153	0.43615
152	---	rs138164693	NA	136134118	G	A	1	1	1252	0	0	0	0	ND	ND	ND
153	---	rs638756	NA	136134172	C	A	0.99787	0.99641	89.116	468.64	694.24	0	0.25834	0.13085	0.04494	0.0035961
154	---	rs191050088	NA	136134650	C	A	0.99671	0.14604	1247.1	412.2	0	0	0.0016462	ND	ND	ND
155	---	rs183232893	NA	136134835	G	A	1	0.0033309	1252	0.006	0	0	0	ND	ND	ND
156	---	rs626792	NA	136134864	C	A	0.93522	0.87801	103.9	510.59	637.49	0.002	0.2869	0.13935	0.047145	0.0234334
157	---	rs626035	NA	136134874	G	T	0.97273	0.94743	99.363	483.5	669.14	0.006	0.27245	0.13245	0.044486	0.0015325
158	---	rs547495	NA	136134895	C	T	0.97273	0.94743	99.363	484.49	668.14	0.006	0.27245	0.13245	0.044486	0.0015329
159	---	rs547643	NA	136135047	T	C	1	0.001	459	713	0	0	0.2472	0.13752	0.045697	0.0026164
160	---	rs62663	NA	136135096	A	G	0.99098	0.00995	80.002	468.02	710.99	0	0.24721	0.13752	0.045697	0.0026665
161	---	rs8176707	NA	136135108	G	A	0.9361	0.75539	1021.6	241.61	15.814	0.015	0.0033338	0.26212	0.073878	0.00038065
162	---	chr9:136135137:D	NA	136135137	GCTTCCTCTCCACAC	G	0.99819	0.97363	85.311	468.49	700.19	0.007	0.25514	0.14555	0.04022	0.0034426
163	---	rs13674534	NA	136135194	G	A	0.99916	0.84615	1246.3	5.64	0	0	0.002262	-0.13106	0.045575	0.60410
164	---	rs49331	NA	136135195	G	A	0.99799	0.99664	89.053	468.71	694.24	0	0.25831	0.13077	0.044942	0.0036164
165	---	rs549443	NA	136135237	A	G	0.89151	0.7555	120.08	528.79	603.05	0.005	0.30711	0.12136	0.047813	0.01114
166	---	rs549446	NA	136135238	T	C	0.86399	0.79851	130.45	531.66	589.83	0.005	0.31653	0.1413	0.046863	0.0025677
167	---	rs14927903	NA	136135321	G	A	1	1	1252	0	0	0	0	ND	ND	ND
168	---	rs624601	NA	136135365	A	G	0.99799	0.99664	89.053	468.71	694.24	0	0.25831	0.13077	0.044942	0.0036164
169	---	rs551322	NA	136135444	G	A	1	80	459	713	0	0	0.2472	0.13752	0.045697	0.0026186
170	---	rs192192638	NA	136135455	C	T	0.96586	0.37061	1202.7	49.07	2.68	0.008	0.019782	-0.21201	0.19199	0.26947
171	---	rs180880130	NA	136135472	C	T	0.99846	0.024821	1250.1	1.97	0	0.002	0.00076957	ND	ND	ND
172	---	rs613423	NA	136135478	A	G	1	0.0033309	1250.4	1.574	0	0	0.00062859	ND	ND	ND
173	---	rs8176704	NA	136135552	G	A	0.99874	0.99787	1250.4	1.574	0	0	0.2472	0.13752	0.045697	0.0026186
174	rs574347	rs574347	NA	136135659	CTG	C	1	1	1252	0	0	0	0	ND	ND	ND
175	rs575725	rs575725	NA	136135752	G	A	0.99798	0.99664	89.052	468.71	694.24	0.001	0.25831	0.13077	0.044942	0.0026186
176	---	rs8176703	NA	136135863	G	T	0.99894	0.21781	1250.7	15.3	0	0	0.00053914	ND	ND	ND
177	---	rs8176702	NA	136136146	G	A	0.99893	0.99712	630.2	512.79	109.01	0.001	0.29186	-0.16821	0.04356	0.00011268
178	---	rs759483	NA	136136196	A	T	0.99999	0.99998	80	459.01	712.99	0	0.24721	0.13751	0.045697	0.0026198
179	---	rs140166545	NA	136136367	C	A	0.99989	0.058102	1251.9	0.137	0	0	0.00090216	ND	ND	ND
180	---	rs687261	NA	136137065	A	G	0.99796	0.99735	378.17	594.14	279.69	0.002	0.46067	0.23505	0.039029	1.7184e-09
181	---	rs184666147	NA	136137088	G	A	0.99996	0.020119	1252	0.44	0	0	0	ND	ND	ND
182	---	rs687289	NA	136137106	G	A	0.99985	0.99973	393.1	591.97	266.92	0	0.44961	0.23365	0.039062	2.2119e-09
191	---	rs2073827	NA	136137133	G	C	1	1	609	527	116	0	0.30312	-0.16228	0.043132	0.00016833
192	rs2073828	rs2073828	NA	136137140	G	A	1	1	609	527	11					

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241	---	rs2519093	NA	136141870	C	T	0.99806	0.99625	647.5	505.95	98.545	0.004	0.28077	0.027212	0.04425	0.53857
242	---	rs144140881	NA	136141877	G	T	1	0.0059976	1252	0.006	0	0	0	ND	ND	ND
243	---	rs113423093	NA	136142000	G	C	0.96439	0.93239	103.39	487.25	661.33	0.03	0.27718	0.12425	0.045443	0.0062532
244	---	rs8176672	NA	136142185	C	T	0.99786	0.99415	874.05	341.66	36.29	0.001	0.16543	0.33692	0.05322	8.7796e-14
245	---	rs514659	NA	136142203	A	C	0.99836	0.99762	397.73	589.42	264.85	0	0.44693	0.23373	0.039049	2.1551e-09
246	---	rs644234	NA	136142217	T	G	0.99837	0.99763	397.73	589.42	264.85	0	0.44693	0.23373	0.039049	2.1556e-09
247	---	rs182705402	NA	136142293	C	T	1	1	1252	0	0	0	0	ND	ND	ND
248	---	rs140796254	NA	136142304	A	T	1	1	1252	0	0	0	0	ND	ND	ND
249	---	rs8176671	NA	136142313	A	C	0.99688	0.99114	874.8	341.48	35.716	0.002	0.1649	0.39753	0.05343	1.0046e-13
250	---	rs143309595	NA	136142339	C	T	1	1	1252	0	0	0	0	ND	ND	ND
251	---	rs643434	NA	136142355	G	A	0.99816	0.99722	397.68	589.26	265.06	0.001	0.44704	0.23399	0.039054	2.0789e-09
252	---	rs8176669	NA	136142463	A	C	0.99711	0.99461	707.9	460.87	83.226	0.005	0.25053	-0.13538	0.045525	0.0029426
253	---	rs148987354	NA	136142465	G	A	0.9999	0.020262	1251.9	0.128	0	0	0	ND	ND	ND
254	---	chr9:136142521:1	NA	136142521	C	CT	0.99487	0.89609	1204.1	47.878	0	0.004	0.019121	-0.0063473	0.15011	0.96627
255	---	chr9:136142540:1	NA	136142540	T	TTG	1	1	1252	0	0	0	0	ND	ND	ND
256	---	rs143746661	NA	136142711	A	G	0.99742	0.11314	1248.8	3.226	0	0	0.0012883	ND	ND	ND
257	---	rs187821258	NA	136142968	T	C	0.99742	0.9934	396.75	589.54	265.71	0.001	0.44767	0.2362	0.039159	1.622e-09
258	---	rs54304	NA	136143000	A	T	0.99582	0.9934	397.73	589.42	264.85	0	0.44693	0.23372	0.039049	2.1595e-09
259	---	rs146903770	NA	136143055	A	G	0.99111	0.28893	1240.3	11.683	0	0	0.0046657	-0.0046294	0.46627	0.99208
260	---	rs613534	NA	136143120	A	G	0.99583	0.99385	399.27	589.45	263.29	0.001	0.4457	0.23496	0.039102	1.8695e-09
261	---	rs543968	NA	136143121	T	C	0.99437	0.9912	400.49	588.95	262.56	0.002	0.44492	0.23385	0.039169	2.3696e-09
262	---	rs139001859	NA	136143192	G	T	0.99825	0.077245	1249.8	2.196	0	0.001	0.000877	ND	ND	ND
263	---	rs544873	NA	136143212	G	A	0.99502	0.9917	402.45	586.52	263.02	0	0.44432	0.2267	0.03913	6.8917e-09
264	---	rs142956930	NA	136143330	A	G	0.99889	0.098024	1250.6	1.391	0	0	0.0055551	ND	ND	ND
265	---	rs54971	NA	136143372	C	T	0.99838	0.99763	397.73	589.42	264.85	0	0.44693	0.23372	0.039049	2.1595e-09
266	---	rs138194969	NA	136143391	G	A	1	1	1252	0	0	0	0	ND	ND	ND
267	---	rs612169	NA	136143442	A	G	0.99839	0.99764	397.73	589.42	264.85	0	0.44693	0.23373	0.039049	2.158e-09
268	---	rs144566139	NA	136143419	G	A	1	1	1252	0	0	0	0	ND	ND	ND
269	---	chr9:136144000:D	NA	136144000	AT	A	0.98756	0.54188	1223.5	28.432	0.053	0.001	0.011397	0.16805	0.27756	0.54488
270	---	rs8176668	NA	136144059	A	T	0.99815	0.9973	611.89	523.75	116.36	0	0.3021	-0.16333	0.04312	0.00015202
271	---	rs574311	NA	136144110	G	A	0.99799	0.99662	83.583	461.25	707.17	0	0.25096	0.13496	0.045414	0.0029614
272	---	rs187551277	NA	136144172	G	A	0.99825	0.077245	1249.8	2.196	0	0.001	0.000877	ND	ND	ND
273	---	rs597988	NA	136144284	T	A	0.97199	0.95467	459.93	572.81	219.14	0.009	0.40386	0.24214	0.040417	2.0856e-09
274	---	rs597974	NA	136144297	A	G	0.96877	0.94921	462.62	571.57	217.79	0.018	0.40222	0.24298	0.040508	1.9938e-09
275	---	rs140914497	NA	136144308	T	C	0.96718	0.94681	463.77	571.11	217.11	0.017	0.40149	0.24164	0.040583	2.6109e-09
276	---	rs76125	NA	136144309	G	A	0.96718	0.94681	463.77	571.11	217.12	0.017	0.4015	0.24163	0.040582	2.6158e-09
277	---	rs8176663	NA	136144427	T	C	0.99836	0.99751	397.61	589.5	268.49	0	0.447	0.23358	0.039052	2.2144e-09
278	---	rs1876662	NA	136144454	A	G	0.99669	0.9908	874.94	341.45	35.619	0.004	0.16481	0.33785	0.053453	9.8386e-14
279	---	rs149899707	NA	136144521	C	T	1	1	1252	0	0	0	0	ND	ND	ND
280	---	rs484775	NA	136144524	G	A	0.9979	0.99644	83.81	461.14	707.27	0.001	0.25093	0.13473	0.045419	0.0030144
281	---	rs6697526	NA	136144533	G	T	0.99595	0.99095	958.35	272.93	20.717	0.007	0.12557	0.060261	0.67375	0.00015164
282	---	rs7036642	NA	136144626	A	G	0.99811	0.99718	61.96	523.69	116.36	0	0.30208	-0.16337	0.043123	0.00015164
283	---	rs8176664	NA	136144640	A	G	0.99996	0.0086711	1262	0.045	0	0	0	ND	ND	ND
284	---	rs5961441	NA	136144639	T	G	0.99516	0.99701	79.64	461.37	710.04	0	0.24791	0.1373	0.045782	0.002708
285	---	rs146907040	NA	136144771	A	G	1	0.0039984	1252	0.004	0	0	0	ND	ND	ND
286	---	rs491626	NA	136144873	C	T	0.99109	0.98649	411.75	583.83	256.42	0.001	0.43797	0.139894	0.039186	3.8392e-07
287	---	rs629488	NA	136144960	G	A	0.99656	0.99462	391.81	591.2	269	0	0.45095	0.2331	0.039058	2.4003e-09
288	---	rs493246	NA	136144994	G	A	0.98929	0.9882	384.57	593.12	274.3	0.01	0.45596	0.23485	0.032666	2.2184e-09
289	---	rs192389301	NA	136145029	C	T	0.99979	0.10628	1251.7	261.01	0	0	0.0010423	ND	ND	ND
290	---	rs494242	NA	136145118	C	T	0.99587	0.99328	391.13	591.58	269.29	0.001	0.45134	0.23314	0.039088	2.4534e-09
291	---	rs495203	NA	136145240	C	T	0.99587	0.99328	391.13	591.58	269.29	0.001	0.45134	0.23314	0.039088	2.4537e-09
292	---	chr9:136145403:D	NA	136145403	AC	A	0.89421	0.82553	562.28	557.59	132.08	0.005	0.32189	0.20554	0.047103	1.2795e-05
293	---	chr9:136145414:1	NA	136145414	C	CA	0.90476	0.60485	1022.7	217.57	11.653	0.032	0.096197	0.337574	0.085683	1.5876e-07
294	---	chr9:136145418:D	NA	136145418	AC	A	0.88378	0.80486	584.03	533.39	145.34	0.034	0.32048	0.17883	0.047069	0.00014515
295	---	chr9:136145424:D	NA	136145424	AC	A	0.99664	0.92861	684.01	483.64	84.323	0.03	0.2605	0.033456	0.047087	0.47739
296	---	rs4911378	NA	136145425	C	A	0.99783	0.92381	684.98	482.46	84.58	0.038	0.2602	0.035049	0.047122	0.457
297	---	rs582118	NA	136145471	A	G	0.99785	0.99653	392.38	591.52	268.09	0	0.45036	0.23391	0.03908	2.1568e-09
298	---	rs582094	NA	136145484	A	T	0.99785	0.99653	392.38	591.52	268.09	0	0.45036	0.23391	0.03908	2.1568e-09
299	---	rs180922067	NA	136145717	A	G	0.99986	0.035747	1251.8	0.179	0	0	0	ND	ND	ND
300	---	chr9:136145907:1	NA	136145907	G	GA	0.99609	0.99346	349.01	565.09	260.79	0.001	0.44874	0.23502	0.039071	1.7968e-09
301	---	rs2769071	NA	136145974	A	G	0.99601	0.99357	392.38	591.18	268.44	0.004	0.4505	0.23674	0.039157	1.4849e-09
302	---	rs147027134	NA	136145979	A	G	1	1	1252	0	0	0	0	ND	ND	ND
303	---	rs677355	NA	136146046	G	A	0.99691	0.99484	394.01	590.7	267.29	0.003	0.44939	0.23489	0.039079	1.8488e-09
304	---	chr9:136146061:1	NA	136146061	T	TA	0.99429	0.99034	398.01	585.8	268.18	0.003	0.44815	0.23506	0.039103	1.8422e-09
305	---	chr9:136146068:1	NA	136146068	A	AT	0.99726	0.99542	393.77	590.73	267.49	0.005	0.44957	0.		

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362	---	rs566183	NA	136151198	A	G	0.99981	0.9996	80.003	460.79	711.21	0	0.24792	0.13644	0.045699	0.0028311	
363	rs568203	rs568203	NA	136151445	A	C	1	1	80	461	711	0	0.248	0.13668	0.045696	0.0027801	
364	---	rs600038	NA	136151806	T	C	0.99859	0.99702	647.51	505.09	99.397	0.005	0.28111	0.026237	0.044172	0.55254	
365	---	rs8176635	NA	136152009	G	A	0.99781	0.99572	509.3	574.27	168.43	0	0.36387	0.093837	0.041158	0.022611	
366	---	rs8176634	NA	136152070	G	C	0.9989	0.99671	822.29	369.72	59.983	0.002	0.19556	-0.20037	0.048669	3.398e-05	
367	---	rs150185195	NA	136152286	C	T	0.98223	0.82631	1155.9	96.035	0.055	0.005	0.038397	-0.26446	0.11394	0.020284	
368	---	rs146783421	NA	136152336	A	C	1	0.003398	1252	0.005	0	0	0	ND	ND	ND	
369	---	rs182406156	NA	136152347	C	T	0.99998	0.0061435	1252	0.026	0	0	0	ND	ND	ND	
370	---	rs8176633	NA	136152354	C	T	0.9965	0.99222	800.56	385.48	65.975	0	0.20664	-0.19025	0.047796	6.878e-05	
371	---	rs145690885	NA	136152368	G	A	0.97486	0.40551	1210.5	41.39	0.098	0.004	0.016608	-0.090541	0.22581	0.68845	
372	---	rs8176632	NA	136152547	C	T	0.99758	0.99186	867.35	348.72	35.923	0.005	0.16796	0.39646	0.053182	8.998e-14	
373	---	rs8176631	NA	136152594	C	T	0.99974	0.041108	1251.7	0.321	0	0	0.00012819	ND	ND	ND	
374	---	rs8176630	NA	136152722	C	T	0.99683	0.98399	1023.4	219.58	9.028	0.003	0.094904	-0.02347	0.069032	0.73387	
375	---	rs183899911	NA	136153355	G	A	0.99981	0.03951	1251.8	0.242	0	0	0	ND	ND	ND	
376	---	rs663367	NA	136153451	A	G	0.99777	0.99649	81.231	464.11	706.66	0	0.25023	0.13739	0.045667	0.0026254	
377	---	rs525148	NA	136153481	T	C	0.9846	0.97189	85.345	472.52	694.13	0.001	0.25687	0.1407	0.045575	0.0202211	
378	---	rs61696148	NA	136153544	A	T	0.99658	0.98329	1026	216.96	9.047	0.001	0.093873	-0.023778	0.069321	0.73159	
379	---	rs1461538449	NA	136153797	C	T	0.99874	0.036203	1250.4	1.576	0	0	0.00062939	ND	ND	ND	
380	---	rs651007	NA	136153875	C	T	0.99996	0.9999	647.01	504.99	100	0	0.28155	0.025238	0.044093	0.56707	
381	---	rs1124061	NA	136153981	C	T	0.99201	0.57166	1234.7	17.302	0	0	0.00069097	-0.35008	0.27863	0.20896	
382	---	rs187861744	NA	136154097	G	C	0.99827	0.51738	1248.4	3.554	0	0	0.0014193	-0.81551	0.63951	0.89853	
383	---	rs79459	NA	136154168	T	C	0.99997	0.99993	647.01	504.99	100	0	0.28155	0.025236	0.044093	0.56709	
384	---	rs9411488	NA	136154246	G	T	0.99751	0.99417	805.67	381.75	64.577	0	0.20404	-0.198	0.047976	3.6737e-05	
385	---	rs649129	NA	136154304	C	T	0.99998	0.99995	647.01	504.99	100.01	0	0.28155	0.025229	0.044092	0.5672	
386	---	rs10751504	NA	136154433	G	C	0.9682	0.93236	708.84	473.58	69.535	0.044	0.24468	0.3069	0.048743	3.0492e-10	
387	---	chr9:136154263:D	NA	136154623	TC	T	0.99812	0.12397	1249.6	2.357	0	0	0.00094129	ND	ND	ND	
388	---	chr9:136154262:D	NA	136154626	CA	C	0.99812	0.12397	1249.6	2.357	0	0	0.00094129	ND	ND	ND	
389	---	rs147375567	NA	136154866	A	G	0.99888	0.50891	1249.9	2.111	0	0	0.00084305	-0.36094	0.78396	0.64523	
390	rs495828	rs495828	NA	136154867	G	T	1	1	647	505	100	0	0.28155	0.025236	0.044092	0.56709	
391	---	rs635634	NA	136155000	C	T	0.99998	0.99995	647.01	504.98	100.01	0	0.28155	0.025221	0.044092	0.56731	
392	---	rs12683493	NA	136155063	C	T	0.99649	0.98298	1026	216.93	9.046	0.003	0.093858	-0.023794	0.069336	0.73147	
393	---	rs150022816	NA	136155093	G	T	1	1	1252	0	0	0	0	ND	ND	ND	
394	---	rs13298002	NA	136155127	G	A	0.99811	0.12465	1249.6	2.365	0	0	0.00094449	ND	ND	ND	
395	rs500428	rs500428	NA	136155343	A	G	1	1	90	474	688	0	0.26118	0.1299	0.044787	0.0037263	
396	---	rs7030248	NA	136155359	G	A	0.99743	0.99482	692.55	484.83	74.628	0	0.25323	0.29602	0.046352	1.6984e-10	
397	---	rs9411490	NA	136155414	A	G	0.99744	0.99405	805.68	381.77	64.549	0.003	0.20402	-0.19798	0.047982	3.6893e-05	
398	---	rs633862	NA	136155444	C	T	0.99743	0.99608	606.76	366.17	0.002	0.046522	0.23387	0.039386	2.8863e-09		
399	---	rs02361	NA	136155589	C	G	0.99994	0.99895	89.94	47.37	68.04	0	0.26116	0.12953	0.044789	0.0037215	
400	---	rs9411491	NA	136156015	T	C	0.99715	0.99328	88.67	381.78	64.553	0.004	0.20403	-0.19793	0.047995	3.7235e-05	
401	---	rs5589807	NA	136156064	G	A	0.9809	0.47025	1218	33.85	0.148	0.003	0.016361	-0.26662	0.23535	0.25745	
402	---	rs9411381	NA	136156097	T	C	0.99683	0.99246	808.65	381.78	64.556	0.017	0.20403	-0.19784	0.04801	3.7788e-05	
403	---	rs7659074	NA	136156230	C	T	0.98471	0.47114	1226.6	2.241	0.002	0.010257	1.0622	0.2959	0.00032328	0.00032328	
404	---	rs144274742	NA	136156244	G	A	0.9999	0.9999	1250.7	0.001	0	0	0	ND	ND	ND	
405	---	rs40502769	NA	136156498	G	G	0.99999	0.0094373	1252	0.018	0	0	0	ND	ND	ND	
406	---	rs144306020	NA	136156542	A	G	0.999	0.01524	1250.7	1.255	0	0.001	0.0005012	ND	ND	ND	
407	---	rs186119325	NA	136156672	G	G	0.9998	0.016521	1251.8	0.246	0	0	0	ND	ND	ND	
408	---	rs575317	NA	136157037	C	A	0.96391	0.9988	67.661	48.34	665.97	0.026	0.27303	0.12273	0.044781	0.0061314	
409	---	rs558240	NA	136157133	G	A	0.99025	0.97154	89.23	233.18	23.573	0.012	0.15189	-0.076444	0.057078	0.18047	
410	---	rs138026250	NA	136157138	C	A	0.99882	0.13924	1251.8	0.224	0	0	0	ND	ND	ND	
411	---	rs146149084	NA	136157335	A	G	0.9998	0.015153	1252	0.023	0	0	0	ND	ND	ND	
412	---	rs181316261	NA	136157449	C	T	0.99919	0.01179	1251.4	0.64	0	0	0	0.000255593	ND	ND	ND
413	---	rs7867220	NA	136157520	C	T	0.99988	0.032044	1251.8	0.154	0	0	0	ND	ND	ND	
414	---	rs183796796	NA	136157596	C	T	0.99988	0.032044	1251.8	0.154	0	0	0	ND	ND	ND	
415	---	rs142673851	NA	136157644	C	T	0.9994	0.14465	1251.3	0.746	0	0	0	ND	ND	ND	
416	---	rs62574607	NA	136157654	T	C	0.99069	0.87367	1181.9	67.059	3.032	0.002	0.029203	0.040896	0.12094	0.73525	
417	---	rs7867359	NA	136157656	C	G	0.99945	0.12814	1251.3	0.687	0	0	0	0.0027436	ND	ND	ND
418	---	rs7853492	NA	136157785	C	A	0.99945	0.14681	1251.3	0.69	0	0	0	0.0027556	ND	ND	ND
419	---	rs7867739	NA	136157924	C	T	0.99942	0.15659	1251.3	0.726	0	0	0	0.0028994	ND	ND	ND
420	---	rs7867864	NA	136158008	C	G	0.99936	0.008981	1252	0.048	0	0	0	0.000255593	ND	ND	ND
421	---	rs14195220	NA	136158268	A	G	0.99937	0.02787	87.87	612.33	351.77	0.02	0.47448	0.08685	0.039936	0.03086	
422	---	rs570480	NA	136158964	A	G	0.99937	0.038212	0.015	5.06	1246.9	0.003	0.25122	-0.1348	0.045602	0.003117	
423	---	rs570567	NA	136159089	A	G	0.99705	0.96869	724.2	454.3	78.469	0.028	0.24012	-0.15394	0.047072	0.0010746	
424	---	rs2476494	NA	136158288	T	A	0.99862	0.99925	193.0	49.145	5.365	0.013	0.0205652	-0.089473	0.12863	0.48669	
425	---	rs26368280	NA	136158400	C	T	0.99925	0.1935	1251.1	0.942	0	0	0	0.0003762	ND	ND	ND
426	---	rs18485103	NA	136158490	C	T	0.99988	0.0090575	1252.0	0.229	0	0	0	0.00031669	ND	ND	ND
427	---	rs9127311	NA	136158544	T	C	0.98747	0.70227	94	28.74	15.747	0.016	0.12487	-0.055147			

Table S2. Associations of imputed genotypes in the ABOlocus with ACE in Takahata population.

index	id*	rsid	chr	pos	Allele		average posterior	maximum call	info	Genotypes						
					A	B				AA	AB	BB	Null	MAF	beta	SE
---	rs73562368	17	136123019	C	T	0.99972	0.0585888	1251.6	0.351	0	0	0.00014018	ND	ND	ND	ND
2	rs11244049	17	136123092	T	C	0.99546	0.98526	907.73	314.42	29.845	0.008	0.14941	0.37435	0.05596	2.2379e-11	
3	rs146031614	17	136123172	T	C	0.99928	0.25135	1251.1	0.9	0	0	0.00035942	ND	ND	ND	
4	rs9328546	17	136123267	C	G	0.93562	0.84338	388.89	376.56	36.519	0.034	0.17956	-0.4023	0.055922	6.2907e-13	
5	rs114204311	17	136123277	G	A	0.99987	0.083469	1251.8	0.166	0	0	0	ND	ND	ND	
6	rs19246092	17	136123600	A	C	0.9999	0.01904	1251.9	0.131	0	0	0	ND	ND	ND	
7	rs183830736	17	136123679	C	T	0.99982	0.10392	1251.8	0.221	0	0	0	ND	ND	ND	
8	rs6597168	17	136123740	T	C	0.99414	0.98988	459.62	583.73	208.63	0.01	0.39976	-0.12847	0.040268	0.0014203	
9	rs4363269	17	136123840	A	G	1	1	634	517	101	0	0.28714	-0.39307	0.04419	5.8492e-19	
10	rs18609467	17	136123994	T	C	1	0.0029988	1252	0.003	0	0	0	ND	ND	ND	
11	rs7025839	17	136124190	G	A	0.99694	0.99335	703.13	474.76	74.109	0.005	0.24879	-0.4253	0.046445	5.3338e-20	
12	rs72775410	17	136124314	G	A	0.96768	0.3896	1204.1	47.501	0.434	0.002	0.019317	0.079419	0.23598	0.73645	
13	rs7362371	17	136124326	T	C	0.99955	0.72665	1250.2	1.76	0	0	0.00072088	-1.5209	0.9160	0.096852	
14	rs9411473	17	136124399	G	A	0.99562	0.98996	702.96	474.93	74.059	0.021	0.24885	-0.42732	0.046516	4.0619e-20	
15	rs9411370	17	136124432	A	G	0.99538	0.98938	702.97	474.94	74.071	0.015	0.24884	-0.42765	0.04653	3.8938e-20	
16	rs78755996	17	136124590	T	A	0.99747	0.075064	1248.8	3.171	0	0	0.0012664	ND	ND	ND	
17	rs115370334	17	136124602	T	G	0.99982	0.017178	1251.8	0.225	0	0	0	ND	ND	ND	
18	rs123380226	17	136124857	C	G	0.993	0.58542	1250.4	1.624	0	0	0.00064856	ND	ND	ND	
19	rs7047076	17	136125329	A	T	0.989	0.97695	702.56	475.81	73.599	0.031	0.24881	-0.43591	0.04683	1.2986e-20	
20	rs193302481	17	136125592	C	T	1	0.0009992	1252	0.002	0	0	0	ND	ND	ND	
21	rs9411474	17	136125716	C	G	0.98636	0.97321	702.37	476.16	73.432	0.031	0.24882	-0.43906	0.04693	8.3172e-21	
22	rs61527302	17	136125994	G	A	0.98778	0.97299	813.69	388.96	49.337	0.007	0.19474	-0.40888	0.050613	6.5466e-16	
23	rs191151099	17	136126026	T	A	1	1	1252	0	0	0	0	ND	ND	ND	
24	rs10901251	17	136126129	A	C	0.97626	0.96443	456.22	577.46	218.27	0.044	0.40497	-0.11346	0.040487	0.0050732	
25	rs148763789	17	136126132	C	T	0.99999	0.0058286	1252	0.012	0	0	0	ND	ND	ND	
26	rs188658949	17	136126153	G	T	1	0.0019992	1252	0.002	0	0	0	ND	ND	ND	
27	rs78320709	17	136126394	C	T	0.989	0.49772	1213.6	20.321	0.068	0.002	0.0081697	0.034413	0.28287	0.90498	
28	rs34266669	17	136126495	C	T	0.98406	0.96903	702.47	476.4	73.112	0.022	0.24866	-0.44461	0.047076	3.5709e-21	
29	rs151316644	17	136126625	G	A	0.99948	0.039007	1251.3	0.653	0	0	0.00026078	ND	ND	ND	
30	rs3205136	17	136126631	C	A	0.98673	0.96021	895.8	322.73	33.451	0.018	0.15561	0.41364	0.055894	1.3571e-13	
31	rs7849280	17	136126636	A	G	0.94202	0.87242	783.14	416.66	52.162	0.033	0.20807	-0.43438	0.052112	7.217e-16	
32	rs116958345	17	136127122	A	C	0.99974	0.013815	1251.7	0.33	0.162	0.002	0.00013179	ND	ND	ND	
33	rs9411475	17	136127268	T	C	0.95823	0.92214	649.81	507.3	94.676	0.044	0.27823	-0.43752	0.046433	4.4015e-21	
34	rs118065674	17	136127346	C	T	0.99999	0.0019982	1252	0.012	0	0	0	ND	ND	ND	
35	rs12216891	17	136127366	C	T	0.97965	0.94627	83.43	376.17	41.661	0.04	0.18351	0.45304	0.053218	1.6973e-17	
36	chr9:136127382::I	17	136127382	A	AGCCCT	0.99991	0.015581	1251.9	1.15	0	0	0	ND	ND	ND	
37	rs13291798	17	136127481	A	G	0.99227	0.25953	1242.2	9.761	0	0.002	0.0038982	-0.28934	0.413	0.48356	
38	rs186319566	17	136127534	C	A	1	1	1252	0	0	0	0	ND	ND	ND	
39	rs9411476	17	136127601	G	A	0.98081	0.95873	741.55	443.51	66.925	0.017	0.23058	-0.42418	0.047964	9.571e-19	
40	rs34085694	17	136127605	C	G	0.99157	0.24806	1241.4	10.634	0.003	0.004	0.0042492	-0.32526	0.41426	0.3942	
41	rs4962113	17	136127641	T	C	0.90386	0.84096	575.07	543.82	133.01	0.099	0.32345	0.10505	0.046389	0.023539	
42	rs112134200	17	136127689	A	G	0.98948	0.18823	1250.1	1.89	0	0	0.00075479	ND	ND	ND	
43	rs146579922	17	136127736	C	T	0.99748	0.11506	1248.8	3.154	0	0	0.0012596	ND	ND	ND	
44	rs115413527	17	136127737	G	A	0.98845	0.26706	1236.9	15.091	0	0	0.0060268	-0.19639	0.4121	0.26168	
45	rs62574563	17	136127764	C	T	0.96103	0.91922	705.84	484.63	92.006	0.071	0.26704	0.091014	0.046916	0.052387	
46	rs62574564	17	136127811	G	A	0.96476	0.92562	683.58	482.47	85.92	0.057	0.26132	0.081085	0.047165	0.085579	
47	rs73660468	17	136127907	G	A	0.99996	0.012682	1251.9	0.504	0	0	0	ND	ND	ND	
48	rs189931088	17	136127934	C	A	0.99995	0.033943	1251.9	0.058	0	0	0	ND	ND	ND	
49	rs34085694	17	136127956	A	C	0.99193	0.23459	1241.2	10.828	0.001	0.001	0.0043251	-0.31627	0.40805	0.43829	
50	rs10901252	17	136128000	G	C	0.99201	0.97662	838.98	369.68	43.30	0.017	0.18223	0.45201	0.052106	4.1441e-18	
51	rs141318395	17	136128175	G	A	0.99976	0.0084847	1251.7	0.299	0	0	0.00019141	ND	ND	ND	
52	rs62574565	17	136128259	G	A	0.98307	0.96009	706.68	467.13	78.136	0.054	0.24897	0.084741	0.047097	0.071975	
53	rs58081338	17	136128329	T	C	0.98293	0.96091	705.84	467.79	78.204	0.057	0.24933	0.084609	0.047075	0.072283	
54	rs62574567	17	136128421	C	G	0.98235	0.95978	706.68	467.6	78.094	0.05	0.24895	0.084946	0.0471	0.071302	
55	rs60484807	17	136128467	A	G	0.98558	0.97444	425.78	609.68	216.5	0.04	0.41642	0.36313	0.041006	8.3212e-19	
56	rs7875599	17	136128546	A	G	0.97543	0.94996	111.38	510.4	630.16	0.056	0.29281	-0.014598	0.044559	0.74321	
57	rs77693339	17	136128558	T	C	0.99453	0.98277	686.78	348.12	305.05	0.016	0.16705	0.51834	0.054001	8.0937e-22	
58	rs12554580	17	136128603	T	C	0.9858	0.97488	425.82	609.68	216.45	0.043	0.41630	0.36301	0.040996	3.8948e-19	
59	rs12554336	17	136128663	A	G	0.96931	0.93122	719.48	460.02	462.25	0.045	0.24159	0.092701	0.048318	0.050543	
60	rs62574565	17	136128664	T	C	0.96954	0.93141	720.32	459.28	73.357	0.049	0.24122	0.092846	0.048342	0.054783	
61	rs11244051	17	136128731	G	A	0.99583	0.14038	1246.8	5.212	0.003	0	0.0020839	ND	ND	ND	
62	rs62541784	17	136130665	G	A	0.91516	0.55494	1077.8	169.01	5.2	0.024	0.16705	0.36198	0.0409	8.7206e-19	
63	rs62641785	17	136130669	T	C	0.89227	0.71802	886.33	334.16	29.466	0.048	0.15699	0.07006	0.06539	6.5243e-22	
64	rs10751502	17	136129079	A	G	0.99154	0.18325	1240.6	11.35	0.007	0.007	0.00454	-0.90886	0.71079	0.20102	
65	rs35698328	17	136129112	C	T	0.99495	0.98413	868.71	348.31	35.078	0.004	0.16706	0.51799	0.053964	8.0877e-22	
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125	---	rs112655244	17	136132608	C	T	0.99795	0.99695	718.35	455.96	77.685	0.007	0.24414	0.077367	0.046072	0.093104
126	---	rs8176725	17	136132617	C	A	0.99526	0.99115	721.82	446.01	83.272	0	0.24409	0.21677	0.045652	0.20507e-06
127	rs2073824	rs2073824	17	136132633	A	G	1	1	327	609	316	0	0.49561	0.22886	0.039285	5.6917e-09
128	---	rs2073825	17	136132707	A	T	0.99883	0.99748	718.55	456.18	77.27	0.001	0.2439	0.077833	0.046096	0.091316
129	---	rs8176722	17	136132754	C	A	0.99867	0.99629	860.62	351.83	39.546	0.001	0.1721	0.4888	0.052404	1.0846-20
130	---	rs8176721	17	136132852	G	A	0.99934	0.062922	1251.2	0.82	0	0.001	0.00032748	ND	ND	ND
131	---	rs8176720	17	136132873	T	C	0.99964	0.9993	428.06	606.21	217.73	0	0.416	0.34913	0.04032	4.7599e-18
132	---	chr9:136132908:1	17	136132908	T	TC	0.99875	0.99767	387.31	595.14	269.54	0.002	0.45297	-0.067771	0.03922	0.025227
133	---	rs75179845	17	136132954	T	C	0.99743	0.99301	862.35	350.51	39.138	0.003	0.17124	0.488	0.052745	2.2012e-20
134	---	rs8176718	17	136132957	C	T	0.99986	0.99962	718.94	456.01	77.051	0	0.24365	0.078603	0.046101	0.088188
135	---	rs8176717	17	136133034	G	T	0.99933	0.99837	719.63	455.38	76.989	0	0.24335	0.078788	0.046125	0.087613
136	---	rs8176716	17	136133065	G	T	0.99863	0.15653	1250.3	72.17	0	0	0.00068371	ND	ND	ND
137	---	rs8176715	17	136133148	T	C	0.99802	0.97381	83.13	476.46	692.39	0.02	0.25668	-0.035566	0.045948	0.4389
138	---	rs191976170	17	136133149	G	A	0.99921	0.24201	1251	0.989	0	0	0.00039497	ND	ND	ND
139	---	rs8176714	17	136133178	G	A	0.99356	0.98601	648.13	505.84	98.016	0.01	0.2803	0.069039	0.044583	0.12149
140	---	rs115166192	17	136133218	C	T	0.99965	0.0086304	1251.6	0.441	0	0	0.00017612	ND	ND	ND
141	---	rs18295414	17	136133346	A	C	1	1	1252	0	0	0	0	ND	ND	ND
142	---	chr9:136133380:D	17	136133380	CGGG	C	0.99867	0.99649	76.608	455.3	720.09	0.004	0.24302	-0.07819	0.046199	0.090558
143	---	rs45610939	17	136133396	A	G	0.9792	0.8157	1146.1	105.73	200	0.003	0.042387	-0.055135	0.10946	0.61446
144	rs512770	rs512770	17	136133506	A	G	1	1	77	456	719	0	0.24361	-0.078689	0.046104	0.087862
145	---	rs4962040	17	136133531	A	G	0.99764	0.99606	629.73	513.26	109.01	0.008	0.29204	0.028954	0.043672	0.50734
146	---	chr9:136133582:D	17	136133582	AG	A	0.99258	0.98658	91.997	483.41	676.59	0.002	0.26654	-0.12006	0.045053	0.0077009
147	---	rs641959	17	136133699	C	A	0.99758	0.99575	89.28	468.68	694.07	0.001	0.25846	-0.092724	0.045021	0.039439
148	---	rs641943	17	136133714	G	A	0.99778	0.99624	89.168	468.59	694.24	0.001	0.25836	-0.09265	0.045023	0.039609
149	---	rs514708	17	136133743	T	C	0.99778	0.99625	89.168	468.59	694.24	0	0.25836	-0.092649	0.045023	0.039611
150	---	rs517414	17	136134034	A	G	0.9996	0.99919	80.042	459.41	712.54	0	0.2474	-0.066682	0.045827	0.058558
151	---	rs9411372	17	136134068	G	A	0.97122	0.94646	730.70	444.16	77.763	0.012	0.23949	-0.4368	0.047436	3.3133e-20
152	---	rs138164693	17	136134118	G	A	1	1	1252	0	0	0	0	ND	ND	ND
153	---	rs638756	17	136134472	C	A	0.99787	0.99641	89.116	468.64	694.24	0	0.25834	-0.092673	0.045026	0.039568
154	---	rs191050088	17	136134650	A	G	0.99671	0.14604	1247.9	4.122	0	0	0.0016462	ND	ND	ND
155	---	rs183228393	17	136134835	G	A	1	0.0033309	1252	0.006	0	0	0	ND	ND	ND
156	---	rs626792	17	136134964	C	A	0.93522	0.87801	103.3	510.59	637.49	0.026	0.2869	-0.051962	0.046684	0.26568
157	---	rs626035	17	136134994	G	T	0.97273	0.94743	99.363	483.5	668.14	0.006	0.27245	-0.1037	0.045359	0.022241
158	---	rs547495	17	136135095	C	T	0.97273	0.94743	99.363	483.49	668.14	0.006	0.27245	-0.1037	0.045359	0.022241
159	---	rs547643	17	136135047	T	C	1	1	80	459	713	0	0.2472	-0.054541	0.045802	0.061979
160	---	rs625593	17	136135096	A	G	0.99988	0.99995	80.002	459.02	712.98	0	0.24721	-0.054542	0.045802	0.062026
161	---	rs8176707	17	136135108	G	T	0.93611	0.7539	1021.6	214.61	15.814	0.015	0.098338	-0.045788	0.075941	0.54655
162	---	chr9:136135137:D	17	136135137	GCTTCTGTCCACAC	G	0.98818	0.97538	85.311	466.49	700.19	0.007	0.25444	-0.077415	0.045792	0.09092
163	---	rs13874634	17	136135194	G	A	0.99916	0.84615	1246.3	5.664	0	0	0.002626	0.049472	0.43827	0.91013
164	---	rs549331	17	136135195	G	A	0.99799	0.99664	89.053	468.71	694.24	0	0.25831	-0.092707	0.045029	0.039509
165	---	rs549443	17	136135237	A	G	0.88915	0.79851	1205.9	528.03	605.05	0.005	0.30711	-0.13429	0.048546	0.0056698
166	---	rs549448	17	136135238	T	C	0.88939	0.79851	130.45	531.66	589.83	0.055	0.31653	-0.13067	0.04747	0.0061278
167	---	rs14279023	17	136135321	G	A	1	1	1252	0	0	0	0	ND	ND	ND
168	---	rs624601	17	136135365	A	G	0.99799	0.99664	89.053	468.71	694.24	0	0.25831	-0.092707	0.045029	0.039509
169	---	rs551322	17	136135444	G	A	1	1	80	459	713	0	0.2472	-0.054541	0.045802	0.061997
170	---	rs192192638	17	136135445	C	T	0.96586	0.37961	1202.7	49.017	258.08	0.001	0.019782	1.0383	0.26064	6.7827e-05
171	---	rs188088103	17	136135472	C	T	0.99846	0.024321	1250.1	1.927	0	0.002	0.00076957	ND	ND	ND
172	---	rs6134243	17	136135478	A	G	1	1	80	459	713	0	0.2472	-0.054541	0.045802	0.061997
173	---	rs8176704	17	136135552	G	A	0.99874	0.1339	1250.4	1.574	0	0	0.00062859	ND	ND	ND
174	---	chr9:136135625:D	17	136135625	CTG	C	1	1	1252	0	0	0	0	ND	ND	ND
175	rs574347	rs574347	17	136135659	C	T	1	1	80	459	713	0	0.2472	-0.054541	0.045802	0.061997
176	---	rs575259	17	136135752	G	A	0.99798	0.99664	89.052	468.71	694.24	0.001	0.25831	-0.092713	0.045029	0.039497
177	---	rs8176703	17	136135863	G	T	0.99884	0.21781	1250.7	1.35	0	0	0.00053914	ND	ND	ND
178	---	rs8176702	17	136136146	G	A	0.99883	0.99712	630.2	512.79	109.01	0.001	0.29186	0.029119	0.043645	0.50466
179	---	rs579483	17	136136196	A	T	0.99999	0.99988	80	459.01	712.99	0	0.24721	-0.085506	0.045803	0.061924
180	---	rs579622	17	136137050	G	A	0.99882	0.21075	1249.7	2.259	0	0	0.00090216	ND	ND	ND
181	---	rs140166545	17	136137078	C	T	0.99796	0.99735	378.17	594.14	279.69	0.002	0.46067	-0.085673	0.03911	0.028483
182	---	rs493211	17	136137088	G	A	0.99996	0.020119	1252	0.044	0	0	0	ND	ND	ND
183	---	rs687289	17	136137133	G	C	1	1	609	527	116	0	0.30312	0.037734	0.043232	0.38275
192	rs2073828	rs2073828	17	136137140	G	A	0.99997	0.019153	1252	0.036	0	0	0	ND	ND	ND
193	---	rs189048957	17	136137228	G	A	1	1	1252	0	0	0	0	ND	ND	ND
194	---	rs150021186	17	136137293	G	A	1	1	1252	0	0	0	0	ND	ND	ND
195	---	rs77513793	17	136137296	T	C	0.99874	0.38576	1249.6	2.428	0	0	0.00096965	ND	ND	ND
196	---	chr9:136137375:L	17	136137375	A	AT	1	1	1252	0	0	0	0	ND	ND	ND
197	---	rs55876802	17	136137547	C	A	1	1	1252	0	0	0	0	ND	ND	ND
198	---	rs143174772	17	136137600	C	T	1	1	1252	0	0	0	0	ND	ND	ND
199	---	rs15115														

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252	---	rs8176669	17	136142463	A	C	0.99711	0.99461	707.9	460.87	83.226	0.005	0.25053	0.09383	0.045625	0.039729	
253	---	rs148087354	17	136142465	G	A	0.9999	0.02062	1251.0	0.128	0	0	0	ND	ND	ND	
254	---	chr9:136142521:1	17	136142521	C	CT	0.99467	0.89609	1204.1	47.878	0	0.004	0.019121	-0.50178	0.15256	0.001056	
255	---	chr13:136142540:1	17	136142540	T	TTG	1	1	1252	0	0	0	0	ND	ND	ND	
256	---	rs143746661	17	136142711	A	G	1	1	1252	0	0	0	0	ND	ND	ND	
257	---	rs187821258	17	136142968	T	C	0.99742	0.11134	1248.8	3.226	0	0	0.0012883	ND	ND	ND	
258	---	rs543040	17	136143000	A	T	0.99562	0.9934	396.75	589.54	265.71	0.001	0.44767	-0.10259	0.039183	0.0088412	
259	---	rs146903770	17	136143055	A	G	0.99111	0.28893	1204.3	11.683	0	0	0.0046657	-1.2328	0.50413	0.014472	
260	---	rs613534	17	136143120	A	G	0.99563	0.99385	399.27	589.45	263.29	0.001	0.4457	-0.10427	0.039263	0.007913	
261	---	rs543968	17	136143121	T	C	0.99437	0.9912	400.49	588.95	262.56	0.002	0.44492	-0.10659	0.039365	0.0067754	
262	---	rs139081859	17	136143192	G	T	0.99825	0.077245	1249.8	2.196	0	0.001	0.000877	ND	ND	ND	
263	---	rs548473	17	136143212	G	A	0.99502	0.9917	402.45	586.52	263.02	0	0.44432	-0.10442	0.039194	0.0077153	
264	---	rs142956930	17	136143330	A	G	0.99889	0.098024	1250.6	1.391	0	0	0.00055551	ND	ND	ND	
265	---	rs545971	17	136143372	C	T	0.99838	0.99763	397.73	589.42	264.85	0	0.44693	-0.10225	0.039161	0.0090282	
266	---	rs138194969	17	136143391	G	A	1	1	1252	0	0	0	0	ND	ND	ND	
267	---	rs612169	17	136143442	A	G	0.99839	0.99764	397.73	589.42	264.85	0	0.44693	-0.10225	0.03916	0.0090285	
268	---	rs144566139	17	136143919	G	A	1	1	1252	0	0	0	0	ND	ND	ND	
269	---	chr9:136144000:D	17	136144000	AT	A	0.98756	0.54188	1223.5	28.432	0.053	0.001	0.011397	0.41596	0.25184	0.098597	
270	---	rs8176668	17	136144059	A	T	0.99815	0.9973	611.89	523.75	116.36	0	0.3021	0.039923	0.043211	0.35553	
271	---	rs574311	17	136144110	G	A	0.99799	0.99662	83.583	461.25	707.17	0	0.25096	-0.093923	0.0455593	0.039165	
272	---	rs187551277	17	136144172	G	A	0.99825	0.077245	1249.8	2.196	0	0.001	0.000877	ND	ND	ND	
273	---	rs597988	17	136144284	T	A	0.97199	0.95467	459.93	572.81	219.04	0.019	0.40388	-0.092999	0.040572	0.021895	
274	---	rs597974	17	136144297	A	G	0.96877	0.94921	462.62	571.57	217.79	0.018	0.40222	-0.093592	0.040683	0.021418	
275	---	rs140114497	17	136144308	T	C	0.96718	0.9468	463.77	571.11	217.11	0.017	0.40149	-0.096203	0.040787	0.01834	
276	---	rs576125	17	136144309	G	A	0.96718	0.94681	463.77	571.1	217.12	0.017	0.4015	-0.0962	0.040786	0.018343	
277	---	rs8176663	17	136144427	T	C	0.99836	0.97367	397.61	589.5	268.49	0	0.447	-0.10235	0.039164	0.0089644	
278	---	rs8176662	17	136144454	A	G	0.99669	0.9908	874.92	341.45	35.619	0.004	0.16481	0.48185	0.053637	2.6219e-19	
279	---	rs149899707	17	136144521	C	T	1	1	1252	0	0	0	0	ND	ND	ND	
280	---	rs488775	17	136144534	G	A	0.99797	0.99644	83.591	461.14	707.27	0.001	0.25093	-0.094063	0.045543	0.038888	
281	---	rs66697526	17	136144593	G	T	0.99595	0.98605	95.35	272.93	207.7	0.017	0.12554	0.061971	0.060295	0.30405	
282	---	rs7036642	17	136144626	G	A	0.9981	0.99718	61.196	523.69	116.36	0	0.30208	0.039927	0.043211	0.35549	
283	---	rs8176661	17	136144640	A	G	0.99996	0.086711	1252	0.045	0	0	0	ND	ND	ND	
284	---	rs596141	17	136144689	T	G	0.99816	0.99701	79.694	461.37	710.94	0	0.24791	-0.083621	0.045896	0.068458	
285	---	rs146907040	17	136144771	A	G	1	0	0.0039984	1252	0.004	0	0	0	ND	ND	ND
286	---	rs491626	17	136144873	C	T	0.99109	0.98649	411.75	583.83	256.42	0.001	0.43797	-0.11815	0.039366	0.0026882	
287	---	rs492488	17	136144960	G	A	0.99656	0.99462	391.81	591.2	269	0	0.45098	-0.094233	0.039206	0.016237	
288	---	rs493246	17	136144994	G	A	0.98929	0.982	384.57	593.12	274.3	0.01	0.45598	-0.097317	0.039427	0.013576	
289	---	rs192389831	17	136145029	C	T	0.99979	0.10628	1251.7	0.261	0	0	0.00010423	ND	ND	ND	
290	---	rs494242	17	136145118	C	T	0.99567	0.99328	391.13	591.58	269.29	0.001	0.45134	-0.094146	0.039244	0.016441	
291	---	rs495203	17	136145240	C	T	0.99567	0.99328	391.13	591.58	269.29	0.001	0.45134	-0.094146	0.039244	0.016441	
292	---	chr9:136145403:D	17	136145403	AC	A	0.88421	0.82553	562.29	557.9	132.08	0.005	0.32819	-0.27512	0.047402	6.4719e-09	
293	---	chr9:136145414:D	17	136145414	C	CA	0.90476	0.80485	1022.7	217.57	11.653	0.032	0.098197	0.53079	0.090195	3.9815e-09	
294	---	chr9:136145418:D	17	136145418	AC	A	0.88378	0.80486	584.03	533.39	134.53	0.004	0.32048	-0.22009	0.047346	3.3413e-06	
295	---	chr9:136145424:D	17	136145424	AC	A	0.96664	0.92861	684.01	483.64	84.323	0.003	0.2605	-0.46763	0.04662	1.655e-23	
296	---	rs9411378	17	136145425	C	A	0.96466	0.92381	684.98	482.46	84.528	0.003	0.2602	-0.46476	0.046865	3.5102e-23	
297	---	rs582118	17	136145471	A	G	0.99785	0.99665	392.39	591.52	268.09	0	0.45036	-0.093554	0.039192	0.016983	
298	---	rs5820918	17	136145484	A	T	0.99785	0.99665	392.39	591.52	268.09	0	0.45036	-0.093554	0.039192	0.016983	
299	---	rs189022067	17	136145717	A	G	0.99986	0.035747	1251.8	1.079	0	0	0	ND	ND	ND	
300	---	chr9:136145907:D	17	136145907	G	GA	0.99609	0.99346	395.19	590.05	266.79	0.001	0.44874	-0.092758	0.03919	0.017939	
301	---	rs2769071	17	136145974	A	G	0.99601	0.99357	392.38	591.18	268.44	0.004	0.4505	-0.09383	0.039183	0.016635	
302	---	rs147027134	17	136145979	A	G	1	1	1252	0	0	0	0	ND	ND	ND	
303	---	rs677355	17	136146046	G	A	0.99691	0.99484	394.01	590.7	267.29	0.003	0.44939	-0.093172	0.039196	0.017449	
304	---	chr9:136146061:D	17	136146061	T	TA	0.99429	0.99034	398.01	585.8	268.18	0.009	0.44815	-0.096895	0.039108	0.013225	
305	---	chr9:136146068:D	17	136146068	A	AT	0.99726	0.99542	397.33	590.73	267.49	0.005	0.44957	-0.093428	0.039176	0.017087	
306	---	rs676969	17	136146077	T	G	0.99628	0.99354	395.58	595.54	271.11	0.005	0.44719	-0.096516	0.030907	0.01354	
307	---	rs676457	17	136146227	A	T	0.99788	0.99613	79.842	461.53	710.62	0.001	0.44962	-0.093651	0.039154	0.016786	
308	---	rs473533	17	136148035	T	C	0.99828	0.99718	270.49	597.73	383.79	0.002	0.45475	-0.078192	0.032924	0.02116e-20	
309	---	rs475419	17	136148231	C	T	0.99867	0.99765	270.55	597.66	383.79	0	0.45478	-0.078008	0.032821	0.047046	
310	---	chr9:136147630:D	17	136147630	G	GA	0.99796	0.98811	82.173	466.32	703.51	0.005	0.25186	-0.095002	0.045949	0.038682	
311	---	rs675201	17	136147662	T	G	0.99796	0.99816	61.715	461.51	710.61	0.002	0.44811	-0.083391	0.045889	0.069184	
312	---	rs7046674	17	136147672	C	T	0.99716	0.99473	61.82	523.29	115.82	0.004	0.30149	0.040452	0.043295	0.35012	
313	---	rs554833	17	136147760	C	T	0.99848	0.99822	393.07	591.1	267.83	0.001	0.44998	-0.093501	0.039145	0.016913	
314	---	rs674302	17	136147798	T	C	0.99727	0.99545	634.51	508.13	271.29	0	0.28153	-0.46233	0.044317	1.7678e-25	
315	---	rs551100	17	136147808	T	C	0.99788	0.0009996	125.00	0.001	0	0	0				

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378	---	rs61696148	17	136153544	A	T	0.99658	0.98329	1026	216.96	9.047	0.001	0.093873	0.059061	0.069996	0.39879
379	---	rs141638449	17	136153707	C	T	0.99874	0.036203	1260.4	1,576	0	0	0.0062030	ND	ND	ND
380	---	rs651007	17	136153875	C	T	0.99996	0.9999	647.01	504.99	100	0	0.28155	-0.46236	0.044194	1.2925e-25
381	---	rs11244061	17	136153981	C	T	0.99201	0.57166	1234.7	17.302	0	0	0.0069097	-0.47242	0.30302	0.11688
382	---	rs187861744	17	136154097	G	C	0.99827	0.51738	1248.4	3.554	0	0	0.0014193	4.8627	1.0473	0.0001943
383	---	rs579459	17	136154168	T	C	0.99997	0.99993	647.01	504.99	100	0	0.28155	-0.46236	0.044194	1.2893e-25
384	---	rs9411488	17	136154246	G	T	0.99751	0.99417	805.67	381.78	64.577	0	0.20404	0.040535	0.048138	0.39975
385	---	rs649129	17	136154304	C	T	0.99998	0.99995	647.01	504.99	100.01	0	0.28155	-0.46236	0.044193	1.2891e-25
386	---	rs10751504	17	136154433	G	C	0.9682	0.93236	708.84	473.58	69.535	0.044	0.24468	0.38052	0.049077	8.9352e-15
387	---	chr9:136154623:D	17	136154623	T	I	0.99812	0.12397	1249.6	2.357	0	0	0.00094129	ND	ND	ND
388	---	chr9:136154626:D	17	136154626	CA	C	0.99812	0.12397	1249.6	2.357	0	0	0.00094129	ND	ND	ND
389	---	rs147375567	17	136154866	A	G	0.99898	0.50891	1249.9	2.111	0	0	0.00084305	-0.24873	1.0283	0.80887
390	rs495828	rs495828	17	136154867	G	T	1	1	647	505	100	0	0.28155	-0.46238	0.044194	1.288e-25
391	---	rs635634	17	136155000	C	T	0.99998	0.99995	647.01	504.98	100.01	0	0.28155	-0.46236	0.044193	1.288e-25
392	---	rs12683493	17	136155063	C	T	0.99649	0.98298	1026	216.93	9.046	0.003	0.093856	0.05928	0.07002	0.39721
393	---	rs150022816	17	136155093	G	T	1	1	1252	0	0	0	0	ND	ND	ND
394	---	rs13298002	17	136155127	G	A	0.99811	0.12465	1249.6	2.365	0	0	0.00094449	ND	ND	ND
395	rs500428	rs500428	17	136155343	A	G	1	1	90	474	688	0	0.26118	-0.066892	0.04489	0.052908
396	---	rs7030248	17	136155359	G	A	0.99743	0.99482	692.55	484.83	74.628	0	0.25323	0.38089	0.046505	2.6058e-16
397	---	rs9411490	17	136155414	A	G	0.99744	0.99405	805.68	381.77	64.549	0.003	0.20402	0.040483	0.048145	0.40043
398	---	rs633862	17	136155444	C	T	0.99743	0.99608	279.07	606.76	366.17	0.002	0.46522	-0.094595	0.039506	0.016647
399	---	rs502361	17	136155589	C	G	0.99994	0.99985	89.94	473.97	688.04	0	0.26116	-0.066917	0.044892	0.052853
400	---	rs9411491	17	136156015	T	C	0.99715	0.99328	805.67	381.78	64.553	0.004	0.20403	0.040355	0.048159	0.40206
401	---	rs55988407	17	136156064	G	A	0.9809	0.47025	1218	33.85	0.149	0.003	0.013631	0.44056	0.24649	0.073883
402	---	rs9411381	17	136156067	T	C	0.99683	0.99246	805.65	381.78	64.556	0.017	0.20403	0.040234	0.048177	0.40365
403	---	rs78590974	17	136156230	C	T	0.98471	0.47114	1226.6	25.202	0.241	0.002	0.010257	2.0613	0.3907	1.3213e-07
404	---	rs142447342	17	136156244	G	A	1	1	1252	0	0	0	0	ND	ND	ND
405	---	rs140502769	17	136156498	G	A	0.99999	0.0094373	1252	0.018	0	0	0	ND	ND	ND
406	---	rs144306320	17	136156542	G	T	0.999	0.013524	1250.7	1.255	0	0.001	0.0005012	ND	ND	ND
407	---	rs189611925	17	136156672	G	T	0.9998	0.016521	1251.8	0.246	0	0	0	ND	ND	ND
408	---	rs557317	17	136157037	C	A	0.98391	0.9689	97.661	488.34	665.97	0.026	0.27303	-0.09539	0.045036	0.034167
409	---	rs558240	17	136157133	G	A	0.99025	0.97154	895.23	333.18	23.573	0.012	0.15189	-0.10138	0.057695	0.078881
410	---	rs138026250	17	136157138	C	A	0.99982	0.13924	1251.8	0.224	0	0	0	ND	ND	ND
411	---	rs146149084	17	136157335	A	G	0.99993	0.015153	1252	0.023	0	0	0	ND	ND	ND
412	---	rs181316261	17	136157449	C	T	0.99919	0.14459	1251	1.009	0	0	0.00040296	ND	ND	ND
413	---	rs7867220	17	136157520	C	T	0.99949	0.11798	1251.4	0.64	0	0	0.00025559	ND	ND	ND
414	---	rs187397696	17	136157596	C	T	0.99988	0.032044	1251.8	0.154	0	0	0	ND	ND	ND
415	---	rs142673851	17	136157644	G	T	0.9994	0.14465	1251.3	0.746	0	0	0.00029792	ND	ND	ND
416	---	rs62574607	17	136157654	T	C	0.99069	0.87367	1181.9	67.059	3.032	0.002	0.029203	0.17883	0.12453	0.15097
417	---	rs7673539	17	136157656	C	G	0.99945	0.12814	1251.3	0.687	0	0	0.00027436	ND	ND	ND
418	---	rs7853493	17	136157785	A	T	0.99945	0.14861	1251.3	0.69	0	0	0.00027556	ND	ND	ND
419	---	rs786739	17	136157924	C	T	0.99942	0.15659	1251.3	0.726	0	0	0.00028994	ND	ND	ND
420	---	rs7867664	17	136158008	C	G	0.99936	0.15691	1251.2	0.8	0	0	0.00031949	ND	ND	ND
421	---	rs141952207	17	136158026	G	C	0.99996	0.008981	1252	0.048	0	0	0	ND	ND	ND
422	---	rs149863406	17	136158155	C	T	0.99918	0.013745	1251	1.032	0	0	0.00041214	ND	ND	ND
423	---	rs28467494	17	136158288	T	C	0.99928	0.18331	1251.1	0.9	0	0	0.00035942	ND	ND	ND
424	---	rs7860182	17	136158327	T	A	0.98662	0.97544	98.488	491.58	661.91	0.02	0.27499	-0.090135	0.044889	0.044647
425	---	rs28368280	17	136158400	C	G	0.99925	0.1935	1251.1	0.942	0	0	0.0003762	ND	ND	ND
426	---	rs184855103	17	136158490	C	T	0.99998	0.009075	1250.7	0.229	0	0	0	ND	ND	ND
427	---	rs1927311	17	136158544	T	C	0.9874	0.97617	104.32	494.88	652.79	0.01	0.28096	-0.094887	0.044506	0.033006
428	---	rs115017869	17	136158862	G	A	0.99911	0.23694	1250.9	1.112	0	0	0.0004409	ND	ND	ND
429	---	rs507480	17	136158964	A	G	0.99937	0.989	3	59.69	118.93	0.001	0.026234	-0.13081	0.12101	0.27972
430	---	rs507567	17	136158998	A	G	0.99705	0.38212	0.015	5.06	1246.9	0.003	0.020328	-0.51462	0.57591	0.37155
431	---	rs509482	17	136159240	G	A	0.98604	0.96899	724.2	454.3	73.46	0.029	0.24012	0.067143	0.047084	0.15386
432	---	rs62574608	17	136159266	A	G	0.92354	0.50729	1110.6	135.9	5.479	0.02	0.058652	0.22538	0.11535	0.050713
433	---	chr9:136159268:D	17	136159268	AGGCC	A	0.92738	0.62775	1063.2	181.65	7.148	0.013	0.078252	0.0096289	0.039572	0.91804
434	---	rs146582653	17	136159450	G	C	0.99949	0.4148	1251.3	0.684	0	0	0.00027316	ND	ND	ND
435	---	rs67318122	17	136159538	A	G	0.9957	0.27916	1246.6	5.365	0.013	0	0.002153	3.5202	0.73093	0.16276-06
436	---	rs512179	17	136159540	C	G	0.99989	0.997	703.95	467.03	81.018	0.001	0.25122	0.077437	0.045708	0.090234
437	---	rs51287	17	136159579	G	A	0.9999	0.99973	703.96	467.03	81.016	0	0.25122	0.077438	0.045708	0.090232
438	---	rs535855	17	136159790	A	G	0.99996	0.99988	703.98	467.01	81.008	0	0.25121	0.077425	0.045705	0.090264
439	---	rs535993	17	136159843	A	G	0.99996	0.99989	703.98	467.01	81.007	0	0.25121	0.077422	0.045705	0.090276
440	---	chr9:136159892:I	17	136159892	TAA	A	0.90366	0.70227	949	287.24	15.747	0.016	0.12729	-0.05678	0.073102	0.43733
441	---	chr9:136159896:I	17	136159896	TAAAGAAATAA	A	0.90474	0.70343	954.66	281.97	15.353	0.016	0.12487	-0.077352	0.073269	0.29109
442	---	rs184760764	17	136160131	G	A	0.999	0.023915	1250.7	1.254	0	0	0.0005009	ND	ND	ND
443	---	rs146174278	17	136160165	G	A	1	1	704	467	81	0	0.2512	0.07740		

Table S3. Associations of ABO gene in 9q32 with P-LIP or ACE levels after adjustment by smoking or Box-Cox transformation of trait values

SNPs (position)	stage*	minor allele	MAF	P-LIP						ACE					
				adjusted by alcohol intake			Box-Cox transformation***			adjusted by smoking			Box-Cox transformation***		
				beta**	SE	p value	beta**	SE	p value	beta**	SE	p value	beta**	SE	p value
rs4363269 (136123840)	1st	G	0.287	0.0849	0.500	> 1e-7	0.00272	0.003	> 1e-7	-2.01	0.218	9.85e-20	-1.12	0.125	1.20e-18
	2nd	G	0.269	-0.220	0.511	> 1e-7	0.000446	0.003	> 1e-7	-1.99	0.192	1.91e-24	-1.14	0.106	3.56e-26
(136131188)	1st	A	0.161	4.18	0.601	5.55e-12	0.0336	0.004	1.85e-16	2.57	0.266	2.67e-21	1.46	0.153	5.77e-21
	2nd	A	0.188	5.12	0.576	1.61e-18	0.0396	0.004	6.45e-27	2.13	0.219	7.15e-22	1.28	0.123	1.03e-24
(136131322)	1st	A	0.167	4.16	0.593	3.75e-12	0.0476	0.004	4.35e-17	2.58	0.261	3.22e-22	1.47	0.15	7.39e-22
	2nd	A	0.198	5.42	0.562	2.04e-21	0.0580	0.004	3.10e-30	2.11	0.214	2.74e-22	1.26	0.12	8.67e-25
(136132633)	1st	G	0.496	1.29	0.447	> 1e-7	0.00990	0.004	> 1e-7	1.17	0.199	6.25e-9	0.660	0.114	8.21e-9
	2nd	A	0.475	-1.82	0.462	> 1e-7	-0.0127	0.003	> 1e-7	-1.17	0.176	4.00e-11	-0.622	0.099	3.68e-10
(136139265)	1st	A	0.440	2.75	0.440	5.63e-10	0.0225	0.003	4.97e-14	-0.593	0.201	> 1e-7	-0.324	0.115	> 1e-7
	2nd	A	0.454	3.38	0.455	1.81e-13	0.0276	0.003	2.47e-21	-0.685	0.178	> 1e-7	-0.417	0.099	> 1e-7
(136148647)	1st	A	0.455	-2.71	0.438	8.70e-10	-0.0215	0.003	3.56e-13	0.371	0.2	> 1e-7	0.228	0.114	> 1e-7
	2nd	A	0.445	-3.01	0.456	5.77e-11	-0.0247	0.003	2.28e-17	0.428	0.179	> 1e-7	0.260	0.099	> 1e-7
(136149229)	1st	G	0.450	2.62	0.433	1.92e-9	0.0220	0.003	8.64e-14	-0.503	0.199	> 1e-7	-0.274	0.114	> 1e-7
	2nd	G	0.458	3.57	0.452	4.97e-15	0.0294	0.003	3.02e-24	-0.630	0.178	> 1e-7	-0.389	0.099	> 1e-7
(136154867)	1st	A	0.281	0.536	0.499	> 1e-7	0.00473	0.003	> 1e-7	-2.34	0.215	2.14e-26	-1.31	0.123	2.30e-25
	2nd	A	0.264	-0.0315	0.509	> 1e-7	0.00214	0.003	> 1e-7	-2.37	0.188	9.84e-35	-1.39	0.104	1.80e-38
(136166346)	1st	A	0.415	-0.834	0.445	> 1e-7	-0.00565	0.003	> 1e-7	-1.50	0.197	5.83e-14	-0.810	0.113	1.11e-12
	2nd	A	0.398	-0.667	0.467	> 1e-7	-0.00632	0.003	> 1e-7	-1.27	0.178	1.52e-12	-0.738	0.099	1.29e-13

* 1st population from the Takahata cohort, 2nd from the Yamagata cohort

** Regression coefficient

*** Optimal lambda value in Box-Cox transformation for P-LIP is -0.4 for all except rs8176746 that has -0.3, whereas that for ACE is 0.8 for all SNPs.

Table S4. Associations of ACE1 in 17q23.3 with ACE level after adjustment by smoking or Box-Cox transformation of trait values

SNPs (position)	stage*	minor allele	MAF	ACE			Box-Cox transformation****		
				adjusted by smoking			Box-Cox transformation****		
				beta**	SE	p value	beta**	SE	p value
rs4459609 (61548948)	1st	C	0.371	3.06	0.192	3.91e-52	1.82	0.110	3.87e-56
	2nd	C	0.356	2.48	0.177	8.81e-42	1.46	0.098	1.28e-47
rs4309 (61559923)	1st	G	0.426	3.31	0.181	1.03e-65	1.95	0.104	3.83e-69
	2nd	G	0.409	3.27	0.163	1.52e-76	1.87	0.091	7.51e-85
rs4311 (61560763)	1st	A	0.349	3.25	0.189	5.08e-59	1.91	0.109	4.21e-62
	2nd	A	0.346	2.54	0.180	2.99e-42	1.53	0.099	2.80e-50
rs4329 (61563458)	1st	A	0.358	3.25	0.189	2.10e-59	1.92	0.108	2.45e-63
	2nd	A	0.342	2.64	0.176	1.00e-47	1.58	0.097	3.11e-55
rs4343 (61566031)	1st	G	0.360	3.23	0.189	9.54e-59	1.91	0.108	1.09e-62
	2nd	G	0.342	2.65	0.176	6.93e-48	1.58	0.097	2.03e-55
rs4353 (61570422)	1st	A	0.440	3.91	0.173	4.09e-94	2.32	0.098	6.72e-102
	2nd	A	0.422	3.51	0.158	1.07e-94	2.04	0.088	4.47e-104
rs4362 (61573761)	1st	A	0.442	3.94	0.173	1.55e-95	2.33	0.098	3.43e-103
	2nd	A	0.426	3.54	0.157	8.97e-97	2.05	0.087	5.77e-106
rs4461142 (61578048)	1st	A	0.434	2.00	0.194	6.56e-24	1.16	0.111	7.77e-25
	2nd	A	0.433	1.87	0.170	6.41e-27	1.09	0.095	5.04e-29

* 1st population from the Takahata cohort, 2nd from the Yamagata cohort

** Regression coefficient

*** Coefficient t-statistic

**** Optimal lambda value for Box-Cox transformation is 0.8 for all SNPs.

Table S5. Associations of ABO gene in 9q32 with P-LIP or ACE levels after adjustment or subgroup by treatment

SNPs (position)	P-LIP								ACE							
	adjusted by treatment history*				subgroup of notreatment				adjusted by treatment history*				subgroup of notreatment			
	stage**	beta***	SE	p value	stage**	beta***	SE	p value	stage**	beta***	SE	p value	stage**	beta***	SE	p value
rs4363269 (136123840)	1st	-0.0325	0.481	> 1e-7	comb	0.525	0.443	> 1e-7	1st	-1.96	0.212	9.62e-20	comb	-1.89	0.160	3.07e-31
	2nd	-0.343	0.564	> 1e-7					2nd	-1.94	0.194	6.18e-23				
rs8176749 (136131188)	1st	4.40	0.575	3.92e-14	comb	4.04	0.524	2.15e-14	1st	2.52	0.258	1.24e-21	comb	2.30	0.192	9.06e-32
	2nd	5.40	0.635	4.53e-17					2nd	2.35	0.223	4.14e-25				
rs8176746 (136131322)	1st	4.37	0.568	2.79e-14	comb	4.17	0.516	1.32e-15	1st	2.53	0.254	1.83e-22	comb	2.32	0.189	3.00e-33
	2nd	5.69	0.620	1.41e-19					2nd	2.30	0.218	3.23e-25				
rs657152 (136139265)	1st	2.65	0.424	5.38e-10	comb	3.01	0.401	7.37e-14	1st	-0.550	0.195	> 1e-7	comb	-0.680	0.151	> 1e-7
	2nd	3.49	0.506	7.98e-12					2nd	-0.651	0.182	> 1e-7				
rs500498 (136148647)	1st	-2.44	0.421	8.35e-9	comb	-2.56	0.399	1.71e-10	1st	0.362	0.194	> 1e-7	comb	0.386	0.151	> 1e-7
	2nd	-3.02	0.507	3.16e-9					2nd	0.376	0.182	> 1e-7				
rs505922 (136149229)	1st	2.53	0.419	2.22e-9	comb	3.02	0.394	3.25e-14	1st	-0.454	0.193	> 1e-7	comb	-0.598	0.149	> 1e-7
	2nd	3.64	0.501	5.85e-13					2nd	-0.597	0.180	> 1e-7				
rs495828 (136154867)	1st	0.250	0.481	> 1e-7	comb	0.640	0.446	> 1e-7	1st	-2.25	0.209	7.13e-26	comb	-2.32	0.157	2.23e-46
	2nd	-0.182	0.562	> 1e-7					2nd	-2.38	0.190	2.07e-34				
rs7025162 (136166346)	1st	-0.806	0.428	> 1e-7	comb	-0.230	0.402	> 1e-7	1st	-1.41	0.191	2.85e-13	comb	-1.22	0.148	2.12e-16
	2nd	-0.678	0.510	> 1e-7					2nd	-1.29	0.178	7.02e-13				

* Treatment history is coded by 0 for untreated, 1 for treated or NA for not available.

** 1st population from the Takahata cohort and 2nd from the Yamagata cohort, whereas comb is combined populations of certainly untreated individuals in both cohorts.

*** Regression coefficient

Table S6. Associations of ACE1 gene in 17q23.2 with ACE level after adjustment or subgroup by treatment

SNPs (position)	ACE							
	adjusted by treatment history*				subgroup of notreatment			
	stage**	beta***	SE	p value	stage**	beta***	SE	p value
rs4459609 (61548948)	1st	3.11	0.210	5.22e-57	comb	2.86	0.148	7.45e-76
rs4309 (61559923)	2nd	2.52	0.179	2.08e-42	comb	3.30	0.135	8.57e-114
rs4311 (61560763)	1st	3.33	0.195	1.91e-70	comb	2.94	0.149	1.50e-78
rs4311 (61560763)	2nd	2.62	0.181	2.25e-80	comb	3.02	0.146	1.00e-85
rs4329 (61563458)	1st	3.29	0.205	5.39e-64	comb	2.94	0.149	1.50e-78
rs4329 (61563458)	2nd	2.70	0.177	1.78e-44	comb	3.02	0.146	1.00e-85
rs4343 (61566031)	1st	3.30	0.204	7.35e-65	comb	3.02	0.146	1.27e-85
rs4343 (61566031)	2nd	2.71	0.177	7.42e-49	comb	3.02	0.146	1.27e-85
rs4353 (61570422)	1st	3.28	0.204	3.13e-64	comb	3.02	0.146	1.27e-85
rs4353 (61570422)	2nd	2.71	0.177	2.88e-49	comb	3.78	0.128	1.49e-155
rs4362 (61573761)	1st	3.99	0.180	2.32e-105	comb	3.77	0.128	6.96e-156
rs4362 (61573761)	2nd	3.60	0.159	1.54e-97	comb	3.77	0.128	6.96e-156
rs4362 (61573761)	1st	4.01	0.179	1.48e-106	comb	3.77	0.128	6.96e-156
rs4362 (61573761)	2nd	3.62	0.158	1.51e-99	comb	3.77	0.128	6.96e-156
rs4461142 (61578048)	1st	3.62	0.158	4.40e-26	comb	3.77	0.128	6.96e-156
rs4461142 (61578048)	2nd	2.03	0.209	4.27e-29	comb	1.89	0.145	7.80e-37

* Treatment history is coded by 0 for untreated, 1 for treated or NA for not available.

** 1st population from the Takahata cohort and 2nd from the Yamagata cohort, whereas comb is combined populations of certainly untreated individuals in both cohorts.

*** Regression coefficient

**** Coefficient t-statistic

Table S7. SNPs associated with P-LIP and/or ACE levels at a genome-wide significance level

locus (chr)	SNPs (build135)	position (GRCh37/hg19)	starnd	major allele	minor allele	class		residue change	note
						genome	functional		
ABO (9q32)	rs4363269	136123840	+	A	G	genomic	3' downstream ~6.7kb		
	rs8176749	136131188	-	G	A	coding	synonymous	Leu310Leu	
	rs8176746	136131322	-	C	A	coding	nonsynonymous	Leu266Met	Allele A is associated with B group
	rs2073824	136132633	+	A	G	intron			
	rs657152	136139265	+	C	A	intron			
	rs500498	136148647	-	G	A	intron			
	rs505922	136149229	-	A	G	intron			Allele A is associated with O group
	rs495828	136154867	-	C	A	genomic	5' upstream ~4.2kb		
	rs7025162	136166346	-	G	A	genomic	5' upstream ~15.7kb		
ACE1 (17q23.3)	rs4459609	61548948	+	A	C	genomic	5' upstream ~5.5kb		
	rs4309	61559923	-	A	G	coding	synonymous	Pro405Pro	
	rs4311	61560763	-	G	A	intron			
	rs4329	61563458	+	G	A	intron			
	rs4343	61566031	+	A	G	intron			
	rs4353	61570422	+	G	A	intron			
	rs4362	61573761	-	G	A	coding	synonymous	Phe555Phe	
	rs4461142	61578048	-	G	A	genomic	3' downstream ~2.3kb		

Table S8. Associations of ABO gene in 9q32 with P-LIP and ACE in combined population

SNPs (position)	minor allele	MAF	P-LIP			ACE		
			beta*	T**	p value	beta*	T**	p value
rs4363269	G	0.276	-0.115	-0.321	> 1e-7	-1.94	-14.1	1.87e-43
rs8176749	A	0.177	4.78	11.6	3.02e-30	2.29	14.0	4.08e-43
rs8176746	A	0.186	4.94	12.2	2.01e-33	-1.07	-8.40	7.06e-17
rs657152	A	0.448	3.05	9.61	1.50e-21	0.417	3.26	> 1e-7
rs500498	A	0.449	-2.76	-8.71	5.16e-18	-0.581	-4.57	> 1e-7
rs505922	G	0.455	3.11	9.88	1.18e-22	-2.31	-17.0	> 1e-7
rs495828	A	0.271	0.101	0.282	> 1e-7	-2.27	-10.8	8.24e-62
rs7025162	A	0.404	-0.715	-2.21	> 1e-7	-1.31	-10.3	1.65e-24

* Regression coefficient

** Coefficient t-statistic

Table S9. Associations of ACE gene in 17q23.2 with ACE in combined population

SNPs (position)	minor allele	MAF	ACE		
			beta*	SE	p value
rs4459609	G	0.276	2.765	0.125	9.462e-101
rs4309	A	0.177	3.27	0.116	5.659e-154
rs4311	A	0.186	2.891	0.126	4.389e-108
rs4329	A	0.448	2.945	0.123	3.04e-115
rs4343	A	0.449	2.939	0.123	6.647e-115
rs4353	G	0.455	3.702	0.112	4.446e-205
rs4362	A	0.271	3.726	0.111	1.527e-208
rs4461142	A	0.404	1.923	0.123	8.542e-053

* Regression coefficient

** Coefficient t-statistic

Table S10. Sample numbers removed by each quality control filter

Initial sample size	1639
Individual missing data rate (imiss) > 0.05	16
Individual heterozygosity > 0.35	6
Cryptic relatedness: likely 1st - 3rd relatives*	314
Population outliers (PCA by EIGENSTRAT)	51
Final sample size	1252

*eliminating sample with higher imiss from each pair.

Text S1. Organization and all contributors list of Yamagata University Genomic Cohort Consortium

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