

**Significant *trans* - associations**  
**linear regression, individual population analysis, 0.001 permutation threshold.**

Gene	Illumina GeneID	SNP Label	Gene chromosome	SNP chromosome	SNP coordinate (NCBI B35)	$-\log_{10}(p\text{-value})$	Adjusted R <sup>2</sup>	Association gradient	permutation threshold (0.001)	Population
SNX27	GI_31742500-S	rs10753434	1	1	223867335	6.9944	0.3786	-0.2697	6.8283	CEU
C19orf24	GI_42476017-S	rs10753434	19	1	223867335	7.1019	0.3837	-0.2616	6.8390	CEU
ARIH1	GI_9966762-S	rs10753434	15	1	223867335	7.0443	0.3810	-0.2005	6.9786	CEU
Hs.132700	Hs.132700-S	rs10865974	15	3	52693320	7.5672	0.4056	-0.0563	6.9235	CEU
SNX27	GI_31742500-S	rs10916133	1	1	223872402	6.9944	0.3786	0.2697	6.8283	CEU
C19orf24	GI_42476017-S	rs10916133	19	1	223872402	7.1019	0.3837	0.2616	6.8390	CEU
ARIH1	GI_9966762-S	rs10916133	15	1	223872402	7.0443	0.3810	0.2005	6.9786	CEU
Hs.132700	Hs.132700-S	rs1108842	15	3	52695120	6.9335	0.3756	-0.0543	6.9235	CEU
LOC401154	GI_42656959-S	rs1113765	4	7	55663543	11.5989	0.5662	-0.3376	7.2717	CEU
LOC401154	GI_42656959-S	rs11238381	4	7	55666110	10.6640	0.5332	0.3305	7.2717	CEU
ZFYVE27	GI_34303918-S	rs11242837	10	6	3206887	7.1921	0.4413	-0.0947	6.8628	CEU
Hs.132700	Hs.132700-S	rs11714565	15	3	52581332	8.3128	0.4628	-0.0575	6.9235	CEU
Hs.132700	Hs.132700-S	rs11720243	15	3	52593058	7.4592	0.4060	0.0561	6.9235	CEU
LOC401118	GI_42657060-S	rs11757159	4	6	32628250	7.1443	0.3858	-0.8205	6.9800	CEU
LOC401154	GI_42656959-S	rs11982736	4	7	55629389	14.7860	0.6623	-0.3743	7.2717	CEU
Hs.475035	Hs.475035-S	rs12085083	1	1	170139974	7.0745	0.3877	0.1047	6.8608	CEU
LOC401118	GI_42657060-S	rs12191360	4	6	32559339	9.8565	0.5027	0.9883	6.9800	CEU
Hs.132700	Hs.132700-S	rs12635140	15	3	52713205	7.5672	0.4056	0.0563	6.9235	CEU
Hs.132700	Hs.132700-S	rs13083798	15	3	52624788	6.9335	0.3756	-0.0543	6.9235	CEU
Hs.132700	Hs.132700-S	rs13085775	15	3	52653410	7.5672	0.4056	-0.0563	6.9235	CEU
Hs.132700	Hs.132700-S	rs17052259	15	3	52568178	7.5571	0.4051	-0.0559	6.9235	CEU
LOC400452	GI_42660576-S	rs17816433	15	21	39524326	13.5733	0.6349	1.7343	7.6952	CEU
Hs.132700	Hs.132700-S	rs1866268	15	3	52694438	7.5672	0.4056	-0.0563	6.9235	CEU
GGPS1	GI_21359876-S	rs194590	1	7	111333793	7.6476	0.4261	-0.1893	6.9375	CEU
CTBP2	GI_4557498-I	rs2040748	10	6	31351764	7.8938	0.4317	-0.4819	7.7086	CEU
LOC400452	GI_42660576-S	rs2056844	15	21	39592330	13.8045	0.6352	1.7345	7.6952	CEU
Hs.132700	Hs.132700-S	rs2072390	15	3	52755549	7.5672	0.4056	0.0563	6.9235	CEU
LOC400452	GI_42660576-S	rs2150411	15	21	39496422	13.5733	0.6349	-1.7343	7.6952	CEU
LOC400452	GI_42660576-S	rs2150413	15	21	39536748	14.2607	0.6545	-1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2150414	15	21	39536858	14.2607	0.6545	-1.7400	7.6952	CEU
LOC401118	GI_42657060-S	rs2157051	4	6	32766602	13.0812	0.6472	-1.1313	6.9800	CEU
Hs.132700	Hs.132700-S	rs2289247	15	3	52702297	7.5672	0.4056	-0.0563	6.9235	CEU
LOC400452	GI_42660576-S	rs2297256	15	21	39480291	13.5267	0.6401	1.7380	7.6952	CEU
Hs.132700	Hs.132700-S	rs2336146	15	3	52601686	7.2168	0.3945	0.0552	6.9235	CEU
Hs.132700	Hs.132700-S	rs2336149	15	3	52667164	6.9909	0.3836	0.0557	6.9235	CEU
LOC401118	GI_42657060-S	rs2395185	4	6	32541145	7.4625	0.4007	-0.8735	6.9800	CEU
LOC401118	GI_42657060-S	rs2516049	4	6	32678378	7.6861	0.4110	-0.8642	6.9800	CEU

SPTLC1	GI_30474870-A	rs25803	9	5	129056512	7.1416	0.3856	-0.2832	7.1331	CEU
SPTLC1	GI_30474870-A	rs25816	9	5	129020335	7.1416	0.3856	0.2832	7.1331	CEU
SPTLC1	GI_30474870-A	rs25821	9	5	129019081	7.1416	0.3856	-0.2832	7.1331	CEU
LOC400452	GI_42660576-S	rs2776311	15	21	39638755	23.1317	0.8306	1.9649	7.6952	CEU
LOC400452	GI_42660576-S	rs2836922	15	21	39438178	13.3075	0.6271	-1.6726	7.6952	CEU
LOC400452	GI_42660576-S	rs2836926	15	21	39458660	14.2607	0.6545	1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2836930	15	21	39467526	14.2607	0.6545	-1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2836931	15	21	39468439	14.7985	0.6822	1.7344	7.6952	CEU
LOC400452	GI_42660576-S	rs2836934	15	21	39486755	14.2607	0.6545	-1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2836942	15	21	39513251	13.5733	0.6349	-1.7343	7.6952	CEU
LOC400452	GI_42660576-S	rs2836943	15	21	39514481	14.2607	0.6545	-1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2836948	15	21	39521547	13.5733	0.6349	1.7343	7.6952	CEU
LOC400452	GI_42660576-S	rs2836950	15	21	39526299	10.1830	0.5214	-1.7602	7.6952	CEU
LOC400452	GI_42660576-S	rs2836952	15	21	39538521	14.2607	0.6545	1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2836956	15	21	39544888	13.1899	0.6171	1.6789	7.6952	CEU
LOC400452	GI_42660576-S	rs2836958	15	21	39545396	13.2279	0.6587	1.6948	7.6952	CEU
LOC400452	GI_42660576-S	rs2836959	15	21	39545538	13.5733	0.6349	1.7343	7.6952	CEU
LOC400452	GI_42660576-S	rs2836961	15	21	39548890	12.3294	0.5968	-1.7760	7.6952	CEU
LOC400452	GI_42660576-S	rs2836966	15	21	39559391	14.2607	0.6545	-1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs2836980	15	21	39609488	14.5527	0.6625	-1.7379	7.6952	CEU
LOC400452	GI_42660576-S	rs2836983	15	21	39611206	14.5527	0.6625	1.7379	7.6952	CEU
LOC400452	GI_42660576-S	rs2836985	15	21	39623039	14.5527	0.6625	-1.7379	7.6952	CEU
LOC401118	GI_42657060-S	rs2858330	4	6	32766693	7.1473	0.3859	-0.8366	6.9800	CEU
Hs.132700	Hs.132700-S	rs2878628	15	3	52559755	7.5571	0.4051	-0.0559	6.9235	CEU
SNX27	GI_31742500-S	rs3014272	1	1	223890175	6.9944	0.3786	-0.2697	6.8283	CEU
C19orf24	GI_42476017-S	rs3014272	19	1	223890175	7.1019	0.3837	-0.2616	6.8390	CEU
ARIH1	GI_9966762-S	rs3014272	15	1	223890175	7.0443	0.3810	-0.2005	6.9786	CEU
SPTLC1	GI_30474870-A	rs30649	9	5	129041704	7.1416	0.3856	0.2832	7.1331	CEU
SPTLC1	GI_30474870-A	rs30656	9	5	129038668	7.1416	0.3856	-0.2832	7.1331	CEU
SPTLC1	GI_30474870-A	rs30658	9	5	129037957	7.1416	0.3856	-0.2832	7.1331	CEU
SPTLC1	GI_30474870-A	rs30662	9	5	129036031	7.1416	0.3856	-0.2832	7.1331	CEU
LOC388461	GI_41150880-S	rs3129719	18	6	32769757	8.1396	0.4314	0.1533	7.3269	CEU
LOC401118	GI_42657060-S	rs3129719	4	6	32769757	15.8032	0.6883	1.1745	6.9800	CEU
LOC388461	GI_41150880-S	rs3129720	18	6	32771609	8.1536	0.4435	0.1549	7.3269	CEU
LOC401118	GI_42657060-S	rs3129720	4	6	32771609	15.2143	0.6864	1.1637	6.9800	CEU
LOC388461	GI_41150880-S	rs3134975	18	6	32760559	8.1396	0.4314	-0.1533	7.3269	CEU
LOC401118	GI_42657060-S	rs3134975	4	6	32760559	15.8032	0.6883	-1.1745	6.9800	CEU
LOC388461	GI_41150880-S	rs3135006	18	6	32775097	7.9449	0.4284	-0.1525	7.3269	CEU
LOC401118	GI_42657060-S	rs3135006	4	6	32775097	15.5425	0.6882	-1.1668	6.9800	CEU
Hs.132700	Hs.132700-S	rs3733041	15	3	52706638	7.3045	0.4098	0.0569	6.9235	CEU
Hs.132700	Hs.132700-S	rs3733045	15	3	52618347	7.5672	0.4056	0.0563	6.9235	CEU
Hs.132700	Hs.132700-S	rs3755798	15	3	52716200	7.4533	0.4057	0.0568	6.9235	CEU
LOC388565	GI_41208836-S	rs3756890	19	6	138765462	7.6271	0.4083	-0.1968	7.5162	CEU
Hs.132700	Hs.132700-S	rs3774366	15	3	52616295	7.5672	0.4056	-0.0563	6.9235	CEU
Hs.132700	Hs.132700-S	rs3796353	15	3	52568270	7.3652	0.4071	0.0566	6.9235	CEU
DPCK	GI_22095394-S	rs4233227	17	1	223888315	7.0510	0.3865	-0.2143	6.8782	CEU
MRPL10	GI_22547113-A	rs4233227	17	1	223888315	7.2872	0.3979	-0.2326	7.1349	CEU
SNX27	GI_31742500-S	rs4233227	1	1	223888315	7.1608	0.3918	-0.2731	6.8283	CEU
C19orf24	GI_42476017-S	rs4233227	19	1	223888315	7.0381	0.3859	-0.2628	6.8390	CEU
ADPRHL2	GI_8923416-S	rs4233227	1	1	223888315	7.0703	0.3875	-0.2295	7.0622	CEU
SNX27	GI_31742500-S	rs4294380	1	1	223885720	6.8485	0.3767	0.2677	6.8283	CEU
C19orf24	GI_42476017-S	rs4294380	19	1	223885720	6.9564	0.3819	0.2596	6.8390	CEU

ARIH1	GI_9966762-S	rs4294380	15	1	223885720	7.0453	0.3863	0.1971	6.9786	CEU
Hs.132700	Hs.132700-S	rs4687629	15	3	52590267	7.3389	0.4058	0.0545	6.9235	CEU
LOC400452	GI_42660576-S	rs4818001	15	21	39496958	13.5733	0.6349	1.7343	7.6952	CEU
Hs.132700	Hs.132700-S	rs6445534	15	3	52718424	7.1526	0.3968	0.0543	6.9235	CEU
C20orf155	GI_34147220-S	rs6484718	20	11	34389333	7.5576	0.4813	-0.3628	6.8928	CEU
LOC400452	GI_42660576-S	rs6517522	15	21	39475715	18.8434	0.7608	-1.9012	7.6952	CEU
LOC388565	GI_41208836-S	rs6570228	19	6	138732788	7.6271	0.4083	-0.1968	7.5162	CEU
LOC401154	GI_42656959-S	rs6593279	4	7	55578179	14.4794	0.6942	-0.3885	7.2717	CEU
Hs.132700	Hs.132700-S	rs6805156	15	3	52584402	7.5571	0.4051	0.0559	6.9235	CEU
LOC401118	GI_42657060-S	rs6903608	4	6	32536263	9.8565	0.5027	-0.9883	6.9800	CEU
LOC388461	GI_41150880-S	rs6916742	18	6	32561169	8.3504	0.4407	0.1625	7.3269	CEU
LOC401118	GI_42657060-S	rs6916742	4	6	32561169	9.7741	0.4995	1.0546	6.9800	CEU
LOC388461	GI_41150880-S	rs6928482	18	6	32734227	8.5641	0.4499	0.1501	7.3269	CEU
LOC401118	GI_42657060-S	rs6928482	4	6	32734227	7.8822	0.4199	0.8875	6.9800	CEU
EPB41L4A	GI_37537711-S	rs701831	5	6	32657379	11.9383	0.5776	-0.3960	7.0086	CEU
LOC401118	GI_42657060-S	rs701831	4	6	32657379	11.2426	0.5539	-1.1701	6.9800	CEU
LOC400452	GI_42660576-S	rs7278297	15	21	39547622	13.5733	0.6349	-1.7343	7.6952	CEU
LOC400452	GI_42660576-S	rs7279497	15	21	39453176	14.5043	0.6547	-1.7401	7.6952	CEU
LOC400452	GI_42660576-S	rs7282582	15	21	39618844	10.6900	0.5341	1.6079	7.6952	CEU
RABIF	GI_31543539-S	rs7287503	1	22	40896855	8.6645	0.4600	-0.2450	7.3833	CEU
LOC400452	GI_42660576-S	rs741795	15	21	39640310	22.4323	0.8154	1.9234	7.6952	CEU
SNX27	GI_31742500-S	rs7555764	1	1	223866754	6.9944	0.3786	-0.2697	6.8283	CEU
C19orf24	GI_42476017-S	rs7555764	19	1	223866754	7.1019	0.3837	-0.2616	6.8390	CEU
ARIH1	GI_9966762-S	rs7555764	15	1	223866754	7.0443	0.3810	-0.2005	6.9786	CEU
Hs.132700	Hs.132700-S	rs7652191	15	3	52585455	8.0746	0.4582	0.0590	6.9235	CEU
KIAA1892	GI_22218618-S	rs769218	9	11	34427255	7.8978	0.4262	0.3666	7.5323	CEU
LOC388461	GI_41150880-S	rs7739203	18	6	32549533	8.2096	0.4345	-0.1321	7.3269	CEU
LOC401118	GI_42657060-S	rs7739203	4	6	32549533	8.1896	0.4337	-0.8066	6.9800	CEU
LOC401118	GI_42657060-S	rs7739357	4	6	32549619	7.8566	0.4744	-0.8685	6.9800	CEU
LOC401118	GI_42657060-S	rs7749092	4	6	32557028	8.5657	0.4558	1.3332	6.9800	CEU
LOC401118	GI_42657060-S	rs7749242	4	6	32557008	9.8565	0.5027	-0.9883	6.9800	CEU
LOC388461	GI_41150880-S	rs7755212	18	6	32549386	7.5892	0.4546	-0.1397	7.3269	CEU
LOC401118	GI_42657060-S	rs7755212	4	6	32549386	8.7016	0.5067	-0.9050	6.9800	CEU
LOC400452	GI_42660576-S	rs8130320	15	21	39502128	14.2607	0.6545	1.7400	7.6952	CEU
LOC400452	GI_42660576-S	rs8132143	15	21	39506128	13.5733	0.6349	-1.7343	7.6952	CEU
LOC400452	GI_42660576-S	rs8132424	15	21	39506282	13.8045	0.6352	-1.7345	7.6952	CEU
LOC400452	GI_42660576-S	rs8134214	15	21	39464237	14.2607	0.6545	-1.7400	7.6952	CEU
CUEDC2	GI_13128995-S	rs845552	10	7	55019716	7.6188	0.4080	0.1579	6.9800	CEU
LDHB	GI_22726178-S	rs845552	12	7	55019716	7.9394	0.4225	0.1470	6.9777	CEU
SCFD1	GI_33469977-A	rs910091	14	11	34393603	6.9576	0.3928	-0.1763	6.8372	CEU
EPB41L4A	GI_37537711-S	rs9268199	5	6	32386613	8.8714	0.4630	0.3558	7.0086	CEU
LOC401118	GI_42657060-S	rs9268199	4	6	32386613	11.6176	0.5668	1.1833	6.9800	CEU
LOC401118	GI_42657060-S	rs9268228	4	6	32395097	7.6619	0.4099	0.9639	6.9800	CEU
LOC401118	GI_42657060-S	rs9268853	4	6	32537621	7.4625	0.4007	0.8735	6.9800	CEU
LOC401118	GI_42657060-S	rs9268858	4	6	32537736	7.4625	0.4007	0.8735	6.9800	CEU
LOC388461	GI_41150880-S	rs9268877	18	6	32539125	8.3781	0.4419	-0.1346	7.3269	CEU
LOC401118	GI_42657060-S	rs9268877	4	6	32539125	8.3619	0.4412	-0.8218	6.9800	CEU
LOC401118	GI_42657060-S	rs9268882	4	6	32539601	7.9439	0.4585	0.8381	6.9800	CEU
LOC401118	GI_42657060-S	rs9268914	4	6	32540534	10.1024	0.5513	-1.0155	6.9800	CEU
LOC388461	GI_41150880-S	rs9268923	18	6	32540813	7.7501	0.4430	-0.1301	7.3269	CEU
LOC401118	GI_42657060-S	rs9268923	4	6	32540813	10.4751	0.5587	-0.9001	6.9800	CEU
LOC401118	GI_42657060-S	rs9268969	4	6	32542327	9.2416	0.5526	-0.8463	6.9800	CEU

LOC401118	GI_42657060-S	rs9268979	4	6	32543022	7.4549	0.4170	0.7654	6.9800	CEU
LOC401118	GI_42657060-S	rs9269065	4	6	32548150	8.4603	0.4823	0.8792	6.9800	CEU
LOC388461	GI_41150880-S	rs9269069	18	6	32548340	8.1222	0.4363	-0.1346	7.3269	CEU
LOC401118	GI_42657060-S	rs9269069	4	6	32548340	8.4945	0.4527	-0.8415	6.9800	CEU
LOC401118	GI_42657060-S	rs9269070	4	6	32548429	7.9420	0.4717	0.9132	6.9800	CEU
LOC388461	GI_41150880-S	rs9269071	18	6	32548445	8.3781	0.4419	0.1346	7.3269	CEU
LOC401118	GI_42657060-S	rs9269071	4	6	32548445	8.3619	0.4412	0.8218	6.9800	CEU
LOC401118	GI_42657060-S	rs9269078	4	6	32548857	9.4545	0.5053	-0.9793	6.9800	CEU
LOC388461	GI_41150880-S	rs9269080	18	6	32548947	8.3781	0.4419	0.1346	7.3269	CEU
LOC401118	GI_42657060-S	rs9269080	4	6	32548947	8.3619	0.4412	0.8218	6.9800	CEU
LOC401118	GI_42657060-S	rs9269081	4	6	32549078	9.1948	0.4763	-0.9577	6.9800	CEU
LOC401118	GI_42657060-S	rs9269190	4	6	32556478	9.8565	0.5027	0.9883	6.9800	CEU
EPB41L4A	GI_37537711-S	rs9270986	5	6	32682038	11.7013	0.5760	-0.3968	7.0086	CEU
LOC401118	GI_42657060-S	rs9270986	4	6	32682038	10.9348	0.5493	-1.1577	6.9800	CEU
LOC388461	GI_41150880-S	rs9272346	18	6	32712350	15.3893	0.6780	0.1753	7.3269	CEU
LOC401118	GI_42657060-S	rs9272346	4	6	32712350	13.8668	0.6370	1.0389	6.9800	CEU
LOC388461	GI_41150880-S	rs9272723	18	6	32717405	14.3162	0.6496	0.1697	7.3269	CEU
LOC401118	GI_42657060-S	rs9272723	4	6	32717405	12.6378	0.6001	0.9976	6.9800	CEU
LOC401118	GI_42657060-S	rs9275141	4	6	32759095	7.1473	0.3859	0.8366	6.9800	CEU
LOC388461	GI_41150880-S	rs9275562	18	6	32785824	7.9932	0.4363	0.1529	7.3269	CEU
LOC401118	GI_42657060-S	rs9275562	4	6	32785824	13.7229	0.6458	1.1288	6.9800	CEU
LOC401118	GI_42657060-S	rs9286790	4	6	32547806	7.1120	0.4180	0.9063	6.9800	CEU
LOC401118	GI_42657060-S	rs9380319	4	6	32785890	7.1715	0.3923	-0.8796	6.9800	CEU
LOC401118	GI_42657060-S	rs9405040	4	6	32547371	7.4625	0.4007	-0.8735	6.9800	CEU
LOC401118	GI_42657060-S	rs9405112	4	6	32553578	7.4625	0.4007	0.8735	6.9800	CEU
LOC388565	GI_41208836-S	rs9495068	19	6	138753988	8.0298	0.4322	0.1932	7.5162	CEU
LOC388565	GI_41208836-S	rs9495070	19	6	138756312	7.6271	0.4083	0.1968	7.5162	CEU
LOC401154	GI_42656959-S	rs9642404	4	7	55631008	7.9797	0.4243	-0.3522	7.2717	CEU
Hs.132700	Hs.132700-S	rs998909	15	3	52780133	7.5672	0.4056	0.0563	6.9235	CEU
LOC401155	GI_42656964-S	rs17802433	9	2	94959504	27.2677	0.8741	0.9000	7.5869	CEU
hmm3588	hmm3588-S	rs1529874	14	2	233175093	12.2414	0.5875	-0.3096	11.7630	CEU
KCNMB3	GI_25952101-I	rs3739038	3	2	238454703	21.7376	0.8217	-0.6067	7.7728	CEU
KIAA1002	GI_33356165-S	rs2270286	12	16	8630130	6.8756	0.3728	0.3186	6.8374	CEU
PRIM1	GI_41349493-S	rs2270286	12	16	8630130	7.1618	0.3866	0.5057	7.1398	CEU
LDHC	GI_9257227-A	rs17075173	11	13	50488434	7.3687	0.3963	-0.7269	7.1269	CEU
GYG	GI_20127456-S	rs7990383	3	13	109909236	7.0764	0.3825	0.2964	7.0155	CEU
PEPD	GI_4557834-S	rs7990383	19	13	109909236	7.3826	0.3970	0.2555	6.9451	CEU
ZNF322B	GI_39841070-S	rs3736781	9	6	26613341	7.6562	0.4097	-0.2954	7.2611	CEU
EPB41L4A	GI_37537711-S	rs3129941	5	6	32445664	11.3258	0.5568	-0.3644	7.0086	CEU
RPL14	GI_16753224-S	rs1047855	4	3	40443900	20.7198	0.7886	0.8941	7.1014	CEU
BMP4	GI_19528651-A	rs16845107	14	3	114610681	10.0488	0.5101	-1.2816	9.1525	CEU
RRAS2	GI_38569462-S	rs17579502	1	9	5910886	7.8651	0.4192	0.7333	7.8247	CEU
MGC24133	GI_31341350-S	rs7302230	1	12	7179699	7.5568	0.4051	1.5356	7.3859	CEU
CACNB2	GI_45359833-I	rs7167599	10	15	97490122	9.6014	0.4988	-0.2415	9.4786	CEU
FLJ23506	GI_13376239-S	rs17610219	19	4	148182469	7.7779	0.4152	-0.2218	7.3020	CEU
PEPD	GI_4557834-S	rs3803236	19	13	109915746	7.1692	0.4611	0.2832	6.9451	CEU
FLT1	GI_32306519-S	rs2271194	13	12	54763961	17.1168	0.7190	0.4180	7.1378	CEU
LOC389472	GI_42657859-S	rs2271194	7	12	54763961	16.6517	0.7085	0.6920	7.0910	CEU
LOC401470	GI_42658911-S	rs2271194	8	12	54763961	14.5914	0.6571	0.6390	7.3419	CEU
hmm32535	hmm32535-S	rs2271194	7	12	54763961	16.1568	0.6969	0.2107	8.0518	CEU
UGT2B11	GI_4507822-S	rs10034124	4	4	69222280	24.7728	0.9389	-2.8313	9.2561	CHB
LOC389972	GI_41149334-S	rs10037055	10	5	176623885	11.8628	0.6857	-0.2549	7.2677	CHB

FACL2	GI_12669906-S	rs10040724	4	5	873594	7.8200	0.5182	0.3571	7.3032	CHB
FLJ14803	GI_14249557-S	rs10040724	7	5	873594	7.8386	0.5191	0.2734	6.9672	CHB
NOP5/NOP58	GI_34222329-S	rs10040724	2	5	873594	8.4284	0.5481	0.2544	6.9808	CHB
LOC389972	GI_41149334-S	rs10077929	10	5	176630450	11.8628	0.6857	0.2549	7.2677	CHB
LOC401154	GI_42656959-S	rs10232851	4	7	55603043	15.0702	0.8004	-0.3902	7.0518	CHB
UGT2B11	GI_4507822-S	rs10471154	4	4	69219098	9.5995	0.6006	-2.0833	9.2561	CHB
LOC401154	GI_42656959-S	rs10481004	4	7	55542880	9.7304	0.6061	0.3436	7.0518	CHB
LOC389972	GI_41149334-S	rs11949767	10	5	176672850	8.8773	0.5689	0.2271	7.2677	CHB
LOC389972	GI_41149334-S	rs12187932	10	5	176575757	11.8628	0.6857	0.2549	7.2677	CHB
LOC377709	GI_37552433-S	rs12468079	8	2	217144583	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs12472569	8	2	217206912	8.5418	0.5534	0.1639	8.1994	CHB
LOC389972	GI_41149334-S	rs12519145	10	5	176488129	8.6779	0.5598	-0.2334	7.2677	CHB
LOC377709	GI_37552433-S	rs12611945	8	2	217119515	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs12612280	8	2	217123766	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs12612749	8	2	217129141	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs12619115	8	2	217189346	8.7148	0.5974	-0.1762	8.1994	CHB
LOC377709	GI_37552433-S	rs12620897	8	2	217161264	9.0808	0.5957	-0.1714	8.1994	CHB
LOC377709	GI_37552433-S	rs12621359	8	2	217119330	9.6979	0.6047	-0.1720	8.1994	CHB
FACL2	GI_12669906-S	rs12655323	4	5	875109	7.7899	0.5250	0.3598	7.3032	CHB
FLJ14803	GI_14249557-S	rs12655323	7	5	875109	7.8012	0.5256	0.2754	6.9672	CHB
NOP5/NOP58	GI_34222329-S	rs12655323	2	5	875109	8.3612	0.5534	0.2560	6.9808	CHB
ARID3A	GI_4885192-S	rs12655323	19	5	875109	7.0077	0.4834	0.2827	6.8830	CHB
FACL2	GI_12669906-S	rs12656150	4	5	875123	7.8200	0.5182	-0.3571	7.3032	CHB
FLJ14803	GI_14249557-S	rs12656150	7	5	875123	7.8386	0.5191	-0.2734	6.9672	CHB
NOP5/NOP58	GI_34222329-S	rs12656150	2	5	875123	8.4284	0.5481	-0.2544	6.9808	CHB
LOC389972	GI_41149334-S	rs12660023	10	5	176633491	11.8628	0.6857	-0.2549	7.2677	CHB
LOC389972	GI_41149334-S	rs12716330	10	5	176531108	11.8628	0.6857	0.2549	7.2677	CHB
LOC388461	GI_41150880-S	rs12918	18	6	32735364	13.5603	0.7375	-0.2729	7.4936	CHB
LOC401118	GI_42657060-S	rs12918	4	6	32735364	9.5885	0.6001	-1.3850	7.4613	CHB
LOC389972	GI_41149334-S	rs1363405	10	5	176569488	11.8628	0.6857	-0.2549	7.2677	CHB
LOC389972	GI_41149334-S	rs1546363	10	5	176533637	11.7306	0.6899	0.2536	7.2677	CHB
LOC377709	GI_37552433-S	rs16856088	8	2	217118220	9.6077	0.6009	0.1705	8.1994	CHB
LOC377709	GI_37552433-S	rs16856102	8	2	217120743	9.1355	0.6569	-0.1884	8.1994	CHB
LOC377709	GI_37552433-S	rs16856126	8	2	217124692	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs16856136	8	2	217128131	9.5549	0.6074	-0.1743	8.1994	CHB
LOC377709	GI_37552433-S	rs1705105	8	2	217131441	9.6979	0.6047	0.1720	8.1994	CHB
EPB41L4A	GI_37537711-S	rs17191234	5	6	32672659	7.6184	0.5079	0.3054	7.1949	CHB
LOC377709	GI_37552433-S	rs2066518	8	2	217113894	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs2066521	8	2	217122924	9.6979	0.6047	-0.1720	8.1994	CHB
ZFP91	GI_25777699-I	rs2074411	11	17	32953876	7.3879	0.4958	0.4225	7.1887	CHB
PREI3	GI_41349448-A	rs2074411	2	17	32953876	7.1606	0.4837	0.3648	7.1576	CHB
LOC389972	GI_41149334-S	rs2336237	10	5	176598485	11.8628	0.6857	0.2549	7.2677	CHB
LOC389972	GI_41149334-S	rs244708	10	5	176522191	11.8628	0.6857	0.2549	7.2677	CHB
LOC377709	GI_37552433-S	rs2627045	8	2	217133513	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs2738286	8	2	217140870	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs2738289	8	2	217169528	9.6979	0.6047	0.1720	8.1994	CHB
LOC400452	GI_42660576-S	rs2776311	15	21	39638755	21.0874	0.8823	2.5836	7.6924	CHB
LOC377709	GI_37552433-S	rs284548	8	2	217136361	10.0309	0.6452	-0.1773	8.1994	CHB
LOC377709	GI_37552433-S	rs284552	8	2	217138550	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs284565	8	2	217184312	9.6077	0.6009	0.1705	8.1994	CHB
LOC377709	GI_37552433-S	rs284567	8	2	217177980	9.6077	0.6009	-0.1705	8.1994	CHB
LOC389972	GI_41149334-S	rs351861	10	5	176474543	10.8669	0.6507	-0.2452	7.2677	CHB

LOC377709	GI_37552433-S	rs3732019	8	2	217187892	8.5418	0.5534	-0.1639	8.1994	CHB
FACL2	GI_12669906-S	rs3734155	4	5	874390	7.6974	0.5203	0.3586	7.3032	CHB
FLJ14803	GI_14249557-S	rs3734155	7	5	874390	7.6594	0.5183	0.2736	6.9672	CHB
NOP5/NOP58	GI_34222329-S	rs3734155	2	5	874390	8.2117	0.5461	0.2539	6.9808	CHB
LOC377709	GI_37552433-S	rs3770480	8	2	217129746	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs3770483	8	2	217120269	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs3770485	8	2	217118416	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs3770487	8	2	217117613	9.6979	0.6047	0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs3770488	8	2	217117579	9.4292	0.6296	-0.1748	8.1994	CHB
LOC377709	GI_37552433-S	rs3806575	8	2	217188440	8.2438	0.5748	0.1710	8.1994	CHB
LOC377709	GI_37552433-S	rs3816783	8	2	217128593	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs4324305	8	2	217155418	9.6077	0.6009	0.1705	8.1994	CHB
LOC401154	GI_42656959-S	rs4563845	4	7	55552324	11.0544	0.6752	0.3354	7.0518	CHB
LOC389972	GI_41149334-S	rs4976640	10	5	176484613	11.8628	0.6857	-0.2549	7.2677	CHB
LOC377709	GI_37552433-S	rs505883	8	2	217131540	9.0327	0.5935	0.1687	8.1994	CHB
LOC377709	GI_37552433-S	rs560025	8	2	217133078	9.6979	0.6047	-0.1720	8.1994	CHB
LOC377709	GI_37552433-S	rs605702	8	2	217172200	9.3198	0.5973	-0.1670	8.1994	CHB
LOC400452	GI_42660576-S	rs6517522	15	21	39475715	10.4710	0.6357	-2.2007	7.6924	CHB
LOC377709	GI_37552433-S	rs654474	8	2	217176114	9.6979	0.6047	0.1720	8.1994	CHB
C1orf27	GI_20127566-S	rs6715047	1	2	70944984	8.2818	0.5410	0.1204	8.0067	CHB
LOC389972	GI_41149334-S	rs6872201	10	5	176653372	8.6779	0.5598	0.2334	7.2677	CHB
FACL2	GI_12669906-S	rs6875688	4	5	871479	7.6974	0.5203	-0.3586	7.3032	CHB
FLJ14803	GI_14249557-S	rs6875688	7	5	871479	7.6594	0.5183	-0.2736	6.9672	CHB
NOP5/NOP58	GI_34222329-S	rs6875688	2	5	871479	8.2117	0.5461	-0.2539	6.9808	CHB
LOC389972	GI_41149334-S	rs6886255	10	5	176650967	11.8628	0.6857	0.2549	7.2677	CHB
LOC389972	GI_41149334-S	rs6890580	10	5	176586126	11.8628	0.6857	0.2549	7.2677	CHB
LOC388461	GI_41150880-S	rs6928482	18	6	32734227	11.2738	0.6654	0.1745	7.4936	CHB
LOC401118	GI_42657060-S	rs6928482	4	6	32734227	7.7622	0.5152	0.8651	7.4613	CHB
LOC400452	GI_42660576-S	rs741795	15	21	39640310	21.0874	0.8823	2.5836	7.6924	CHB
UGT2B11	GI_4507822-S	rs7435827	4	4	69243710	11.0625	0.6578	2.6051	9.2561	CHB
LOC389972	GI_41149334-S	rs7719125	10	5	176621772	11.8628	0.6857	-0.2549	7.2677	CHB
ALDH1B1	GI_25777729-S	rs7739357	9	6	32549619	7.2970	0.5544	0.2018	7.0320	CHB
ANAPC11	GI_19923724-S	rs7764237	17	6	30105965	6.9264	0.5332	-0.2676	6.5271	CHB
CD1C	GI_4502646-S	rs9262564	1	6	31117116	7.7413	0.5401	-0.4358	7.6371	CHB
CACNA1H	GI_10864076-S	rs9270657	16	6	32673999	8.0626	0.5303	-0.1228	7.3606	CHB
EPB41L4A	GI_37537711-S	rs9270657	5	6	32673999	12.2948	0.6998	0.3911	7.1949	CHB
EPB41L4A	GI_37537711-S	rs9272218	5	6	32710216	9.4784	0.6223	-0.3508	7.1949	CHB
LOC388461	GI_41150880-S	rs9272346	18	6	32712350	14.2539	0.7562	0.1823	7.4936	CHB
LOC401118	GI_42657060-S	rs9272346	4	6	32712350	10.5679	0.6394	0.9424	7.4613	CHB
LOC388461	GI_41150880-S	rs9272723	18	6	32717405	12.4898	0.7059	0.1748	7.4936	CHB
LOC401118	GI_42657060-S	rs9272723	4	6	32717405	9.3152	0.5884	0.8974	7.4613	CHB
LOC389972	GI_41149334-S	rs9313749	10	5	176530300	11.8628	0.6857	0.2549	7.2677	CHB
LOC389972	GI_41149334-S	rs9885210	10	5	176501794	11.8628	0.6857	0.2549	7.2677	CHB
POMZP3	GI_23510405-A	rs2286428	7	7	75699023	12.2408	0.6980	0.9772	9.3542	CHB
LOC283520	GI_27482429-S	rs722896	13	2	70955116	7.3574	0.4942	0.1581	7.0816	CHB
LOC283520	GI_27482429-S	rs2075221	13	2	70955866	7.3574	0.4942	-0.1581	7.0816	CHB
LOC401155	GI_42656964-S	rs17802433	9	2	94959504	19.6544	0.8756	0.9351	7.4464	CHB
KCNMB3	GI_25952101-I	rs3739038	3	2	238454703	14.9962	0.8069	-0.6132	7.1872	CHB
APOL1	GI_21735617-A	rs2295320	22	14	23752607	7.5358	0.5036	-0.6110	6.7211	CHB
RPL30	GI_15812218-S	rs6089219	8	20	60399713	6.9881	0.4742	-0.1359	6.7902	CHB
RPL35A	GI_16117790-S	rs6089219	3	20	60399713	6.8553	0.4669	-0.1066	6.8323	CHB
RPL24	GI_17017969-S	rs6089219	3	20	60399713	7.5319	0.5034	-0.0980	7.0568	CHB

Hs.483001	Hs.483001-S	rs1538264	5	10	120393426	8.7422	0.5628	0.0529	7.4517	CHB
MICA	GI_4557750-S	rs7575	6	22	17797804	7.4144	0.4972	0.7768	7.2698	CHB
ChGn	GI_40255140-S	rs7023244	8	9	14809370	7.3086	0.4916	0.2749	7.2630	CHB
LOC201294	GI_29736784-S	rs2269700	17	9	115243117	7.9811	0.5263	0.3031	7.5925	CHB
Hs.486750	Hs.486750-S	rs1551118	6	12	79263898	7.2042	0.4860	-0.0677	6.9923	CHB
LOC389972	GI_41149334-S	rs3733875	10	5	176569846	11.1771	0.6620	-0.2544	7.2677	CHB
HIST1H3H	GI_15718725-S	rs2271194	6	12	54763961	7.5884	0.5063	0.2273	7.2973	CHB
FLT1	GI_32306519-S	rs2271194	13	12	54763961	12.3665	0.7020	0.4372	7.2192	CHB
LOC389472	GI_42657859-S	rs2271194	7	12	54763961	12.9292	0.7193	0.7286	7.0419	CHB
LOC401470	GI_42658911-S	rs2271194	8	12	54763961	11.5018	0.6734	0.7210	7.2314	CHB
hmm32535	hmm32535-S	rs2271194	7	12	54763961	9.2507	0.5856	0.1890	7.6540	CHB
UGT2B11	GI_4507822-S	rs10034124	4	4	69222280	28.1888	0.9557	-2.5800	8.8707	JPT
LOC389972	GI_41149334-S	rs10037055	10	5	176623885	11.1854	0.6623	-0.2448	7.2502	JPT
LOC389972	GI_41149334-S	rs10077929	10	5	176630450	11.1854	0.6623	0.2448	7.2502	JPT
LOC401154	GI_42656959-S	rs10232851	4	7	55603043	8.0479	0.5557	-0.2974	7.2421	JPT
GPD2	GI_4504084-S	rs10411185	2	19	8009881	7.1733	0.4843	-0.3124	6.9394	JPT
PIGS	GI_23397651-S	rs10490798	17	3	47303748	6.8304	0.4735	-0.2084	6.6648	JPT
DKFZP564I0422	GI_13899258-S	rs11102778	12	1	114566166	8.1884	0.5537	-0.4308	8.1146	JPT
FBXL4	GI_21536437-S	rs11102778	6	1	114566166	8.4431	0.5661	-0.2721	8.3858	JPT
CLK3	GI_4557478-A	rs11102778	15	1	114566166	7.6552	0.5267	-0.8226	7.4599	JPT
Hs.96557	Hs.96557-S	rs11102778	8	1	114566166	7.4513	0.5160	-0.4466	7.0729	JPT
Magmas	GI_27363460-S	rs11130126	16	3	47281898	7.4138	0.5054	-0.2407	7.2322	JPT
OSBPL8	GI_22035617-S	rs11242837	12	6	3206887	8.0135	0.6032	0.2760	7.0403	JPT
Magmas	GI_27363460-S	rs11713693	16	3	47049910	7.3420	0.5373	0.2584	7.2322	JPT
Magmas	GI_27363460-S	rs11915074	16	3	47284761	7.4138	0.5054	-0.2407	7.2322	JPT
Magmas	GI_27363460-S	rs11920354	16	3	47237250	7.4138	0.5054	0.2407	7.2322	JPT
Magmas	GI_27363460-S	rs11920853	16	3	47283187	7.3491	0.4938	0.2357	7.2322	JPT
LOC389972	GI_41149334-S	rs11949767	10	5	176672850	10.1076	0.6215	0.2209	7.2502	JPT
LOC389972	GI_41149334-S	rs12187932	10	5	176575757	8.2229	0.5382	0.2107	7.2502	JPT
LOC377709	GI_37552433-S	rs12472569	8	2	217206912	8.9009	0.5875	0.1787	7.7060	JPT
LOC389972	GI_41149334-S	rs12519145	10	5	176488129	8.1493	0.5346	-0.2051	7.2502	JPT
LOC377709	GI_37552433-S	rs12616410	8	2	217193418	8.7359	0.5799	-0.1822	7.7060	JPT
LOC389972	GI_41149334-S	rs12660023	10	5	176633491	10.8341	0.6762	-0.2447	7.2502	JPT
LOC389972	GI_41149334-S	rs12716330	10	5	176531108	8.2229	0.5382	0.2107	7.2502	JPT
LOC389972	GI_41149334-S	rs1363405	10	5	176569488	8.0553	0.5384	-0.2106	7.2502	JPT
LOC389972	GI_41149334-S	rs1546363	10	5	176533637	8.0553	0.5384	0.2106	7.2502	JPT
LOC377709	GI_37552433-S	rs1705105	8	2	217131441	7.7666	0.5238	0.2011	7.7060	JPT
Magmas	GI_27363460-S	rs17410853	16	3	47338975	7.3491	0.4938	-0.2357	7.2322	JPT
Hs.505053	Hs.505053-S	rs2010300	21	15	38867423	8.4992	0.5600	-0.0650	8.0872	JPT
LOC377709	GI_37552433-S	rs2066518	8	2	217113894	7.7666	0.5238	0.2011	7.7060	JPT
PIGS	GI_23397651-S	rs2240796	17	3	47355503	6.8304	0.4735	-0.2084	6.6648	JPT
LOC377709	GI_37552433-S	rs2272243	8	2	217190270	8.8616	0.5857	-0.1876	7.7060	JPT
LOC389972	GI_41149334-S	rs2336237	10	5	176598485	8.0553	0.5384	0.2106	7.2502	JPT
TK1	GI_4507518-S	rs238233	17	17	4826706	7.1831	0.4930	0.3422	7.1287	JPT
LOC401118	GI_42657060-S	rs2395522	4	6	32772700	10.2313	0.6821	0.9489	7.2431	JPT
LOC389972	GI_41149334-S	rs244708	10	5	176522191	8.0553	0.5384	0.2106	7.2502	JPT
LOC400452	GI_42660576-S	rs2776311	15	21	39638755	24.2805	0.9212	2.8295	8.5824	JPT
LOC377709	GI_37552433-S	rs284552	8	2	217138550	7.7666	0.5238	-0.2011	7.7060	JPT
LOC377709	GI_37552433-S	rs284565	8	2	217184312	7.7666	0.5238	0.2011	7.7060	JPT
LOC377709	GI_37552433-S	rs284567	8	2	217177980	7.7666	0.5238	-0.2011	7.7060	JPT
LOC283887	GI_42660747-S	rs2947066	16	10	32871691	7.0726	0.4870	-0.1002	6.8029	JPT
LOC377709	GI_37552433-S	rs3732019	8	2	217187892	9.2027	0.5922	-0.1914	7.7060	JPT

LOC377709	GI_37552433-S	rs3755140	8	2	217188555	8.8167	0.5748	0.1854	7.7060	JPT
LOC377709	GI_37552433-S	rs3770480	8	2	217129746	7.7666	0.5238	0.2011	7.7060	JPT
LOC401118	GI_42657060-S	rs4248168	4	6	32767721	11.1448	0.6608	0.8995	7.2431	JPT
LOC401118	GI_42657060-S	rs4713581	4	6	32768001	11.1448	0.6608	0.8995	7.2431	JPT
LOC401118	GI_42657060-S	rs4713582	4	6	32768029	11.1448	0.6608	0.8995	7.2431	JPT
LOC389972	GI_41149334-S	rs4976640	10	5	176484613	8.0553	0.5384	-0.2106	7.2502	JPT
LOC388461	GI_41150880-S	rs5000634	18	6	32771542	8.4505	0.5491	0.1572	7.1838	JPT
LOC401118	GI_42657060-S	rs5000634	4	6	32771542	9.1497	0.5811	0.8891	7.2431	JPT
LOC401118	GI_42657060-S	rs5002702	4	6	32767136	9.7756	0.6348	-0.8917	7.2431	JPT
LOC401118	GI_42657060-S	rs6457619	4	6	32771964	9.3597	0.5903	-0.8299	7.2431	JPT
LOC401118	GI_42657060-S	rs6457620	4	6	32771977	11.1448	0.6608	0.8995	7.2431	JPT
LOC388461	GI_41150880-S	rs6457622	18	6	32772141	7.2049	0.5393	-0.1456	7.1838	JPT
LOC400452	GI_42660576-S	rs6517522	15	21	39475715	24.2805	0.9212	-2.8295	8.5824	JPT
LOC389972	GI_41149334-S	rs6872201	10	5	176653372	10.8198	0.6577	0.2380	7.2502	JPT
LOC389972	GI_41149334-S	rs6886255	10	5	176650967	11.1854	0.6623	0.2448	7.2502	JPT
LOC389972	GI_41149334-S	rs6890580	10	5	176586126	8.0553	0.5384	0.2106	7.2502	JPT
LOC388461	GI_41150880-S	rs6916742	18	6	32561169	7.2441	0.4963	0.1498	7.1838	JPT
LOC388461	GI_41150880-S	rs6928482	18	6	32734227	8.3601	0.5621	0.1610	7.1838	JPT
LOC401118	GI_42657060-S	rs6928482	4	6	32734227	10.5834	0.6577	0.9765	7.2431	JPT
LOC400452	GI_42660576-S	rs741795	15	21	39640310	24.2805	0.9212	2.8295	8.5824	JPT
UGT2B11	GI_4507822-S	rs7435827	4	4	69243710	11.3464	0.6767	2.3770	8.8707	JPT
Magmas	GI_27363460-S	rs7632501	16	3	47243310	7.4138	0.5054	0.2407	7.2322	JPT
LOC389972	GI_41149334-S	rs7719125	10	5	176621772	11.1854	0.6623	-0.2448	7.2502	JPT
LOC388461	GI_41150880-S	rs7745040	18	6	32772310	8.4505	0.5491	0.1572	7.1838	JPT
LOC401118	GI_42657060-S	rs7745040	4	6	32772310	9.1497	0.5811	0.8891	7.2431	JPT
GPD2	GI_4504084-S	rs882951	2	19	7999265	7.0338	0.4848	-0.3127	6.9394	JPT
MGC955	GI_13129101-S	rs892086	1	19	10698677	7.1254	0.4817	-0.2334	7.0651	JPT
LOC401118	GI_42657060-S	rs9268831	4	6	32535726	9.6106	0.6010	-0.8448	7.2431	JPT
Hs.442025	Hs.442025-S	rs9269070	6	6	32548429	7.0640	0.5620	-0.0690	6.9110	JPT
EPB41L4A	GI_37537711-S	rs9270657	5	6	32673999	8.8144	0.5836	0.3087	7.2238	JPT
LOC388461	GI_41150880-S	rs9271568	18	6	32698441	7.2770	0.5433	0.1318	7.1838	JPT
LOC401118	GI_42657060-S	rs9271586	4	6	32698877	7.7846	0.5515	-0.7099	7.2431	JPT
LOC401118	GI_42657060-S	rs9271589	4	6	32699106	8.1484	0.5518	0.6734	7.2431	JPT
LOC401118	GI_42657060-S	rs9271591	4	6	32699119	8.5961	0.5918	0.6957	7.2431	JPT
EPB41L4A	GI_37537711-S	rs9272218	5	6	32710216	8.8759	0.5864	-0.2974	7.2238	JPT
LOC388461	GI_41150880-S	rs9272346	18	6	32712350	8.3399	0.5611	0.1618	7.1838	JPT
LOC401118	GI_42657060-S	rs9272346	4	6	32712350	10.5269	0.6556	0.9701	7.2431	JPT
LOC401118	GI_42657060-S	rs9272723	4	6	32717405	7.5753	0.5405	0.8416	7.2431	JPT
LOC401118	GI_42657060-S	rs9275224	4	6	32767856	11.1930	0.6801	-0.9368	7.2431	JPT
LOC401118	GI_42657060-S	rs9275292	4	6	32771267	11.6570	0.6962	0.9442	7.2431	JPT
LOC389972	GI_41149334-S	rs9313749	10	5	176530300	8.2229	0.5382	0.2107	7.2502	JPT
LOC389972	GI_41149334-S	rs9885210	10	5	176501794	8.0553	0.5384	0.2106	7.2502	JPT
LOC401155	GI_42656964-S	rs17802433	9	2	94959504	24.6330	0.9288	0.8192	7.5639	JPT
KCNMB3	GI_25952101-I	rs3739038	3	2	238454703	18.0430	0.8650	-0.5503	7.2900	JPT
C6orf80	GI_31083115-S	rs1771487	6	1	224708856	7.3101	0.4999	-0.3142	7.1396	JPT
GCSH	GI_31542839-S	rs10459872	1	16	79652452	7.4271	0.4979	-0.4121	7.1040	JPT
SLC19A2	GI_27734718-S	rs3790267	1	20	19513693	7.1427	0.4992	-0.3579	7.0473	JPT
CLN3	GI_4502888-S	rs3790267	16	20	19513693	6.8345	0.4820	-0.2952	6.7855	JPT
DIA1	GI_6552326-A	rs3790267	22	20	19513693	7.3972	0.5131	-0.3085	7.0000	JPT
ZNF322B	GI_39841070-S	rs3736781	9	6	26613341	9.6549	0.6116	-0.3937	7.1494	JPT
HIF1AN	GI_8923568-S	rs9367097	10	6	41293436	6.9928	0.4826	-0.1035	6.8946	JPT
RPL14	GI_16753224-S	rs1047855	4	3	40443900	11.1422	0.6694	0.7699	7.6449	JPT



LOC401495	GI_42659169-S	rs7037849	9	9	137249657	10.9302	0.6617	-0.4000	8.4647	JPT
NAP1L5	GI_24371267-S	rs1852450	4	12	16289001	7.1721	0.5009	-0.2258	6.8448	JPT
hmm30082	hmm30082-S	rs2166807	4	12	32025905	6.9689	0.4732	0.2492	6.7734	JPT
hmm30082	hmm30082-S	rs3759301	4	12	32026708	7.1099	0.4809	0.2583	6.7734	JPT
LOC389972	GI_41149334-S	rs3733875	10	5	176569846	8.4430	0.5573	-0.2198	7.2502	JPT
FLT1	GI_32306519-S	rs2271194	13	12	54763961	13.8332	0.7533	0.4153	7.3486	JPT
LOC389472	GI_42657859-S	rs2271194	7	12	54763961	15.6374	0.7973	0.7990	7.5532	JPT
LOC401470	GI_42658911-S	rs2271194	8	12	54763961	14.5168	0.7710	0.8526	7.2210	JPT
hmm32535	hmm32535-S	rs2271194	7	12	54763961	8.1926	0.5452	0.1290	7.5717	JPT
LOC387805	GI_41149627-S	rs11065706	11	12	109618251	9.7424	0.4983	0.3727	9.2056	YRI
C9orf74	GI_13569869-S	rs11769548	9	7	55428976	7.2704	0.4025	0.2161	7.1743	YRI
LOC388461	GI_41150880-S	rs12918	18	6	32735364	11.2883	0.5555	-0.2191	7.1764	YRI
LOC401118	GI_42657060-S	rs12918	4	6	32735364	8.3208	0.4394	-0.9367	7.1702	YRI
CYBA	GI_4557504-S	rs1603705	16	4	106576743	7.9385	0.4397	-0.2825	7.1655	YRI
LOC400452	GI_42660576-S	rs2037925	15	21	39621801	9.0283	0.4695	-1.4538	8.5168	YRI
LOC387805	GI_41149627-S	rs2338754	11	12	108954453	9.3943	0.4904	-0.3324	9.2056	YRI
LOC400452	GI_42660576-S	rs2776311	15	21	39638755	22.4023	0.8149	1.9538	8.5168	YRI
LOC400452	GI_42660576-S	rs2836985	15	21	39623039	10.5242	0.5280	-1.5286	8.5168	YRI
LOC400452	GI_42660576-S	rs2836988	15	21	39629845	9.0283	0.4695	-1.4538	8.5168	YRI
LOC401118	GI_42657060-S	rs3129719	4	6	32769757	19.5878	0.7688	1.0231	7.1702	YRI
LOC401118	GI_42657060-S	rs3129720	4	6	32771609	17.2878	0.7290	1.0042	7.1702	YRI
LOC401118	GI_42657060-S	rs3134975	4	6	32760559	19.5878	0.7688	-1.0231	7.1702	YRI
LOC401118	GI_42657060-S	rs3135006	4	6	32775097	19.1021	0.7657	-1.0236	7.1702	YRI
LOC388461	GI_41150880-S	rs6906021	18	6	32734289	7.5532	0.4217	0.1490	7.1764	YRI
LOC401118	GI_42657060-S	rs6906021	4	6	32734289	8.4983	0.4648	0.7426	7.1702	YRI
LOC401118	GI_42657060-S	rs6928482	4	6	32734227	8.2917	0.4381	0.6261	7.1702	YRI
LOC400452	GI_42660576-S	rs741795	15	21	39640310	14.5530	0.6625	1.9287	8.5168	YRI
LOC401118	GI_42657060-S	rs7744001	4	6	32734064	7.9873	0.4419	0.6454	7.1702	YRI
Hs.444908	Hs.444908-S	rs7919631	1	10	127457290	7.7706	0.4149	0.1160	7.5676	YRI
LOC388461	GI_41150880-S	rs9272346	18	6	32712350	15.4353	0.6792	0.1599	7.1764	YRI
LOC401118	GI_42657060-S	rs9272346	4	6	32712350	9.8780	0.5036	0.6620	7.1702	YRI
LOC388461	GI_41150880-S	rs9272723	18	6	32717405	9.4451	0.4865	0.1275	7.1764	YRI
C20orf12	GI_18491021-S	rs17802433	20	2	94959504	7.6127	0.4077	0.4761	7.2277	YRI
FJX1	GI_18765710-S	rs17802433	11	2	94959504	7.3504	0.3955	0.5940	7.1970	YRI
AK2	GI_26665888-I	rs17802433	1	2	94959504	7.4536	0.4003	0.3735	7.2961	YRI
CSRP2BP	GI_29570789-I	rs17802433	20	2	94959504	7.8902	0.4203	0.2660	7.2467	YRI
LOC401155	GI_42656964-S	rs17802433	9	2	94959504	36.7748	0.9407	0.8429	7.5260	YRI
UAP1	GI_34147515-S	rs2043449	1	2	203942196	7.5526	0.4049	-0.3052	6.9956	YRI
KCNMB3	GI_25952101-I	rs3739040	3	2	238454290	10.7860	0.5376	-0.4880	7.4546	YRI
KCNMB3	GI_25952101-I	rs3739038	3	2	238454703	23.3168	0.8278	-0.5972	7.4546	YRI
EGLN2	GI_18141577-A	rs7222388	19	17	54447817	8.1175	0.4305	0.3988	8.0166	YRI
PARC	GI_24307990-S	rs11578336	6	1	163690368	8.1484	0.4318	-0.1743	7.2101	YRI
FLJ10579	GI_8922531-S	rs12435621	15	14	97182294	7.9215	0.4217	0.1925	7.4600	YRI
COPS7A	GI_7705329-S	rs4647958	12	20	48034038	7.5069	0.4028	0.1493	7.0327	YRI
ATP9B	GI_41327759-S	rs643788	18	11	118472968	7.4540	0.4003	-0.1331	7.1528	YRI
ZNF322B	GI_39841070-S	rs3736781	9	6	26613341	14.4872	0.6543	-0.3385	7.1958	YRI
FLJ33979	GI_22749298-S	rs2444007	11	15	42834417	12.8642	0.6136	0.7662	8.5900	YRI
LOC154428	GI_18564351-S	rs1351419	6	4	71382148	7.9075	0.4211	0.0502	7.1641	YRI
MAP2K5	GI_21729898-A	rs3924661	15	10	4919527	7.4791	0.4240	-0.1760	7.1276	YRI