

CHI3L2	GI_11993934-S	rs942697	1	111503545	5615	6.81	0.3696	6.99	5.98	4.82	3.94	CEU
CHRNE	GI_38327653-S	rs2302321	17	4728587	13594	10.4232	0.5243	9.2	7.05	5.42	4.3	CEU
COPS7A	GI_7705329-S	rs714774	12	6489545	221475	5.4052	0.2975	6.37	5.35	4.5	3.74	CEU
CP110	GI_34222317-S	rs1475963	16	19412783	59263	6.262	0.3424	6.13	5.37	4.32	3.58	CEU
CPEB4	GI_32698754-S	rs10516107	5	173280762	36825	10.8238	0.539	5.97	5.68	4.58	3.82	CEU
CPNE1	GI_23397697-A	rs6060535	20	33698936	21264	12.4017	0.5927	7.23	5.63	4.29	3.36	CEU
CRIP1	GI_41350204-S	rs1451156	2	46744253	19148	8.1268	0.4309	6.93	5.72	4.54	3.84	CEU
CRNKL1	GI_30795219-S	rs2073123	20	19978870	15505	5.7006	0.3133	7.28	5.53	4.5	3.73	CEU
CSTB	GI_20357564-S	rs1041456	21	44030783	12403	8.2125	0.4403	6.75	5.76	4.68	3.87	CEU
CSTF3	GI_4557494-S	rs3758741	11	33063192	29	5.3589	0.295	5.85	5.3	4.38	3.58	CEU
CTNS	GI_4826681-S	rs161370	17	3477002	34422	6.562	0.3574	6.7	5.44	4.54	3.81	CEU
CTSH	GI_23110956-A	rs1036938	15	77024302	22891	8.4809	0.4464	6.3	5.45	4.56	3.74	CEU
CTSK	GI_23110958-S	rs1136774	1	147551270	30875	7.2458	0.3906	6.89	5.43	4.24	3.45	CEU
DCLRE1B	GI_24431998-S	rs878129	1	114171237	2889	8.7891	0.4595	7.2	5.63	4.44	3.67	CEU
DCTD	GI_4503276-S	rs7277	4	184186662	176	8.596	0.4513	6.89	5.77	4.65	3.8	CEU
DERP6	GI_44662825-I	rs4562	17	7104463	8022	8.5043	0.4474	7.01	5.91	4.56	3.78	CEU