

Analysis Summary: Parkinsons Disease (Population Controls)

Phenotype Description

Cases answered "Yes" or reported a diagnosis of Parkinson's disease to at least one of the questions below or joined 23andMe as part of Parkinson's Disease Research Initiative and provided no survey responses (that is, if they didn't provide any information to the contrary, we assumed they truly have PD):

- (PD) Background Survey ("Has a doctor ever diagnosed you with Parkinson's disease?" Answer = Yes)
- Your Medical History ("Have you ever been diagnosed by a doctor with Parkinson's disease"? Answer = Yes)
- Research Snippets ("Has a doctor ever diagnosed you with Parkinson's disease?" Answer = Yes)
- Your Health Profile ("Have you ever been diagnosed or treated for any of the following conditions?" Answer = Parkinson's disease)
- Roots into the Future: Health History ("Has a doctor ever told you that you have any of these conditions?" Answer = Parkinson's disease)
- (PD) Baseline Survey ("Has a doctor ever diagnosed you with Parkinson's disease?" Answer = Yes)

Cases were further filtered by removing respondents 1) who were discordant between the first two questions above (that is, removing those who said "No" to the PD Background Survey question but said "Yes" to the Your Medical History question); 2) whose diagnosis changed because their symptoms disappeared or the cause of the symptoms is unknown (according to the questions below);

- (PD) Background Survey ("Is your diagnosis still Parkinson's disease or has your diagnosis changed?" Answer = My diagnosis changed)
- (PD) Background Survey ("Which of the following best describes how your diagnosis changed?" Answer = I no longer have the symptoms that led to the Parkinson's diagnosis, I still have the symptoms but the cause is unknown, My diagnosis changed)

or 3) whose diagnosis changed to another diagnosis that could be confounded with PD based on the following questions:

- (PD) Background Survey ("Which of the following have you been diagnosed with? Please check all that apply." Answers = Progressive Supranuclear Palsy (PSP); Multiple system atrophy (MSA), Shy-Drager syndrome, or striatonigral degeneration; Spinal cerebellar ataxia (SCA), cerebellar degeneration, olivopontocerebellar atrophy (OPCA); Cortical basal ganglionic degeneration, Parkinsonism due to medications, drug-induced parkinsonism)
- (PD) Background Survey ("Which of the following have you been diagnosed with? Please check all that apply." Answers = Frontotemporal dementia (FTD); Picks disease)
- (PD) Background Survey ("Which of the following have you been diagnosed with? Please check all that apply." Answers = Dystonic tremor; Rubral tremor; Primary writing tremor; Cerebellar tremor; Multiple sclerosis (MS))

Because PD is rare, we did not require controls to affirmatively deny a diagnosis of PD. Instead, population controls were used with the following groups excluded:

- Anyone defined as a case by the first set of criteria described above
- Anyone who was diagnosed with "Another form of parkinsonism (other than Parkinson's disease)", "Dementia, cognitive impairment, or senility", "A tremor disorder" according to this question:
 - (PD) Background Survey ("Has a doctor ever diagnosed you with any of the following? Please check all that apply.)
- Anyone reporting a diagnosis of Alzheimer's disease by answering "Yes" to the following question:
 - Your Medical History ("Have you ever been diagnosed by a doctor with Alzheimer's disease?")
- Anyone reporting a diagnosis of memory loss by answering "Yes" to the following question:
 - Your Medical History ("Have you ever been diagnosed by a doctor with memory loss?")
- Anyone with three or more early symptoms of Parkinson's as defined by the following questions below:
 - (PD) Background Survey ("Do you have trembling or shaking of any part of your body?" Yes is + 1)
 - (PD) Background Survey ("Have you noticed or have people told you that your speech or voice has become softer than before?" Yes is + 1)
 - (PD) Background Survey ("When walking, do you swing your arms less than you used to?" Yes is + 1)
 - (PD) Background Survey ("When standing or walking do you stoop or bend forward more than you used to?" Yes is + 1)
 - (PD) Background Survey ("Do you have trouble with your balance or fall sometimes?" Yes is + 1)
 - (PD) Background Survey ("Have you noticed any of the following changes in your handwriting? Please check all that apply." Answers = Handwriting became slower, Handwriting became smaller, Handwriting became shakier (each answer is + 1))
 - (PD) Background Survey ("Have you noticed any of the following changes in the way you walk? Please check all that apply." Answers = Dragging on or both feet, Feet shuffling, Walking more slowly, Taking smaller steps than before, Steps becoming faster and faster, Feet getting stuck as if glued to the floor (each answer is + 1))

Phenotype Statistics

The following table shows demographics of unrelated, European individuals included in the GWAS.

Phenotype	Group	Total	M	F	(0,30]	(30,45]	(45,60]	(60,Inf]
PD_all_pop_con	case	9619	5875	3744	33	330	1959	7297
PD_all_pop_con	control	324522	174938	149584	44241	93985	87425	98871

The following table shows the phenotypic distribution across 23andMe genotyping platforms for individuals included in the GWAS.

Phenotype	Group	Total	v1/v2	v3	v4
PD_all_pop_con	case	9619	2947	6300	372
PD_all_pop_con	control	324522	21205	233307	70010

Null Model with Covariates

The following table shows results of fitting a model for the trait based on just the covariates. Principal coordinates have been standardized, so these effect sizes are in units of standard deviations.

	Estimate	Std. Error	z value	Pr(> z)	LRT	Pr(>Chi)
age	0.0673	0.000731	92.1	0.0	10129.9	0.0
sexF	-0.3477	0.021686	-16.0	7.4×10^{-58}	261.9	6.6×10^{-59}
pc.0	-0.0111	0.009583	-1.2	0.25	1.3	0.25
pc.1	-0.1161	0.012860	-9.0	1.8×10^{-19}	88.4	5.5×10^{-21}
pc.2	-0.0647	0.011090	-5.8	5.5×10^{-9}	35.2	3.0×10^{-9}
pc.3	0.0442	0.010795	4.1	4.3×10^{-5}	16.7	4.4×10^{-5}
pc.4	0.0609	0.012376	4.9	8.6×10^{-7}	26.2	3.1×10^{-7}

SNP-level QC information

The following table shows results for QC filters on the genotyped data:

	failed	passed
no filters	0	1030430
not V1/V2-only, chrM, chrY	33033	997397
parent-offspring test	2129	995294
MAF > 0%	3203	992310
HWE > 1e-20	48225	945039
gt.rate > 90%	30775	925423
batch effects	71627	889734

The following table shows results for QC filters on the imputed dosage data:

	failed	passed
no filters	0	13733809
MAF > 0%	0	13733809
imputation quality	0	13733809
batch effects	1123116	12610693

The following table shows results for QC filters on the merged association test results:

	passed	total
imputed only	11813480	11813480
both passed	797198	12610678
genotyped only	92536	12703214
no test result	-1446	12701768
failed to converge	-263549	12438219

Genetic Association Tests

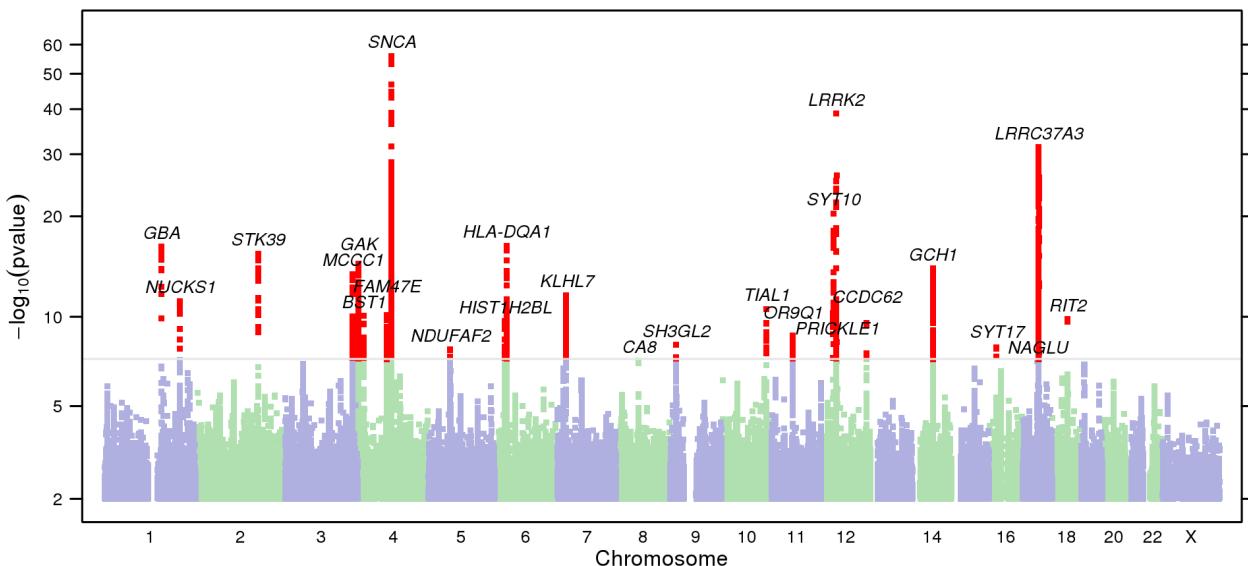
We performed logistic regression assuming an additive model for allelic effects, using the model:

PD_all_pop_con ~ age + sex + pc.0 + pc.1 + pc.2 + pc.3 + pc.4 + genotype

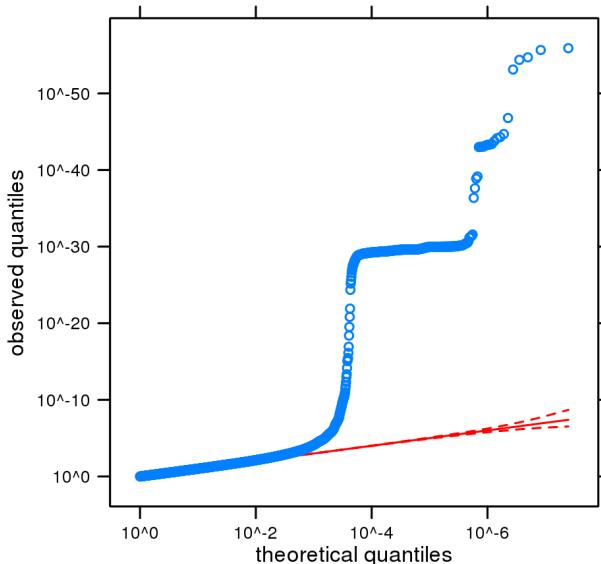
This genome-wide association analysis includes data from 9619 cases and 324522 controls of European ancestry, filtered to remove close relatives.

The results in this report have been adjusted for a genomic control inflation factor $\lambda=1.131$. The equivalent inflation factor for 1000 cases and 1000 controls $\lambda_{1000}= 1.007$, and for 10000, $\lambda_{10000}= 1.070$.

Manhattan Plot



Q-Q Plot of GWAS Results



Index SNPs for Strongest Associations

cytoband	assay.name	scaffold	position	alleles	src	pvalue	OR	95% CI	gene.context
4q22.1	rs356182	chr4	90626111	A/G	I	1.2×10^{-56}	1.318	[1.274,1.363]	GPRIN3---[]--SNCA
12q12	rs28903073	chr12	40653510	A/G	I	1.3×10^{-39}	0.321	[0.276,0.373]	[LRRK2]
17q21.31	rs365825	chr17	43705601	A/G	I	4.2×10^{-32}	0.786	[0.754,0.819]	LRRC37A3---[]--CRHR1
12p11.1	rs144847051	chr12	33301805	A/G	I	4.1×10^{-21}	0.447	[0.383,0.521]	PKP2---[]--SYT10
6p21.32	rs17425622	chr6	32571961	C/T	I	3.8×10^{-17}	1.235	[1.174,1.298]	HLA-DRB5---[]--HLA-DQA1
1q22	rs2230288	chr1	155206167	C/T	I	4.7×10^{-17}	1.674	[1.495,1.873]	[GBA]
2q24.3	rs13016703	chr2	169155595	G/T	I	2.7×10^{-16}	0.827	[0.791,0.865]	STK39---[]--CERS6
4p16.3	rs35541465	chr4	873286	D/I	I	2.8×10^{-15}	0.865	[0.834,0.896]	[GAK]
14q22.2	rs11158026	chr14	55348869	C/T	I	8.8×10^{-15}	0.873	[0.844,0.904]	[GCH1]
3q27.1	rs9858038	chr3	182730291	A/G	I	3.0×10^{-14}	1.172	[1.124,1.221]	DCUN1D1---[]--MCCC1
7p15.3	rs10256359	chr7	23123877	C/T	I	2.1×10^{-12}	0.887	[0.858,0.917]	FAM126A---[]--KLHL7
1q32.1	rs823118	chr1	205723572	C/T	I	6.3×10^{-12}	1.117	[1.082,1.153]	NUCKS1---[]--RAB7L1
10q26.11	rs188789342	chr10	121372328	C/T	I	2.6×10^{-11}	1.488	[1.331,1.663]	TIAL1---[]--BAG3
4q21.1	rs6812193	chr4	77198986	C/T	I	7.3×10^{-11}	0.897	[0.868,0.927]	[FAM47E]
4p15.32	rs4266290	chr4	15737118	C/G	I	8.4×10^{-11}	1.128	[1.088,1.171]	BST1---[]--CD38
18q12.3	rs4130047	chr18	40678235	C/T	I	1.5×10^{-10}	0.897	[0.868,0.927]	[RIT2]
6p22.1	rs4713118	chr6	27676794	A/G	I	2.0×10^{-10}	1.121	[1.083,1.162]	ZNF184---[]--HIST1H2BL
12q24.31	rs11060180	chr12	123303586	A/G	I	2.6×10^{-10}	0.903	[0.875,0.932]	[CCDC62]
11q12.1	rs687432	chr11	57694260	A/C	I	2.0×10^{-9}	1.113	[1.075,1.153]	CTNN1D1---[]--OR9Q1
12q12	rs148294058	chr12	43049382	A/G	I	5.2×10^{-9}	0.637	[0.551,0.736]	PRICKLE1---[]--ADAMTS20
9p22.2	rs2209440	chr9	17736342	G/T	I	7.5×10^{-9}	0.912	[0.884,0.941]	[SH3GL2]
16p12.3	rs11343	chr16	19279464	G/T	I	1.0×10^{-8}	1.096	[1.062,1.131]	SYT17---[]--CLEC19A
5q12.1	rs162227	chr5	60260543	C/T	I	1.4×10^{-8}	0.860	[0.817,0.905]	[NDUFAF2]
8q12.1	rs144074972	chr8	61163895	C/T	I	6.1×10^{-8}	Inf	[0.022, Inf]	[CA8]
17q21.2	rs601999	chr17	40698158	C/T	I	6.7×10^{-8}	1.095	[1.060,1.132]	GLU---[]--HSD17B1

Quality Statistics for Index SNPs

assay.name	is.v2	is.v3	is.v4	gt.rate	hw.p.value	p.date	freq.b	avg.rsqr	min.rsqr	p.batch	dose.b	qc.mask
rs356182	FALSE	FALSE	FALSE				0.8767	0.8611	0.20	0.3572	v2v3v4	
rs28903073	TRUE	TRUE	TRUE	0.9998	5.4×10 ⁻⁵	2.3×10 ⁻¹⁰	0.9956	0.9748	0.9575	0.00053	0.9956	v2v3v4
rs365825	FALSE	FALSE	FALSE				0.9407	0.8998	1.5×10 ⁻⁶	0.2265	v2v3v4	
rs144847051	FALSE	FALSE	FALSE				0.7436	0.6980	0.0047	0.9926	v2v3v4	
rs17425622	FALSE	FALSE	FALSE				0.8535	0.7666	0.22	0.8528	v2v3v4	
rs2230288	FALSE	TRUE	TRUE	0.9823	0.012	0.0	0.0036	0.9545	0.8988	0.027	0.0128	v2v3v4
rs13016703	FALSE	FALSE	TRUE	0.9999	0.28	0.19	0.8747	0.9958	0.9751	0.037	0.8733	v2v3v4
rs35541465	FALSE	FALSE	FALSE				0.9445	0.8958	7.1×10 ⁻⁶	0.7323	v2v3v4	
rs11158026	FALSE	FALSE	TRUE	1.0000	0.92	0.64	0.3317	0.9714	0.9591	0.20	0.3303	v2v3v4
rs9858038	FALSE	FALSE	FALSE				0.9839	0.9695	0.87	0.8029	v2v3v4	
rs10256359	FALSE	FALSE	FALSE				0.9648	0.8961	0.35	0.3695	v2v3v4	
rs823118	FALSE	FALSE	FALSE				0.9979	0.9954	0.61	0.5585	v2v3v4	
rs188789342	FALSE	FALSE	FALSE				0.8154	0.7608	0.17	0.0176	v2v3v4	
rs6812193	TRUE	TRUE	TRUE	1.0000	0.018	0.014	0.3677	1.0000	0.9999	0.21	0.3677	v2v3v4
rs4266290	FALSE	FALSE	FALSE				0.9966	0.9924	0.69	0.7474	v2v3v4	
rs4130047	TRUE	TRUE	TRUE	1.0000	0.54	0.15	0.6854	0.9999	0.9995	0.30	0.6854	v2v3v4
rs4713118	FALSE	TRUE	FALSE	0.9995	4.3×10 ⁻²⁹	0.85	0.2517	0.9960	0.9857	0.00011	0.2527	v2v3v4
rs11060180	TRUE	TRUE	TRUE	0.9997	0.47	0.38	0.4498	0.9990	0.9971	0.69	0.4489	v2v3v4
rs687432	FALSE	FALSE	FALSE				0.9756	0.9138	0.056	0.7042	v2v3v4	
rs148294058	FALSE	FALSE	FALSE				0.5194	0.4686	0.094	0.9830	v2v3v4	
rs2209440	TRUE	TRUE	TRUE	0.9990	0.29	0.22	0.5236	0.9979	0.9927	0.33	0.5244	v2v3v4
rs111343	TRUE	TRUE	TRUE	0.9999	0.29	2.4×10 ⁻⁵	0.4432	0.9982	0.9938	0.00024	0.4435	v2v3v4
rs162227	FALSE	FALSE	FALSE				0.9206	0.8771	4.2×10 ⁻⁵	0.8965	v2v3v4	
rs144074972	FALSE	FALSE	FALSE				0.9007	0.7877	0.81	0.9993	v2v3v4	
rs601999	FALSE	FALSE	FALSE				0.9862	0.9781	0.11	0.3272	v2v3v4	

SNP Statistics in the GWAS Sample

assay.name	AA.0	AB.0	BB.0	im.num.0	dose.b.0	AA.1	AB.1	BB.1	im.num.1	dose.b.1
rs356182				324522	0.3570				9619	0.4127
rs28903073	7	2616	321831	324522	0.9960	4	237	9374	9619	0.9872
rs365825				324522	0.2271				9619	0.1925
rs144847051				324522	0.9928				9619	0.9862
rs17425622				324522	0.8521				9619	0.8678
rs2230288	295942	2056	0	324522	0.0128	6407	22	0	9619	0.0210
rs13016703	1087	15357	53557	324522	0.8743	9	93	270	9619	0.8541
rs35541465				324522	0.7355				9619	0.7084
rs11158026	31249	31042	7718	324522	0.3313	166	175	31	9619	0.3036
rs9858038				324522	0.8032				9619	0.8271
rs10256359				324522	0.3700				9619	0.3422
rs823118				324522	0.5569				9619	0.5836
rs188789342				324522	0.0176				9619	0.0244
rs6812193	129661	150504	44346	324522	0.3686	4130	4351	1136	9619	0.3445
rs4266290				324522	0.7471				9619	0.7660
rs4130047	31948	140038	152532	324522	0.6858	1103	4276	4240	9619	0.6630
rs4713118	131746	85842	15592	324522	0.2528	3349	2441	505	9619	0.2738
rs11060180	97845	160924	65659	324522	0.4504	3124	4797	1695	9619	0.4258
rs687432				324522	0.7042				9619	0.7239
rs148294058				324522	0.9833				9619	0.9785
rs2209440	73044	162130	88997	324522	0.5252	2397	4767	2450	9619	0.5030
rs111343	101021	159837	63634	324522	0.4426	2791	4703	2125	9619	0.4655
rs162227				324522	0.8978				9619	0.8855
rs144074972				324522	0.9994				9619	1.0000
rs601999				324522	0.3280				9619	0.3466

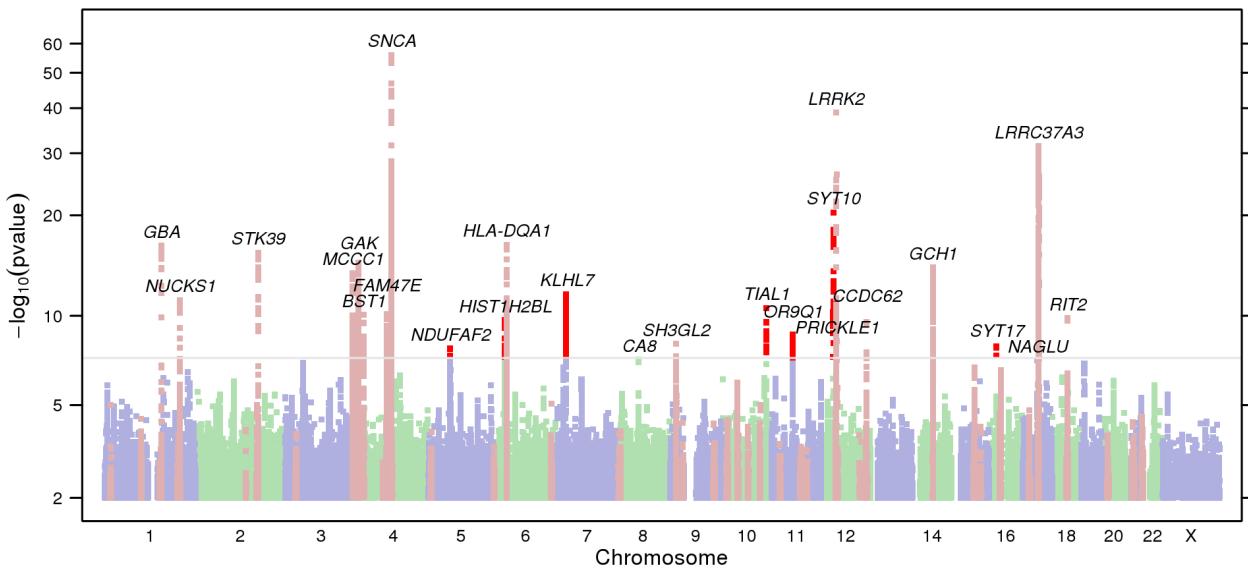
Annotations from NHGRI GWAS Catalog

The following table shows, for each index SNP, all entries in the NHGRI GWAS Catalog that are within 500kb and in at least moderate linkage disequilibrium ($r^2 > 0.5$).

region	position	our.name	our.pval	dist	rsqr	assay.name	pvalue	pubmed.id	trait	genes
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	11490	0.758	rs356219	6.0×10 ⁻⁶⁵	22438815	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	11490	0.758	rs356219	2.0×10 ⁻⁴⁷	21292315	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	15229	0.520	rs356220	3.0×10 ⁻¹¹	24511991	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	15229	0.520	rs356220	8.0×10 ⁻³⁵	22451204	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	15229	0.520	rs356220	2.0×10 ⁻¹⁹	21738487	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	15229	0.520	rs356220	3.0×10 ⁻⁸	21084426	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	15229	0.520	rs356220	9.0×10 ⁻¹⁶	21044948	Parkinson's disease	SNCA
4q22.1	90626111	rs356182	1.2×10 ⁻⁵⁶	15229	0.520	rs356220	3.0×10 ⁻¹¹	20711177	Parkinson's disease	SNCA
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-192160	0.707	rs11012	6.0×10 ⁻⁸	20070850	Parkinson's disease	PLEKHM1, MAPT, IMP5
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-189199	0.707	rs17631303	1.0×10 ⁻⁸	23544013	Ovarian cancer in BRCA1 mutation carriers	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	9249	0.993	rs2942168	1.0×10 ⁻²⁸	21292315	Parkinson's disease	MAPT
17q21.31	43705601	rs365825	4.2×10 ⁻³²	13542	1.000	rs393152	2.0×10 ⁻¹⁶	19915575	Parkinson's disease	MAPT, C17orf69, ...
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218082	0.993	rs12185268	3.0×10 ⁻¹⁴	21738487	Parkinson's disease	MAPT
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218618	0.985	rs12373124	5.0×10 ⁻¹⁰	22693459	Male-pattern baldness	Intergenic
17q21.31	43705601	rs365825	4.2×10 ⁻³²	219696	0.803	rs17690703	6.0×10 ⁻⁹	24429156	Idiopathic pulmonary fibrosis	SPPL2C, CRHR1, ...
17q21.31	43705601	rs365825	4.2×10 ⁻³²	272226	0.993	rs1864325	5.0×10 ⁻¹¹	22504420	Bone mineral density	MAPT
17q21.31	43705601	rs365825	4.2×10 ⁻³²	351166	0.993	rs1981997	9.0×10 ⁻¹⁴	23583980	Interstitial lung disease	MAPT
17q21.31	43705601	rs365825	4.2×10 ⁻³²	375463	0.985	rs8070723	2.0×10 ⁻¹¹⁸	21685912	Progressive supranuclear palsy	MAPT
17q21.31	43705601	rs365825	4.2×10 ⁻³²	375463	0.985	rs8070723	7.0×10 ⁻¹²	21044948	Parkinson's disease	MAPT
17q21.31	43705601	rs365825	4.2×10 ⁻³²	48186	0.964	rs9303525	8.0×10 ⁻¹⁵	22504418	Intracranial volume	MAPT, GRN, ...
17q21.31	43705601	rs365825	4.2×10 ⁻³²	481891	0.993	rs17577094	8.0×10 ⁻⁹	24842889	Parkinson's disease	MAPT
1q22	155206167	rs2230288	4.7×10 ⁻¹⁷	-98000	0.889	rs12726330	5.0×10 ⁻⁸	22451204	Parkinson's disease	GBA
2q24.3	16915595	rs13016703	2.7×10 ⁻¹⁶	-38570	0.697	rs2102808	4.0×10 ⁻¹⁰	21292315	Parkinson's disease	STK39
2q24.3	16915595	rs13016703	2.7×10 ⁻¹⁶	-12560	1.000	rs9917256	3.0×10 ⁻⁶	22451204	Parkinson's disease	Intergenic
4p16.3	873286	rs35541465	2.8×10 ⁻¹⁵	65801	0.677	rs6599388	4.0×10 ⁻¹²	21292315	Parkinson's disease	GAK
3q27.1	182730291	rs9858038	3.0×10 ⁻¹⁴	24918	0.708	rs2270968	1.0×10 ⁻¹⁶	24816252	Blood metabolite levels	MCCC1
3q27.1	182730291	rs9858038	3.0×10 ⁻¹⁴	29782	1.000	rs10513789	3.0×10 ⁻¹⁰	21738487	Parkinson's disease	MCCC1, LAMP3
3q27.1	182730291	rs9858038	3.0×10 ⁻¹⁴	90984	0.594	rs1171441	8.0×10 ⁻¹²	21292315	Parkinson's disease	MCCC1, LAMP3
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	-33765	0.570	rs823094	2.0×10 ⁻⁷	23449627	Pubertal anthropometrics	NUCKS1, RAB7L1

1q32.1	205723572	rs823118	6.3×10^{-12}	-4040	0.995	rs823114	3.0×10^{-6}	24842889	Parkinson's disease	PARK16
4q21.1	77198986	rs6812193	7.3×10^{-11}	0	1.000	rs6812193	8.0×10^{-10}	21738487	Parkinson's disease	SCARB2
4q21.1	77198986	rs6812193	7.3×10^{-11}	0	1.000	rs6812193	4.0×10^{-7}	19915575	Parkinson's disease	STBD1
4p15.32	15737118	rs4266290	8.4×10^{-11}	819	0.583	rs4538475	3.0×10^{-9}	19915576	Parkinson's disease	BST1
18q12.3	40678235	rs4130047	1.5×10^{-10}	-4855	1.000	rs12456492	2.0×10^{-10}	22451204	Parkinson's disease	RIT2
18q12.3	40678235	rs4130047	1.5×10^{-10}	0	1.000	rs4130047	2.0×10^{-7}	21738487	Parkinson's disease	RIT2, SYT4
6p22.1	27676794	rs4713118	2.0×10^{-10}	344026	0.502	rs172166	5.0×10^{-6}	23247143	Cardiac Troponin-T levels	OR2B7P

Replication of GWAS Catalog Results



The following table shows, for each GWAS Catalog result for similar traits, our association test result for our best available proxy (distance < 100kb, $r^2 > 0.8$).

region	position	our.name	our.pval	dist	rsqr	assay.name	pvalue	pubmed.id	trait	genes
4p16.3	852313	rs1564282	1.3×10^{-9}	0	1.000	rs1564282	7.0×10^{-7}	18985386	Parkinson's disease (familial)	GAK, DGKQ
1p36.13	19139517	rs12063142	0.24	0	1.000	rs12063142	5.0×10^{-7}	20070850	Parkinson's disease	TAS1R2
1p21.2	101880005	rs2338971	0.23	0	1.000	rs2338971	5.0×10^{-10}	24511991	Parkinson's disease	HUSEYO
1q21.3	154855055	rs1630500	0.0086	0	1.000	rs1630500	2.0×10^{-8}	24842889	Parkinson's disease	GBA
1q22	155108167	rs12726330	4.9×10^{-16}	0	1.000	rs12726330	5.0×10^{-8}	22451204	Parkinson's disease	GBA
1q22	156030037	rs34372695	1.3×10^{-5}	0	1.000	rs34372695	4.0×10^{-12}	21292315	Parkinson's disease	SYT11
1q32.1	205713378	rs823128	7.7×10^{-6}	0	1.000	rs823128	7.0×10^{-8}	19915575	Parkinson's disease	PARK16, NUCKS1
1q32.1	205719532	rs823114	6.6×10^{-12}	0	1.000	rs823114	3.0×10^{-6}	24842889	Parkinson's disease	PARK16
1q32.1	205752665	rs947211	6.8×10^{-6}	0	1.000	rs947211	2.0×10^{-12}	19915576	Parkinson's disease	PARK16, SLC45A3, NUCKS1, RAB7L1, SLC41A1, PM20D1
1q32.1	205764640	rs823156	2.5×10^{-6}	0	1.000	rs823156	1.0×10^{-7}	21738487	Parkinson's disease	SLC41A1
2q21.3	135539967	rs6430538	0	1.000	rs6430538	2.0×10^{-7}	22451204	Parkinson's disease	Intergenic	
2q24.3	169117025	rs2102808	6.5×10^{-16}	0	1.000	rs2102808	4.0×10^{-10}	21292315	Parkinson's disease	STK39
2q24.3	169143035	rs9917256	8.0×10^{-15}	0	1.000	rs9917256	3.0×10^{-6}	22451204	Parkinson's disease	Intergenic
3p24.1	28705764	rs7617877	0.0095	0	1.000	rs7617877	3.0×10^{-6}	21044948	Parkinson's disease	Intergenic
3q27.1	182760073	rs10513789	1.1×10^{-13}	0	1.000	rs10513789	3.0×10^{-10}	21738487	Parkinson's disease	MCCC1, LAMP3
3q27.1	182821275	rs11711441	3.3×10^{-9}	0	1.000	rs11711441	8.0×10^{-12}	21292315	Parkinson's disease	MCCC1, LAMP3
4p16.3	858332	rs11248051	2.4×10^{-9}	0	1.000	rs11248051	3.0×10^{-9}	20711177	Parkinson's disease	GAK
4p16.3	939087	rs6599388	0	1.000	rs6599388	4.0×10^{-12}	21292315	Parkinson's disease	GAK	
4p16.3	939113	rs6599389	8.7×10^{-9}	0	1.000	rs6599389	4.0×10^{-8}	21738487	Parkinson's disease	GAK
4p16.3	964359	rs11248060	2.3×10^{-11}	0	1.000	rs11248060	3.0×10^{-9}	22451204	Parkinson's disease	DGKQ
4p16.3	964359	rs11248060	2.3×10^{-11}	0	1.000	rs11248060	3.0×10^{-12}	22438815	Parkinson's disease	GAK, DGKQ
4p16.3	964359	rs11248060	2.3×10^{-11}	0	1.000	rs11248060	2.0×10^{-6}	20070850	Parkinson's disease	DGKQ
4p15.32	15737101	rs11724635	2.3×10^{-7}	0	1.000	rs11724635	1.0×10^{-16}	21292315	Parkinson's disease	BST1
4p15.32	15737348	rs4698412	2.5×10^{-7}	0	1.000	rs4698412	3.0×10^{-7}	22451204	Parkinson's disease	BST1
4p15.32	15737348	rs4698412	2.5×10^{-7}	0	1.000	rs4698412	2.0×10^{-6}	21084426	Parkinson's disease	BST1
4p15.32	15737937	rs4538475	2.1×10^{-6}	0	1.000	rs4538475	3.0×10^{-9}	19915576	Parkinson's disease	BST1
4q13.2	68447249	rs2242330	0.70	0	1.000	rs2242330	2.0×10^{-6}	17052657	Parkinson's disease	BRDG1
4q21.1	77198986	rs6812193	7.3×10^{-11}	0	1.000	rs6812193	8.0×10^{-10}	21738487	Parkinson's disease	SCARB2
4q21.1	77198986	rs6812193	7.3×10^{-11}	0	1.000	rs6812193	4.0×10^{-7}	19915575	Parkinson's disease	STBD1
4q22.1	90637601	rs356219	2.2×10^{-56}	0	1.000	rs356219	6.0×10^{-65}	22438815	Parkinson's disease	SNCA
4q22.1	90637601	rs356219	2.2×10^{-56}	0	1.000	rs356219	2.0×10^{-47}	21292315	Parkinson's disease	SNCA
4q22.1	90639515	rs11931074	2.1×10^{-21}	0	1.000	rs11931074	7.0×10^{-17}	19915576	Parkinson's disease	SNCA
4q22.1	90641340	rs356220	7.6×10^{-54}	0	1.000	rs356220	3.0×10^{-11}	24511991	Parkinson's disease	SNCA
4q22.1	90641340	rs356220	7.6×10^{-54}	0	1.000	rs356220	8.0×10^{-35}	22451204	Parkinson's disease	SNCA
4q22.1	90641340	rs356220	7.6×10^{-54}	0	1.000	rs356220	2.0×10^{-19}	21738487	Parkinson's disease	SNCA
4q22.1	90641340	rs356220	7.6×10^{-54}	0	1.000	rs356220	3.0×10^{-8}	21084426	Parkinson's disease	SNCA
4q22.1	90641340	rs356220	7.6×10^{-54}	0	1.000	rs356220	9.0×10^{-16}	21044948	Parkinson's disease	SNCA
4q22.1	90641340	rs356220	7.6×10^{-54}	0	1.000	rs356220	3.0×10^{-11}	20711177	Parkinson's disease	SNCA
4q22.1	90678541	rs2736990	9.9×10^{-44}	0	1.000	rs2736990	7.0×10^{-8}	20070850	Parkinson's disease	SNCA
4q22.1	90678541	rs2736990	9.9×10^{-44}	0	1.000	rs2736990	2.0×10^{-16}	19915575	Parkinson's disease	SNCA
4q22.1	90780902	rs6532194	0.014	0	1.000	rs6532194	5.0×10^{-11}	22438815	Parkinson's disease	SNCA
4q22.1	90797301	rs6532197	1.8×10^{-7}	0	1.000	rs6532197	1.0×10^{-7}	19915575	Parkinson's disease	MMRN1
5p15.31	9332281	rs7702187	0.11	0	1.000	rs7702187	8.0×10^{-6}	165252231	Parkinson's disease	SEMA5A
5q35.3	179794208	rs10464059	0.66	0	1.000	rs10464059	3.0×10^{-6}	20070850	Parkinson's disease	
6p21.32	32387809	rs2395163	1.6×10^{-10}	0	1.000	rs2395163	3.0×10^{-11}	22451204	Parkinson's disease	LOC642072
6p21.32	32409530	rs3129882	0.00032	0	1.000	rs3129882	5.0×10^{-10}	24511991	Parkinson's disease	HLA

6p21.32	32409530	rs3129882	0.00032	0	1.000	rs3129882	2.0×10 ⁻¹⁰	20711177	Parkinson's disease	HLA-DRA
6q25.2	154726988	rs2275336	0.43	0	1.000	rs2275336	9.0×10 ⁻⁷	22451204	Parkinson's disease	CNKS3
8p23.1	11698747	rs1296028	0.0048	0	1.000	rs1296028	1.0×10 ⁻⁶	22451204	Parkinson's disease	Intergenic
9p22.2	17731921	rs1536076	2.3×10 ⁻⁷	0	1.000	rs1536076	5.0×10 ⁻⁷	22451204	Parkinson's disease	SH3GL2
9p13.3	35269819	rs10121009	0.56	0	1.000	rs10121009	3.0×10 ⁻⁶	21812969	Parkinson's disease	UNC13B
9q33.1	122059709	rs4837628	0.57	0	1.000	rs4837628	1.0×10 ⁻⁶	20070850	Parkinson's disease	DBC1
10p13	15561543	rs7077361	0.0083	0	1.000	rs7077361	5.0×10 ⁻⁶	19915575	Parkinson's disease	ITGA8
10q11.21	45161109	rs1480597	0.62	0	1.000	rs1480597	2.0×10 ⁻⁶	17052657	Parkinson's disease	Intergenic
10q22.1	71580120	rs17497526	0.26	0	1.000	rs17497526	6.0×10 ⁻⁷	23793441	Parkinson's disease	COL13A1
10q24.32	104591393	rs17115100	0.98	0	1.000	rs17115100	7.0×10 ⁻⁸	19915575	Parkinson's disease	CYP17A1, C10orf32, CNNM2, SFXN2
11p14.3	22114093	rs11026412	0.34	0	1.000	rs11026412	3.0×10 ⁻⁶	22451204	Parkinson's disease	Intergenic
11q14.1	84417846	rs10501570	0.91	0	1.000	rs10501570	7.0×10 ⁻⁶	17052657	Parkinson's disease	DLG2
11q21	96108428	rs7118648	0.78	0	1.000	rs7118648	2.0×10 ⁻⁷	23793441	Parkinson's disease	CCDC82
12q12	40352996	rs10877840	0.76	0	1.000	rs10877840	1.0×10 ⁻⁶	22451204	Parkinson's disease	SLC2A13
12q12	40428561	rs1994090	0.27	0	1.000	rs1994090	3.0×10 ⁻⁸	19915576	Parkinson's disease	LRRK2
12q12	40620808	rs1491942	7.9×10 ⁻⁸	0	1.000	rs1491942	6.0×10 ⁻¹⁵	22438815	Parkinson's disease	LRRK2
12q12	40620808	rs1491942	7.9×10 ⁻⁸	0	1.000	rs1491942	6.0×10 ⁻¹⁴	21292315	Parkinson's disease	LRRK2
12q12	40734202	rs34637584		0	1.000	rs34637584	2.0×10 ⁻²⁸	21738487	Parkinson's disease	LRRK2
12q12	40757328	rs34778348		0	1.000	rs34778348	3.0×10 ⁻²¹	22438815	Parkinson's disease	LRRK2
12q12	41365640	rs1442190	2.5×10 ⁻⁹	0	1.000	rs1442190	2.0×10 ⁻²⁷	24842889	Parkinson's disease	LRRK2
12q23.3	10694987	rs4964469	0.86	0	1.000	rs4964469	2.0×10 ⁻⁷	21084426	Parkinson's disease	Intergenic
12q24.31	123296294	rs12817488	3.8×10 ⁻¹⁰	0	1.000	rs12817488	3.0×10 ⁻¹³	21292315	Parkinson's disease	CCDC62, HIP1R
14q22.2	54290830	rs12431733	0.043	0	1.000	rs12431733	3.0×10 ⁻⁶	19915575	Parkinson's disease	BMP4
15q22.2	62001132	rs10519131	1.2×10 ⁻⁵	0	1.000	rs10519131	1.0×10 ⁻⁶	22451204	Parkinson's disease	Intergenic
15q25.1	81626725	rs3935740	0.92	0	1.000	rs3935740	2.0×10 ⁻⁷	23793441	Parkinson's disease	TMC3
16p11.2	31095171	rs11865038	4.2×10 ⁻⁵	0	1.000	rs11865038	4.0×10 ⁻⁷	22451204	Parkinson's disease	POL3S
17p11.2	17715101	rs11868035	1.6×10 ⁻⁵	0	1.000	rs11868035	6.0×10 ⁻⁸	21738487	Parkinson's disease	SREBF1, RAI1
17q21.31	43513441	rs11012	1.9×10 ⁻²⁸	0	1.000	rs11012	6.0×10 ⁻⁸	20070850	Parkinson's disease	PLEKHM1, MAPT, IMP5
17q21.31	43714850	rs2942168	4.0×10 ⁻³⁰	0	1.000	rs2942168	1.0×10 ⁻²⁸	21292315	Parkinson's disease	MAPT
17q21.31	43719143	rs393152	4.0×10 ⁻³⁰	0	1.000	rs393152	2.0×10 ⁻¹⁶	19915575	Parkinson's disease	MAPT, C17orf69, KIAA1267, LOC644246
17q21.31	43923683	rs12185268	9.9×10 ⁻³¹	0	1.000	rs12185268	3.0×10 ⁻¹⁴	21738487	Parkinson's disease	MAPT
17q21.31	44081064	rs8070723	7.0×10 ⁻³⁰	0	1.000	rs8070723	7.0×10 ⁻¹²	21044948	Parkinson's disease	MAPT
17q21.31	44187492	rs17577094	5.2×10 ⁻³⁰	0	1.000	rs17577094	8.0×10 ⁻⁹	24842889	Parkinson's disease	MAPT
17q21.31	44788310	rs183211	3.5×10 ⁻²¹	0	1.000	rs183211	3.0×10 ⁻⁷	21812969	Parkinson's disease	NSF
17q21.31	44828931	rs199533	4.4×10 ⁻²⁶	0	1.000	rs199533	1.0×10 ⁻⁶	20711177	Parkinson's disease	MAPT
17q21.31	44828931	rs199533	4.4×10 ⁻²⁶	0	1.000	rs199533	1.0×10 ⁻¹⁴	19915575	Parkinson's disease	NSF
17q21.31	44856641	rs199515	3.3×10 ⁻²⁵	0	1.000	rs199515	3.0×10 ⁻¹⁷	22451204	Parkinson's disease	WNT3
17q21.31	44859144	rs415430	4.3×10 ⁻²⁵	0	1.000	rs415430	7.0×10 ⁻⁷	21812969	Parkinson's disease	WNT3
17q21.31	44865603	rs199498	2.8×10 ⁻²³	0	1.000	rs199498	8.0×10 ⁻⁷	24511991	Parkinson's disease	MAPT
18q12.3	40673380	rs12456492	2.3×10 ⁻¹⁰	0	1.000	rs12456492	2.0×10 ⁻¹⁰	22451204	Parkinson's disease	RIT2
18q12.3	40678235	rs4130047	1.5×10 ⁻¹⁰	0	1.000	rs4130047	2.0×10 ⁻⁷	21738487	Parkinson's disease	RIT2, SYT4
20p12.1	13296912	rs1223271	0.83	0	1.000	rs1223271	5.0×10 ⁻⁶	19915575	Parkinson's disease	C20orf82
21q21.1	16914905	rs2823357	0.50	0	1.000	rs2823357	6.0×10 ⁻⁷	21738487	Parkinson's disease	USP25
21q22.3	43286352	rs2839398	0.36	0	1.000	rs2839398	5.0×10 ⁻⁶	22451204	Parkinson's disease	PRDM15

Nearby Nonsynonymous SNPs

region	position	our.name	our.pval	dist	rsqr	assay.name	gene	aa.chg
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-198304	0.594	rs12949256	ARHGPAP27	A117T
17q21.31	43705601	rs365825	4.2×10 ⁻³²	217341	0.993	rs62621252	SPPL2C	S224P
17q21.31	43705601	rs365825	4.2×10 ⁻³²	217665	0.993	rs62054815	SPPL2C	A332T
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218053	0.993	rs12185233	SPPL2C	R461P
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218082	0.993	rs12185268	SPPL2C	I471V
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218472	0.993	rs12373123	SPPL2C	S601P
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218529	0.993	rs12373139	SPPL2C	G620R
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218599	0.985	rs12373142	SPPL2C	P643R
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218618	0.985	rs12373124	SPPL2C	H649H
17q21.31	43705601	rs365825	4.2×10 ⁻³²	218630	0.985	rs12373140	SPPL2C	Q653Q
17q21.31	43705601	rs365825	4.2×10 ⁻³²	355174	0.993	rs63750417	MAPT	P202L
17q21.31	43705601	rs365825	4.2×10 ⁻³²	355422	0.993	rs62063786	MAPT	D285N
17q21.31	43705601	rs365825	4.2×10 ⁻³²	355435	0.993	rs62063787	MAPT	V289A
17q21.31	43705601	rs365825	4.2×10 ⁻³²	355677	0.993	rs17651549	MAPT	R370W
17q21.31	43705601	rs365825	4.2×10 ⁻³²	361799	0.993	rs10445337	MAPT	S447P
17q21.31	43705601	rs365825	4.2×10 ⁻³²	398810	0.942	rs199706121	MAPT	?1735?
17q21.31	43705601	rs365825	4.2×10 ⁻³²	402885	0.756	rs200384907	KANSL1	?1225?
17q21.31	43705601	rs365825	4.2×10 ⁻³²	403305	0.993	rs34579536	KANSL1	I108T
17q21.31	43705601	rs365825	4.2×10 ⁻³²	404931	0.993	rs36076725	KANSL1	F917F
17q21.31	43705601	rs365825	4.2×10 ⁻³²	404940	0.993	rs35833914	KANSL1	D914D
17q21.31	43705601	rs365825	4.2×10 ⁻³²	411518	0.993	rs34043286	KANSL1	S718P
6p21.32	32571961	rs17425622	3.8×10 ⁻¹⁷	38440	0.634	rs9272785	HLA-DQA1	A210T
1q22	155206167	rs2230288	4.7×10 ⁻¹⁷	57448	0.646	rs199893292	WDHD1	?1761?
14q22.2	55348869	rs11158026	8.8×10 ⁻¹⁵	24918	0.708	rs2270968	MCCC1	H464P

Nearby Expression QTLs

region	position	our.name	our.pval	dist	rsqr	assay.name	eqtl.dist	eqtl.gene	eqtl.pval	eqtl.rsqr	tissue	pubmed.id
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-194166	0.720	rs34465449	0	CRHR1-IT1	2.3×10 ⁻¹³	0.395	Whole_Blood	23715323
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-192160	0.707	rs11012	201708	MGC57346	6.7×10 ⁻²³	0.297	B-Cell	22446964
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-166164	0.563	rs117169618	0	CRHR1	2.1×10 ⁻¹¹	0.362	Lung	23715323
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-156075	0.722	rs17631676	42490	LRRC37A4	6.0×10 ⁻²⁰⁸	0.475	Monocyte	20502693
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-156075	0.722	rs17631676	165623	MGC57346	1.8×10 ⁻⁵⁹	0.613	Monocyte	22446964
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-156075	0.722	rs17631676	378322	HS_554608	0.00057	0.042	B-Cell	22446964
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-156075	0.722	rs17631676	170002	C17ORF69	7.7×10 ⁻⁵	0.055	B-Cell	22446964
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-150348	0.722	rs56192752	0	MAPT	3.0×10 ⁻⁶	0.196	Lung	23715323

17q21.31	43705601	rs365825	4.2×10 ⁻³²	-132705	0.676	rs5026246	0	CRHR1	2.3×10 ⁻¹⁰	0.397	Nerve_Tibial	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-131952	0.781	rs188710438	0	CRHR1-IT1	6.0×10 ⁻⁸	0.333	Esophagus_Mucosa	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-51133	0.892	rs149133346	0	DND1P1	4.8×10 ⁻⁷	0.309	Esophagus_Muscularis	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-43581	0.697	rs116954807	0	ARHGPAP27	6.1×10 ⁻⁷	0.238	Skin_Sun_Exposed_Lower_leg	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	-14964	0.963	rs147670174	0	CRHR1-IT1	4.8×10 ⁻²⁸	0.699	Lung	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	10554	0.685	rs112550936	0	798G7.5	4.9×10 ⁻⁸	0.336	Esophagus_Mucosa	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	13542	1.000	rs393152	580720	NMT1	0.00045	0.156	Lymphoblastoid	19644074	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	45997	0.978	chr17:43751598:D	0	LRRC37A4P	4.1×10 ⁻³³	0.799	Thyroid	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	45997	0.978	chr17:43751598:D	0	LRRC37A4P	6.7×10 ⁻²⁰	0.725	Heart_Left_Ventricle	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	147132	0.993	rs113871181	0	LRRC37A4P	1.1×10 ⁻³²	0.706	Whole_Blood	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	150738	0.578	rs117631329	446394	KANSL1	5.3×10 ⁻⁸	0.077	Lymphoblastoid	24037378	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	313506	0.689	rs111825734	0	LRRC37A	6.6×10 ⁻⁷	0.318	Heart_Left_Ventricle	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	376831	0.963	rs112310745	0	MAPT	3.2×10 ⁻⁶	0.344	Artery_Aorta	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	464476	0.725	chr17:44170077:D	0	669E14.6	1.5×10 ⁻⁶	0.228	Thyroid	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	491826	0.704	chr17:44197427:I	0	LRRC37A2	3.7×10 ⁻¹⁹	0.634	Nerve_Tibial	23715323	
17q21.31	43705601	rs365825	4.2×10 ⁻³²	491826	0.704	chr17:44197427:I	0	RP11-	259G18.2	4.1×10 ⁻³³	0.799	Thyroid	23715323
17q21.31	43705601	rs365825	4.2×10 ⁻³²	495757	0.971	chr17:44201358:D	0	LRRC37A4P	3.5×10 ⁻²³	0.851	Artery_Aorta	23715323	
6p21.32	32571961	rs17425622	3.8×10 ⁻¹⁷	-12057	0.573	rs35743245	246695	BPG246D15.9	1.2×10 ⁻⁶	0.061	Lymphoblastoid	24037378	
6p21.32	32571961	rs17425622	3.8×10 ⁻¹⁷	1601	0.602	rs602457	-1018080	LST1	0.00022	0.048	Monocyte	22446964	
6p21.32	32571961	rs17425622	3.8×10 ⁻¹⁷	8656	0.660	rs3997872	116324	HLA-DRB3	2.2×10 ⁻⁴⁸	0.136	Monocyte	20502693	
1q22	155206167	rs2230288	4.7×10 ⁻¹⁷	231544	0.830	rs35777901	-326859	RAG1AP1	0.00022	0.048	Monocyte	22446964	
2q24.3	169155595	rs13016703	2.7×10 ⁻¹⁶	1312	0.768	rs12616285	-346138	STK39	8.6×10 ⁻⁷	0.083	B-Cell	22446964	
4p16.3	873286	rs35591465	2.8×10 ⁻¹⁵	65801	0.677	rs6599388	0	IDUA	8.6×10 ⁻⁸	0.342	Esophagus_Muscularis	23715323	
14q22.2	55348869	rs11158026	8.8×10 ⁻¹⁵	3994	0.826	rs10136545	0	SAMD4A	3.6×10 ⁻⁶	0.188	Whole_Blood	23715323	
3q27.1	182730291	rs9858038	3.0×10 ⁻¹⁴	-3194	0.600	rs9841498	820265	PARL	0.00031	0.046	Monocyte	22446964	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	-4560	0.988	rs7776649	0	KLHL7-AS1	2.8×10 ⁻⁸	0.261	Muscle_Skeletal	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	-2573	0.915	rs6953728	0	KLHL7-AS1	5.5×10 ⁻²⁴	0.684	Adipose_Subcutaneous	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	16063	0.971	rs10235786	0	KLHL7-AS1	1.6×10 ⁻¹³	0.528	Esophagus_Mucosa	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	19236	0.971	rs10233039	2209	KLHL7-AS1	6.4×10 ⁻¹⁰⁰	0.705	Lymphoblastoid	24037378	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	19236	0.971	rs10233039	0	KLHL7-AS1	9.3×10 ⁻²³	0.651	Skin_Sun_Exposed_Lower_leg	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	19236	0.971	rs10233039	0	KLHL7-AS1	2.4×10 ⁻¹⁸	0.534	Artery_Tibial	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	21418	0.730	rs10256996	107	KLHL7	4.6×10 ⁻⁶	0.016	Monocyte	20502693	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	21418	0.730	rs10256996	0	KLHL7-AS1	5.0×10 ⁻⁸	0.369	Heart_Left_Ventricle	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	21960	0.618	rs1005786	0	AC005082.12	3.9×10 ⁻⁸	0.261	Lung	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	28776	0.947	rs73272048	0	KLHL7-AS1	3.7×10 ⁻¹⁷	0.508	Lung	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	29998	0.947	rs764533	0	NUPL2	1.8×10 ⁻⁶	0.249	Nerve_Tibial	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	46899	0.941	rs4140959	50710	NUPL2	3.7×10 ⁻⁴⁴	0.126	Monocyte	20502693	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	50889	0.935	rs6948469	0	NUPL2	5.9×10 ⁻⁹	0.409	Heart_Left_Ventricle	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	64232	0.877	rs6962213	0	NUPL2	3.0×10 ⁻⁶	0.196	Lung	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	70728	0.787	rs955187	45526	NUPL2	1.4×10 ⁻⁹	0.123	B-Cell	22446964	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	70728	0.787	rs955187	45526	NUPL2	6.1×10 ⁻¹⁹	0.246	Monocyte	22446964	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	87475	0.883	rs6964665	0	KLHL7-AS1	1.2×10 ⁻²¹	0.640	Thyroid	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	96358	0.832	rs10254544	0	KLHL7-AS1	2.1×10 ⁻⁹	0.489	Stomach	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	96573	0.832	rs10242104	0	KLHL7-AS1	7.7×10 ⁻¹⁸	0.660	Esophagus_Muscularis	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	97993	0.758	rs7808488	0	AC005082.12	4.9×10 ⁻⁷	0.294	Esophagus_Mucosa	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	119433	0.634	rs858272	9530	GPNMB	2.3×10 ⁻¹⁷	0.412	Brain_Frontal_Cortex	20485568	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	126951	0.780	rs6970694	0	KLHL7-AS1	2.1×10 ⁻²⁰	0.659	Nerve_Tibial	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	136711	0.670	rs7807879	0	KLHL7-AS1	1.2×10 ⁻⁶	0.204	Whole_Blood	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	138558	0.629	rs1624451	0	AC005082.12	9.2×10 ⁻¹¹	0.458	Esophagus_Muscularis	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	148267	0.658	rs10227559	0	NUPL2	2.4×10 ⁻⁸	0.263	Muscle_Skeletal	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	171023	0.625	rs858273	0	AC005082.12	3.0×10 ⁻¹⁰	0.459	Heart_Left_Ventricle	23715323	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	172656	0.610	rs199355	17487	GPNMB	1.6×10 ⁻⁶	0.079	Monocyte	22446964	
7p15.3	23123877	rs10256359	2.1×10 ⁻¹²	172656	0.610	rs199355	54781	RAB7L1	6.6×10 ⁻¹¹	0.109	Lymphoblastoid	24037378	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	-33765	0.570	rs823094	0	SLC41A1	3.1×10 ⁻⁶	0.259	Esophagus_Mucosa	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	-33765	0.570	rs823094	0	SLC41A1	3.9×10 ⁻⁶	0.187	Whole_Blood	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	-4040	0.995	rs823114	-32107	NUCKS1	7.1×10 ⁻⁵	0.195	T-cell	19644074	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	-4040	0.995	rs823114	-32107	NUCKS1	2.3×10 ⁻¹⁶	0.215	B-Cell	22446964	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	0	1.000	rs823118	0	NUCKS1	1.3×10 ⁻⁹	0.337	Thyroid	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	730	0.610	rs823121	5005	NUCKS1	1.2×10 ⁻²⁹	0.086	Monocyte	20502693	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	12040	0.568	rs2014202	0	RAB7L1	3.1×10 ⁻⁸	0.255	Whole_Blood	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	12319	0.568	rs7522056	2507	RAB7L1	6.1×10 ⁻⁸	0.100	Monocyte	22446964	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	12319	0.568	rs7522056	0	RAB7L1	6.1×10 ⁻⁹	0.346	Nerve_Tibial	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	12319	0.568	rs7522056	0	RAB7L1	1.8×10 ⁻⁹	0.327	Skin_Sun_Exposed_Lower_leg	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	12319	0.568	rs7522056	0	RAB7L1	1.4×10 ⁻⁶	0.231	Adipose_Subcutaneous	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	15694	0.989	rs708723	-51841	NUCKS1	2.1×10 ⁻⁹	0.120	Monocyte	22446964	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	34252	0.620	rs1775148	0	RAB7L1	7.5×10 ⁻⁸	0.430	Artery_Aorta	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	34252	0.620	rs1775148	0	RAB7L1	1.9×10 ⁻¹⁰	0.333	Artery_Tibial	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	35623	0.549	rs1772159	0	PM20D1	5.3×10 ⁻¹⁹	0.592	Adipose_Subcutaneous	23715323	
1q32.1	205723572	rs823118	6.3×10 ⁻¹²	38834	0.552	rs823154	-3940	SLC41A1	3.3×10 ⁻⁵	0.0			

6p22.1	27676794	rs4713118		243000	0.608	rs200997	480698	ZNF323	0.214	Brain Temporal Cortex	20485568
6p22.1	27676794	rs4713118	2.0×10^{-10}	303426	0.588	rs149970	254616	ZNF187	4.4×10^{-6}	Monocyte	20502693
6p22.1	27676794	rs4713118	2.0×10^{-10}	314213	0.533	rs149979	0	ZKSCAN8	8.6×10^{-6}	Muscle_Skeletal	23715323
6p22.1	27676794	rs4713118	2.0×10^{-10}	356380	0.508	rs9295755	0	OR2B8P	1.3×10^{-7}	Skin_Sun_Exposed_Lower_leg	23715323
11q12.1	57694260	rs687432	2.0×10^{-9}	36792	0.968	rs7102929	-2151688	OR5L1	2.5×10^{-5}	Monocyte	22446964
11q12.1	57694260	rs687432	2.0×10^{-9}	39828	0.815	rs12804093	-414908	UBE2L6	0.00040	B-Cell	22446964
16p12.3	19279464	rs11343	1.0×10^{-8}	-652	0.774	rs12445464	258	SYT17	9.5×10^{-9}	Monocyte	20502693
16p12.3	19279464	rs11343	1.0×10^{-8}	-84	1.000	rs1051547	-1121	SYT17	1.3×10^{-5}	B-Cell	22446964
5q12.1	60260543	rs162227	1.4×10^{-8}	-171107	0.733	rs921898	0	ELOVL7	3.8×10^{-6}	Nerve_Tibial	23715323

Nearby Clinical Variants

source	region	our.name	our.pval	dist	rsqr	assay.name	gene	phenotype	accession
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Parkinson's disease	SNOMED CT49049000
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Lewy body dementia	Office of Rare Diseases3243
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Gaucher disease, perinatal lethal	NCBI curation
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Gaucher's disease, type 1	SNOMED CT62201009
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Subacute neuronopathic Gaucher's disease	SNOMED CT5963005
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Acute neuronopathic Gaucher's disease	SNOMED CT12246008
clinvar	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA	Gaucher disease type 3C	NCBI curation
omim	1q22	rs2230288	4.7×10^{-17}	0	1.000	rs2230288	GBA		
clinvar	3q27.1	rs9858038	3.0×10^{-14}	24918	0.708	rs2270968	MCCC1	3 Methylcrotonyl-CoA carboxylase 1 deficiency	Office of Rare Diseases5665

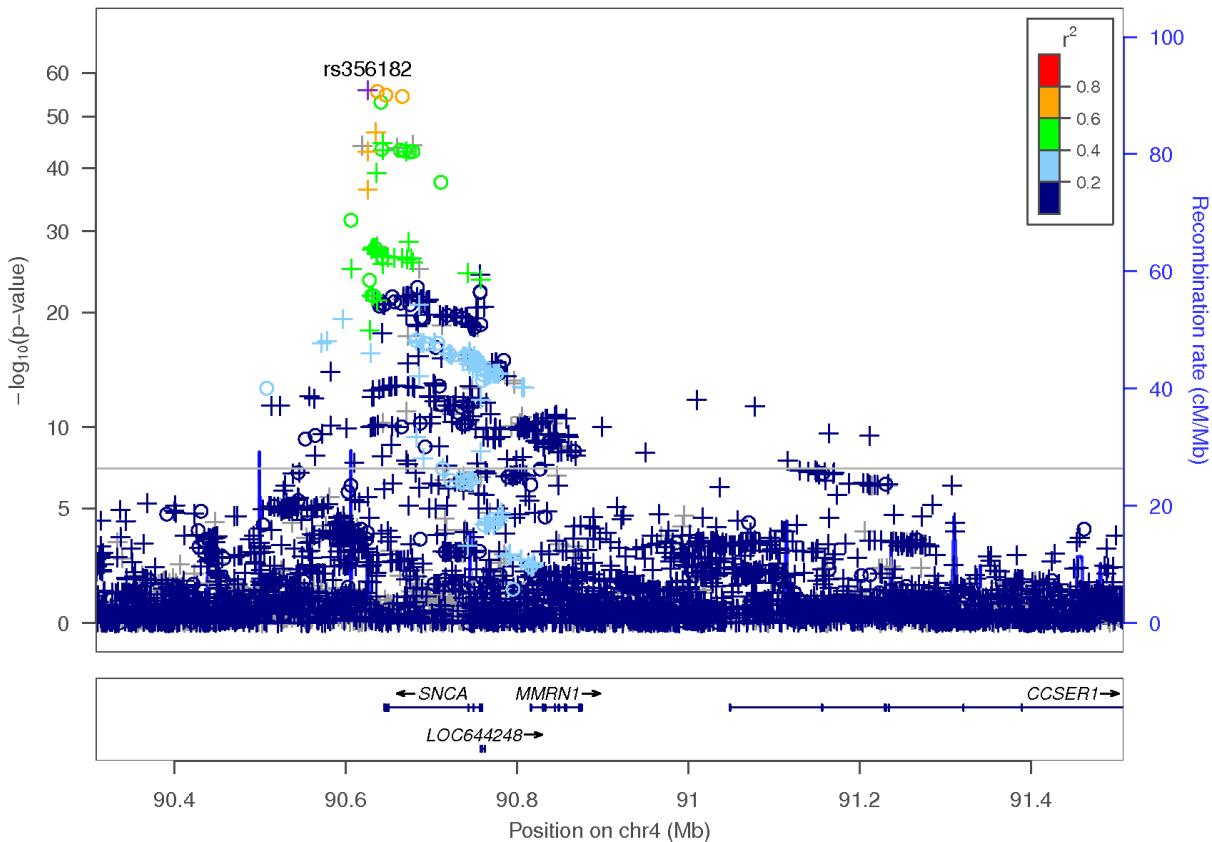
Shared Associations with Other 23andMe Phenotypes

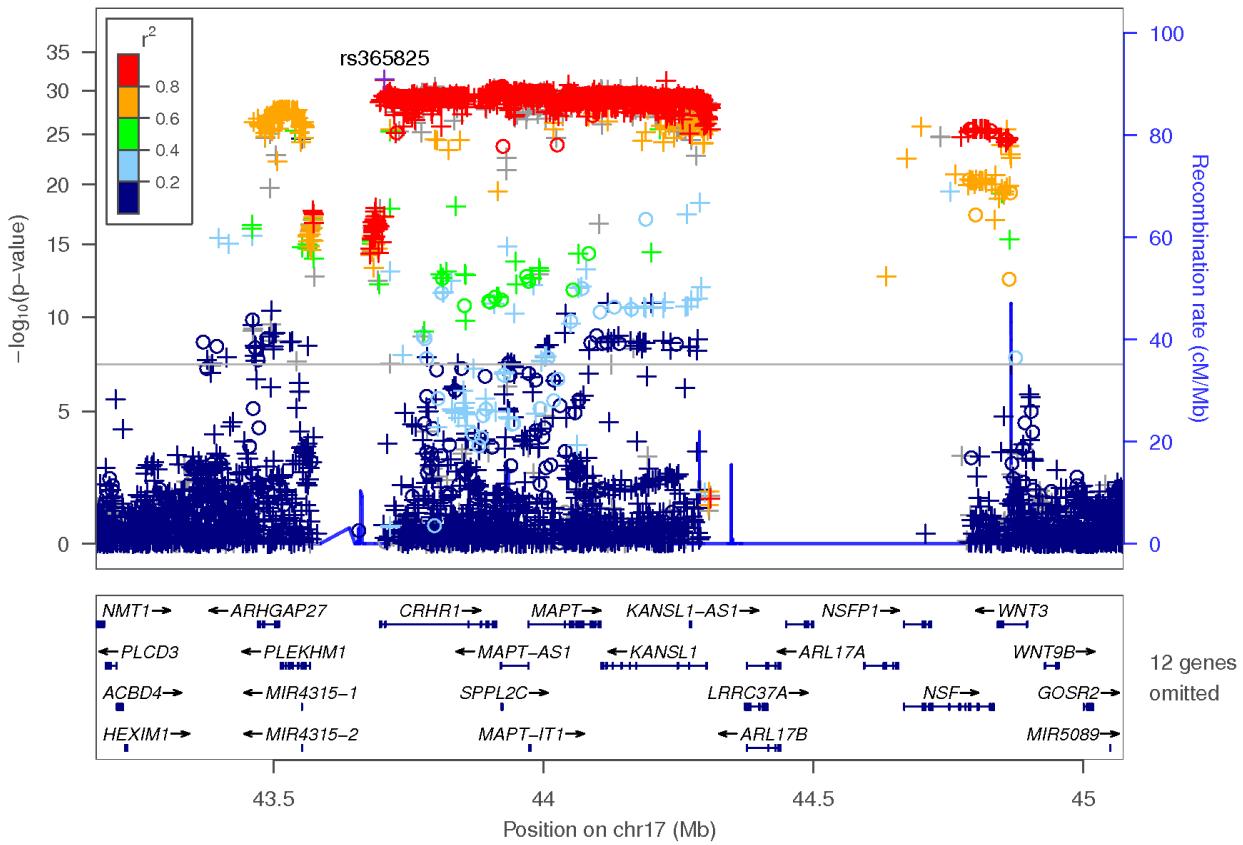
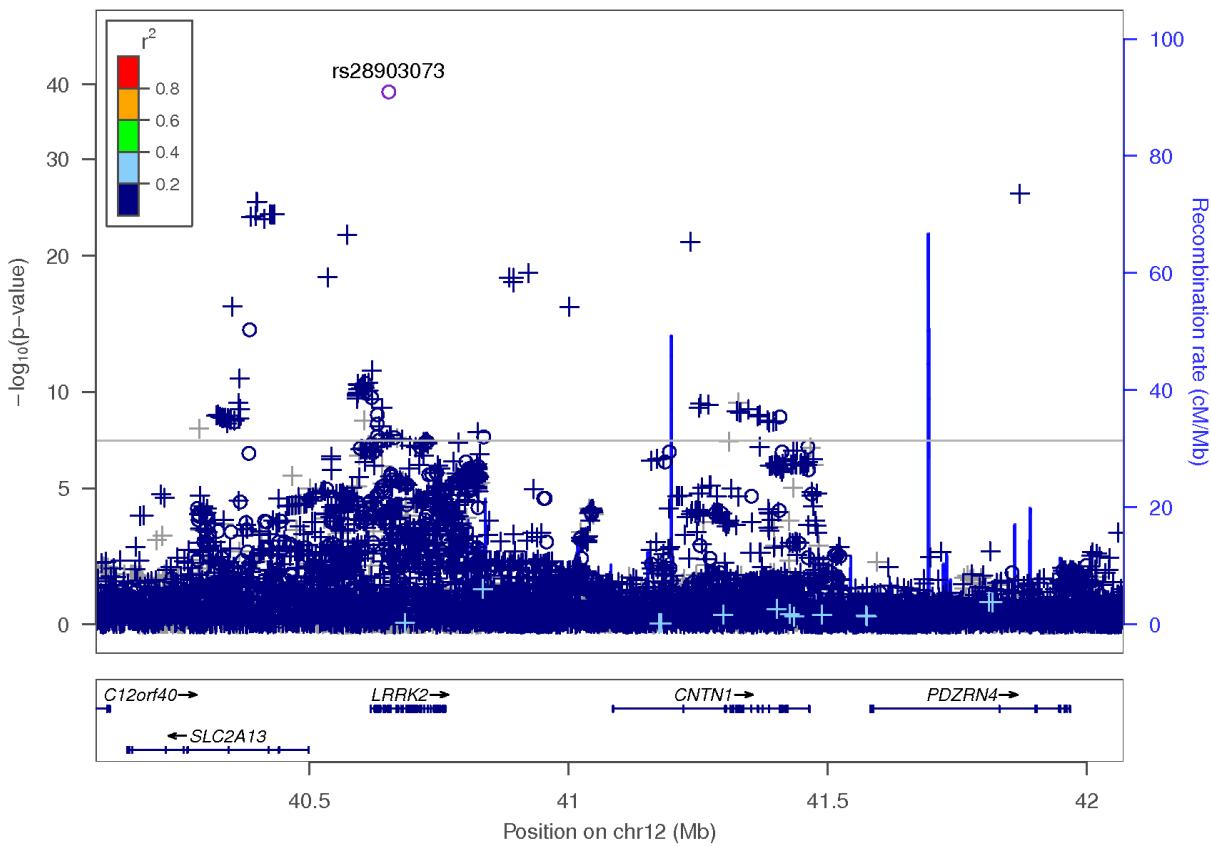
region	position	our.name	our.pval	dist	rsqr	assay.name	phenotype	pvalue	gene.context	match
1p36.23	8445360	rs3765971	1.2×10^{-6}	14887	0.872	rs138050288	any_animal_allergy	1.6×10^{-10}	[RERE]	
1p36.23	8445360	rs3765971	1.2×10^{-6}	14887	0.872	rs138050288	any_plant_allergy	5.6×10^{-12}	[RERE]	
1p36.23	8445360	rs3765971	1.2×10^{-6}	14887	0.872	rs138050288	dust_mites_allergy	1.2×10^{-8}	[RERE]	
1p36.11	25806112	rs35065124	1.4×10^{-5}	-75195	0.527	rs642255	hair_curl'	5.9×10^{-7}	[RHCE]	TRUE
1p36.11	25736369	rs111941366	1.1×10^{-5}	0	1.000	rs111941366	iqb.blood_donation_frequency	2.1×10^{-20}	[RHCE]	TRUE
1p36.11	25736369	rs111941366	1.1×10^{-5}	17656	0.941	rs909832	iqb.bruise_easily	5.7×10^{-14}	RHCE-[]-TMEM57	FALSE
1p36.11	25736369	rs111941366	1.1×10^{-5}	89580	0.566	rs200391466	high_cholesterol_broad	2.8×10^{-7}	[TMEM57]	
3p21.31	48327563	rs34618864	7.3×10^{-5}	19392	0.787	rs34644650	blood_pressure_meds	4.0×10^{-9}	NME6-[]-SPINK8	
3p21.31	49013383	rs6768433	2.6×10^{-7}	-28876	0.831	rs200143221	iqb.carry_a_tune	4.5×10^{-8}	[ARIH2]	
3p21.31	49013383	rs6768433	2.6×10^{-7}	-28876	0.831	rs200143221	motionsick_combined	3.8×10^{-8}	[ARIH2]	
3p21.1	53094194	rs11712910	4.4×10^{-5}	-59150	0.874	rs2581792	hemorrhoids	6.4×10^{-9}	[SFMBT1]	FALSE
3q13.32	118291124	rs6438460	1.6×10^{-5}	37631	0.509	rs6799936	motionsick_combined	5.6×10^{-9}	[]---IGSF11	FALSE
3q25.1	151111173	rs12498042	7.1×10^{-7}	19168	0.653	rs10935845	iqb.sound_sleeper	3.6×10^{-7}	[MED12L]	TRUE
3q27.1	182811490	rs3772722	1.6×10^{-9}	38074	0.546	rs3772715	iqb.fruit_juice_frequency	6.6×10^{-7}	[LAMP3]	FALSE
4p16.3	976598	rs17783233	3.3×10^{-12}	-24651	0.534	rs34311866	pd_family_history	7.1×10^{-7}	[TMEM175]	TRUE
4p16.3	1841330	rs115564335	3.3×10^{-5}	-47050	0.642	rs10012764	linebacker	3.6×10^{-7}	TACC3-[]-FGFR3	TRUE
4p15.31	17853921	rs16895888	1.3×10^{-5}	20168	0.784	rs7671110	height	2.1×10^{-54}	NCAPG-[]-LCORL	TRUE
4p15.31	17853921	rs16895888	1.3×10^{-5}	77397	0.775	rs13146142	shoe_size_normalized	1.9×10^{-17}	[LCORL]	TRUE
4p15.31	17940197	rs71603391	6.4×10^{-5}	89639	0.829	rs112093994	weight	6.3×10^{-10}	LCORL-[]	
4q21.1	77336203	rs13104972	2.8×10^{-7}	17827	0.568	rs59136511	gallstones_or_gall_bladder_removal	1.6×10^{-7}	SHROOM3	TRUE
4q22.1	90626111	rs356182	1.2×10^{-56}	15229	0.520	rs356220	any_neurological	5.4×10^{-12}	GPRIN3-[]-SNCA	TRUE
4q22.1	90626111	rs356182	1.2×10^{-56}	15229	0.520	rs356220	pd_family_history	4.6×10^{-9}	GPRIN3-[]-SNCA	TRUE
4q22.1	90711770	rs2737029	2.4×10^{-38}	32478	0.508	rs2619344	updrs_constipation	9.0×10^{-7}	[SNCA]	FALSE
4q22.1	90781538	rs10025824	1.7×10^{-5}	34485	0.576	rs1442140	panic	6.0×10^{-7}	SNCA-[]-MMRN1	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	-41821	0.820	rs35225200	iqb.rotator_cuff_tear	4.3×10^{-7}	SLC39A8	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	iqb.excess_hair	3.8×10^{-8}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	any_animal_allergy	2.3×10^{-7}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	bmi	7.7×10^{-30}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	high_plasma_glucose	9.0×10^{-7}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	iqb.low_hdl	3.6×10^{-14}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	obesity	3.0×10^{-9}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	weight	2.3×10^{-17}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	height	6.0×10^{-11}	[SLC39A8]	TRUE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	iqb.arthritis_any	2.2×10^{-7}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	0	1.000	rs13107325	osteoarthritis_broad	1.3×10^{-8}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	9373	0.931	rs13135092	iqb.stretch_marks	3.0×10^{-10}	[SLC39A8]	FALSE
4q24	103188709	rs13107325	8.2×10^{-7}	9373	0.931	rs13135092	stretch_marks_any	4.1×10^{-9}	[SLC39A8]	FALSE
5q14.3	87721684	rs35182117	3.7×10^{-6}	-75557	0.666	rs71639107	any_psychiatric	2.1×10^{-9}	MEF2C	
5q14.3	87721684	rs35182117	3.7×10^{-6}	-75557	0.666	rs71639107	depression	6.1×10^{-9}	MEF2C	
5q14.3	87721684	rs35182117	3.7×10^{-6}	-71805	0.641	rs113828108	morning_person	2.4×10^{-10}	MEF2C	
5q31.1	134335383	rs299366	4.2×10^{-5}	37302	0.606	rs31198	height	3.4×10^{-17}	[LOC100996485]	TRUE
6p22.1	27676794	rs4713118	2.0×10^{-10}	17245	0.839	rs67869107	any_immune	5.8×10^{-7}	HIST1H2BL	
6p22.1	27850873	rs494887	5.7×10^{-6}	-75199	0.511	rs200484	glasses_any	7.4×10^{-8}	[HIST1H2BL]	TRUE
6p22.1	27690453	rs9468206	9.7×10^{-9}	97609	0.869	rs200502	urine_control	2.1×10^{-7}	HIST1H4J	FALSE
6p22.1	27968350	rs78094982	6.6×10^{-7}	-57135	0.715	rs151313790	any_neurological	7.3×10^{-7}	OR2B2-[]-OR2B6	TRUE
6p22.1	28595107	rs11970409	4.5×10^{-6}	-16821	0.651	rs34388845	bmi	6.8×10^{-10}	TRIM27	TRUE
6p21.33	31184877	rs9263960	1.2×10^{-5}	43533	0.601	rs1986997	weight	6.3×10^{-15}	HCG27-[]-HLA-C	TRUE
6p21.32	32398853	rs377537858	2.4×10^{-6}	10203	0.596	rs9268652	hepatitis_b	3.1×10^{-9}	[HLA-DRA]	
6p21.32	32427748	rs9268831	1.0×10^{-5}	-4511	0.522	rs9268803	iqb.poison_oak	5.3×10^{-8}	DRB5	FALSE
6p21.32	32406843	rs3763323	1.8×10^{-5}	22739	0.853	rs9268851	multiple_sclerosis	5.3×10^{-85}	HLA-DRA-[]-HLA-	TRUE
6p21.32	32572755	rs9270936	7.3×10^{-6}	-6178	0.773	rs4329147	iqb.childhood_ear_infections	9.5×10^{-12}	HLA-DQA1	FALSE
6p21.32	32572904	rs574170	2.8×10^{-9}	-3742	0.549	rs28752504	depression	2.8×10^{-7}	HLA-DQA1	FALSE

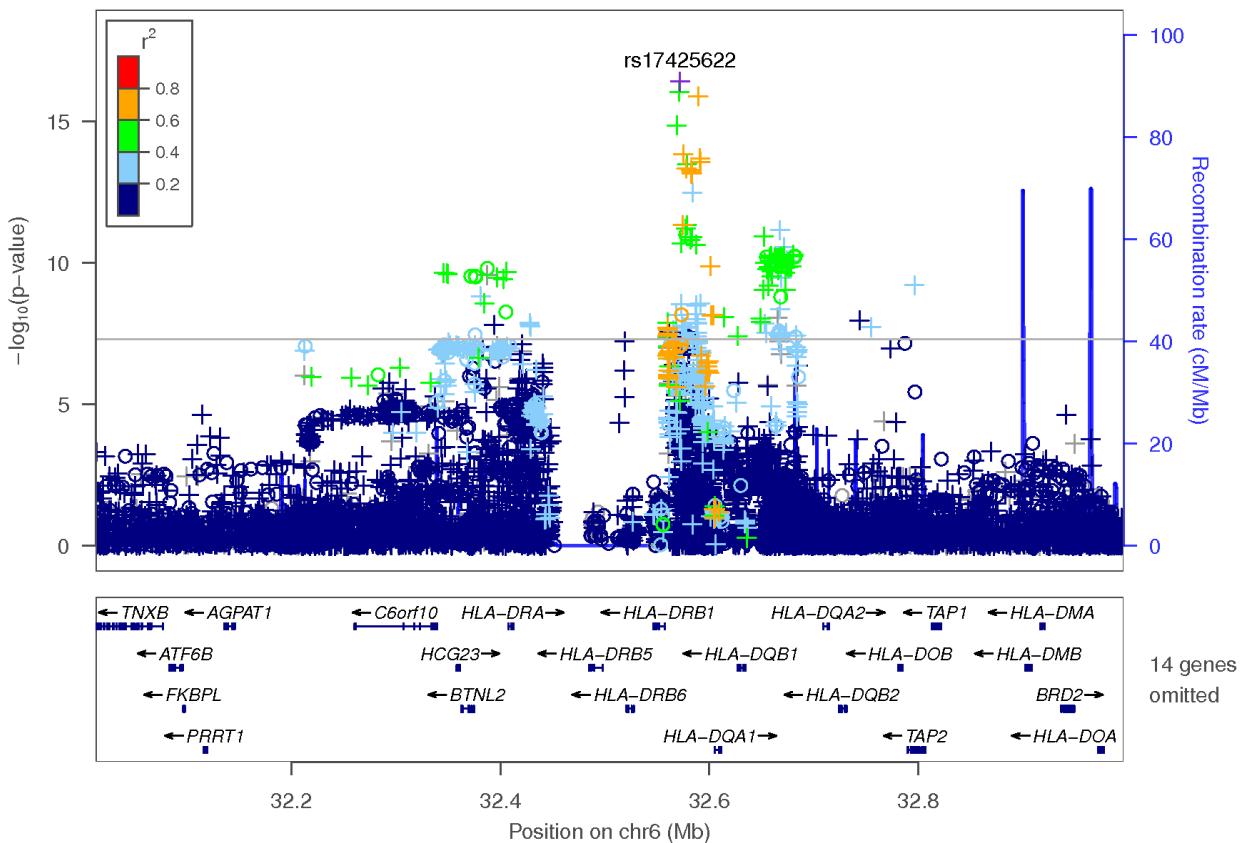
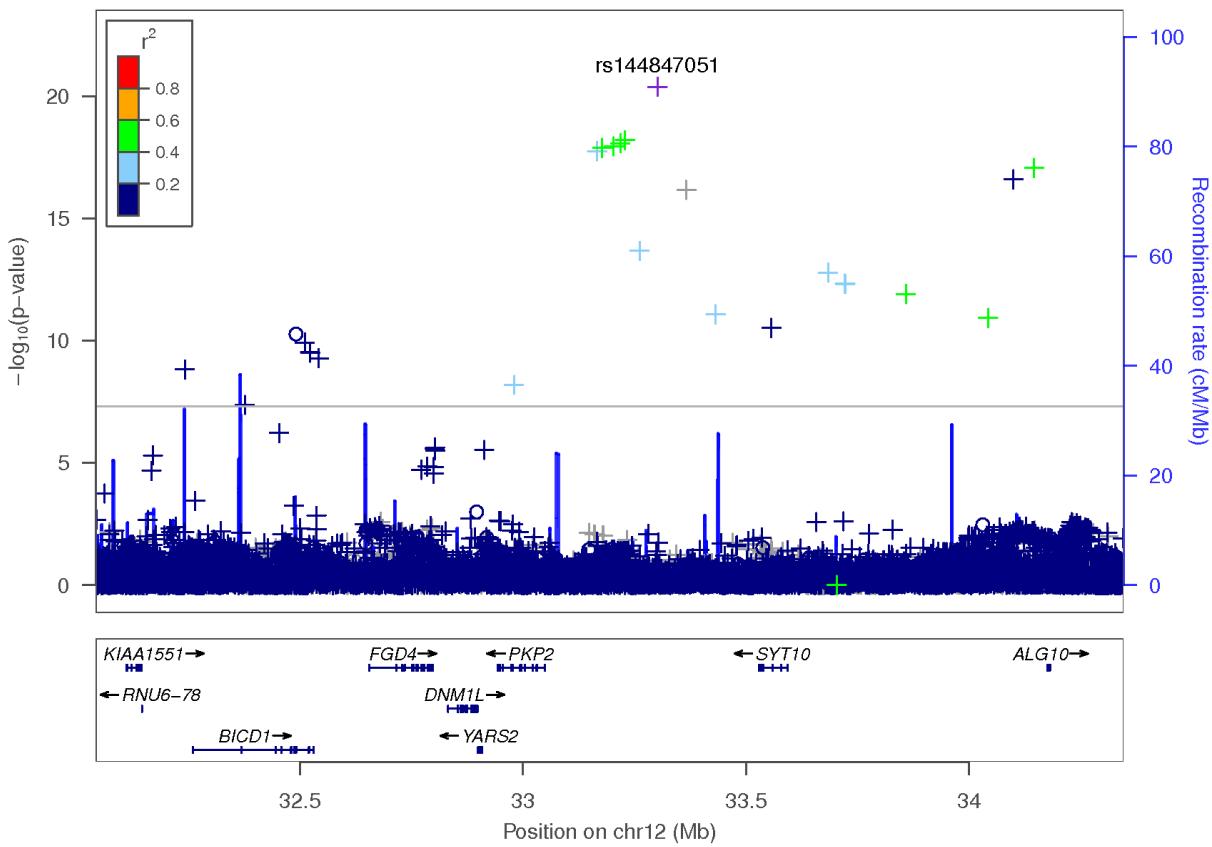
6p21.32	32571961	rs17425622	3.8×10^{-17}	1304	0.523	rs601148	iqb.took_meds_prednisone_or_steroid	6.7×10^{-8}	HLA-DRB5--[]-- HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32582577	rs3104415	4.2×10^{-6}	35	0.927	rs3104414	rheumatoid_arthritis	2.8×10^{-39}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32583603	rs28752515	2.0×10^{-5}	716	0.608	rs28383241	heart_metabolic_disease	2.2×10^{-8}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32583813	rs1846190	7.6×10^{-8}	3200	0.681	rs557011	vitiligo	6.2×10^{-26}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32587416	rs9271386	1.6×10^{-5}	-7	0.752	rs9271385	juvenile_asthma	3.2×10^{-11}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32591332	rs9271600	2.7×10^{-9}	-2909	0.510	rs9271455	any_psychiatric	7.3×10^{-8}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32591332	rs9271600	2.7×10^{-9}	-2778	0.517	rs9271465	iqb.bronchitis_frequency	1.6×10^{-15}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32571961	rs17425622	3.8×10^{-17}	17454	0.658	rs9271509	blood_sugar_meds	2.1×10^{-14}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32591332	rs9271600	2.7×10^{-9}	5	0.989	rs9271601	any_skin_cancer	7.6×10^{-10}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	TRUE
6p21.32	32587553	rs9271404	3.1×10^{-9}	3830	0.536	rs9271602	milk_allergy	7.0×10^{-7}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32572755	rs9270936	7.3×10^{-6}	19167	0.617	rs4959108	high_cholesterol_broad	8.4×10^{-18}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32602872	rs9272245	4.7×10^{-5}	-3801	0.703	rs9272050	plantar_warts	9.6×10^{-31}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32602872	rs9272245	4.7×10^{-5}	-476	0.958	rs9272226	iqb.plantar_warts	1.8×10^{-40}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32566149	rs9270664	1.3×10^{-6}	37338	0.513	rs3104367	allergic_asthma	3.4×10^{-13}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32566149	rs9270664	1.3×10^{-6}	37338	0.513	rs3104367	any_asthma	1.0×10^{-40}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32566149	rs9270664	1.3×10^{-6}	37338	0.513	rs3104367	iqb.took_meds_asthma_or_lung	5.3×10^{-39}	HLA-DQA1 HLA-DRB5--[]-- HLA-DQA1	FALSE
6p21.32	32571961	rs17425622	3.8×10^{-17}	31837	0.629	rs1391371	iqb.acne_as_teensager	1.2×10^{-9}	HLA-DQA1 HLA-DQA1--[]--HLA- DQB1	TRUE
6p21.32	32626021	rs6905775	8.7×10^{-6}	251	0.534	rs9273363	t1d	1.1×10^{-201}	HLA-DQA1--[]--HLA- DQB1	FALSE
6p21.32	32569659	rs9270810	3.6×10^{-7}	56833	0.530	rs9273370	hypothyroidism	4.5×10^{-34}	DQB1 HLA-DQB1	FALSE
6p21.32	32636757	rs3134998	5.9×10^{-5}	-8790	0.595	rs1063346	juvenile_allergic_asthma	6.3×10^{-13}	[HLA-DQB1] HLA-DQB1--[]-- HLA-DQB1	FALSE
6p21.32	32663986	rs6457619	4.1×10^{-5}	-10260	0.520	rs9275170	acid_reflux_meds	6.8×10^{-10}	HLA-DQA2 HLA-DQB1--[]-- HLA-DQB1	FALSE
6p21.32	32667638	rs9275348	4.2×10^{-9}	462	0.508	rs2647045	any_non_melanoma_skin_cancer	4.9×10^{-10}	HLA-DQA2 HLA-DQB1--[]-- HLA-DQB1	TRUE
6p21.32	32658933	rs9275218	1.1×10^{-5}	14821	0.539	rs28772279	lightheaded	3.7×10^{-7}	HLA-DQA2 HLA-DQB1--[]-- HLA-DQB1	FALSE
6p21.32	32587117	rs28383313	9.7×10^{-9}	90795	0.505	rs9275563	iqb.oral_corticosteroid_ever	3.5×10^{-7}	HLA-DQA2 HLA-DQB1--[]-- HLA-DQB1	FALSE
6p21.32	32587117	rs28383313	9.7×10^{-9}	90795	0.505	rs9275563	iqb.pneumonia_frequency	1.4×10^{-11}	HLA-DQA2 HLA-DQB1--[]-- HLA-DQB1	FALSE
6p21.32	32684935	rs9275642	6.5×10^{-5}	0	1.000	rs9275642	basal_cell_carcinoma	7.7×10^{-11}	HLA-DQA2 HLA-DQB1	TRUE
6q21	112240786	rs12661198	9.3×10^{-5}	-55022	0.618	rs139971631	mpn_or_v617f	2.0×10^{-7}	FYN---[]--WISP3	
7p15.3	23586607	rs111254487	7.2×10^{-5}	-92994	0.665	rs112316332	iqb.math_class	4.4×10^{-7}	[IGFBP3]	FALSE
9p22.2	16718436	rs10756788	2.9×10^{-6}	17727	0.926	rs13284478	hair_curl	5.0×10^{-8}	[BNC2] MUC19--[]-- MUC19	TRUE
12q12	40824058	rs149557347	7.7×10^{-7}	-183	0.932	rs189621487	iqb.rotator_cuff_tear	2.9×10^{-7}	MUC19 MUC19	FALSE
12q12	41870804	rs187043470	7.6×10^{-27}	0	1.000	rs187043470	any_neurological	8.4×10^{-8}	[PDZRN4]	TRUE
12q12	41870804	rs187043470	7.6×10^{-27}	0	1.000	rs187043470	pd_family_history	6.7×10^{-16}	[PDZRN4]	TRUE
16p11.2	28620572	rs750155	2.1×10^{-5}	-81186	0.543	rs189278832	iqb.stretch_marks	7.3×10^{-11}	[NPIPL1]	FALSE
16p11.2	28620572	rs750155	2.1×10^{-5}	-81186	0.543	rs189278832	stretch_marks_any	7.8×10^{-11}	[NPIPL1]	FALSE
16p11.2	28564146	rs11364090	2.1×10^{-5}	-18559	0.558	rs231976	mosquito_severe	1.7×10^{-7}	[NPIPL1]	
16p11.2	28620572	rs750155	2.1×10^{-5}	-13040	0.512	rs4115668	any_allergy	5.9×10^{-8}	[NPIPL1,SULT1A2]	TRUE
16p11.2	28620572	rs750155	2.1×10^{-5}	-13040	0.512	rs4115668	any_plant_allergy	3.8×10^{-8}	[NPIPL1,SULT1A2]	TRUE
16p11.2	28620572	rs750155	2.1×10^{-5}	0	1.000	rs750155	seasonal_allergies_broad	6.1×10^{-8}	[NPIPL1,SULT1A1]	TRUE
16p11.2	28927818	rs12919058	2.3×10^{-5}	-61902	0.675	rs7187333	bmi	2.6×10^{-17}	[NPIPL1]	FALSE
16p11.2	28927818	rs12919058	2.3×10^{-5}	-56618	0.671	rs199766562	weight	9.6×10^{-21}	[NPIPL1]	
16p11.2	28927818	rs12919058	2.3×10^{-5}	-29025	0.760	rs6565259	iqb.ingrown_hair_bumps	1.4×10^{-7}	[NPIPL1,ATP2A1]	FALSE
16p11.2	28927818	rs12919058	2.3×10^{-5}	-8477	0.814	rs7206214	shoe_size_normalized	5.5×10^{-11}	[NPIPL1,RABEP2]	FALSE
16p11.2	30914626	rs1046276	5.6×10^{-6}	-85954	0.602	rs7192475	number_of_cavities	6.3×10^{-7}	ZNF629--[]--BCL7C	FALSE
16p11.2	30980130	rs4346218	2.6×10^{-7}	-93487	0.596	rs897984	iqb.took_meds_prednisone_or_steroid	2.2×10^{-7}	ZNF629--[]--BCL7C	FALSE
16p11.2	31037020	rs11150604	7.1×10^{-6}	-85454	0.795	rs13336347	high_blood_pressure	3.4×10^{-8}	[FBXL19]	
16p11.2	31084394	rs11281612	7.0×10^{-6}	-90571	0.610	rs11150602	heart_metabolic_disease	6.2×10^{-11}	[SETD1A]	
16p11.2	31084394	rs11281612	7.0×10^{-6}	-70215	0.837	rs9926533	heart_metabolic_disease_burden	5.6×10^{-11}	[STX1B]	
16p11.2	31084394	rs11281612	7.0×10^{-6}	-36315	0.884	rs10871454	warfarin_dosage	1.7×10^{-28}	[STX4]	
16p11.2	31084394	rs11281612	7.0×10^{-6}	3953	0.879	rs749671	warfarin_sensitivity	2.3×10^{-9}	[ZNF646]	
16q12.2	52635000	rs12922061	5.9×10^{-6}	-54500	0.619	rs3095602	breast_cancer	1.6×10^{-19}	[TOX3]	TRUE
16q12.2	52635000	rs12922061	5.9×10^{-6}	-54500	0.619	rs3095602	breast_or_ovarian_cancer	1.9×10^{-17}	[TOX3]	TRUE
16q12.2	52644895	rs3104798	2.0×10^{-5}	-3709	0.937	rs3104793	iqb.restless_leg_syndrome	3.3×10^{-15}	TOX3--[]--CHD9	FALSE
17p11.2	17715101	rs11868035	1.6×10^{-5}	-18346	0.936	rs11649804	myringotomy	2.9×10^{-7}	[RAI1]	TRUE
17p11.2	17853454	rs80777530	3.5×10^{-5}	-91207	0.823	rs11658309	iqb.strep_throat	4.7×10^{-7}	[TOM1L2]	TRUE
17q21.2	40698158	rs601999	6.7×10^{-8}	0	1.000	rs601999	body_type	3.5×10^{-7}	NAGLU--[]-HSD17B1 PLEKHM1--[]-	TRUE
17q21.31	43515927	rs35354512	1.7×10^{-28}	56969	0.591	rs5026246	personality_assertiveness	3.1×10^{-8}	ARL17A LRRK37A3--[]-- CRHR1	TRUE
17q21.31	43705601	rs365825	4.2×10^{-32}	88227	0.985	rs4479288	nearsightedness	3.7×10^{-7}	LRRK37A3--[]-- CRHR1	TRUE
17q21.31	43807911	rs192006062	5.0×10^{-29}	26760	0.501	rs71363552	extraversion_combined	9.7×10^{-8}	LRRK37A3--[]-- CRHR1	
17q21.31	43807911	rs192006062	5.0×10^{-29}	26760	0.501	rs71363552	iqb.talk_versus_listen	5.2×10^{-10}	LRRK37A3--[]-- CRHR1	
17q21.31	43849572	17:43849572:AT_A	4.8×10^{-30}	0	1.000	17:43849572:AT_A	unibrow	2.9×10^{-32}	LRRK37A3--[]-- CRHR1	FALSE
17q21.31	43924337	rs12373168	3.0×10^{-31}	-73405	0.912	rs113518470	beighton_hypermobility_qtl	4.7×10^{-11}	LRRK37A3--[]-- CRHR1	TRUE
17q21.31	43924337	rs12373168	3.0×10^{-31}	-67998	0.564	rs117631329	any_neurological	8.0×10^{-10}	CRHR1	TRUE
17q21.31	43924337	rs12373168	3.0×10^{-31}	960	0.790	rs17690703	iqb.forward_splits	1.0×10^{-8}	SPPL2C[]--MAPT	TRUE
17q21.31	43879703	rs72834575	9.1×10^{-5}	50997	0.879	rs62054826	iqb.shaves_legs	3.0×10^{-8}	SPPL2C[]--MAPT	TRUE
17q21.31	43931855	17:43931855:GAGAG	4.4×10^{-22}	0	1.000	17:43931855:GAGAG	iqb.self_rate_attractiveness	4.7×10^{-7}	SPPL2C[]--MAPT	TRUE
17q21.31	44041771	rs2664006	6.0×10^{-11}	-2406	0.928	rs242554	iqb.nail_biting_frequency	2.9×10^{-7}	[MAPT]	FALSE
17q21.31	44083948	rs8067056	4.9×10^{-15}	0	1.000	rs8067056	can_lift_left_eyebrow	2.3×10^{-10}	[MAPT]	TRUE
17q21.31	44046934	rs113925422	1.9×10^{-31}	44952	0.907	rs9891103	lift_one_eyebrow	6.3×10^{-9}	[MAPT]	TRUE
17q21.31	44046934	rs113925422	1.9×10^{-31}	60672	0.905	rs367721874	iqb.pee_middle_night	5.4×10^{-10}	[KANSL1]	

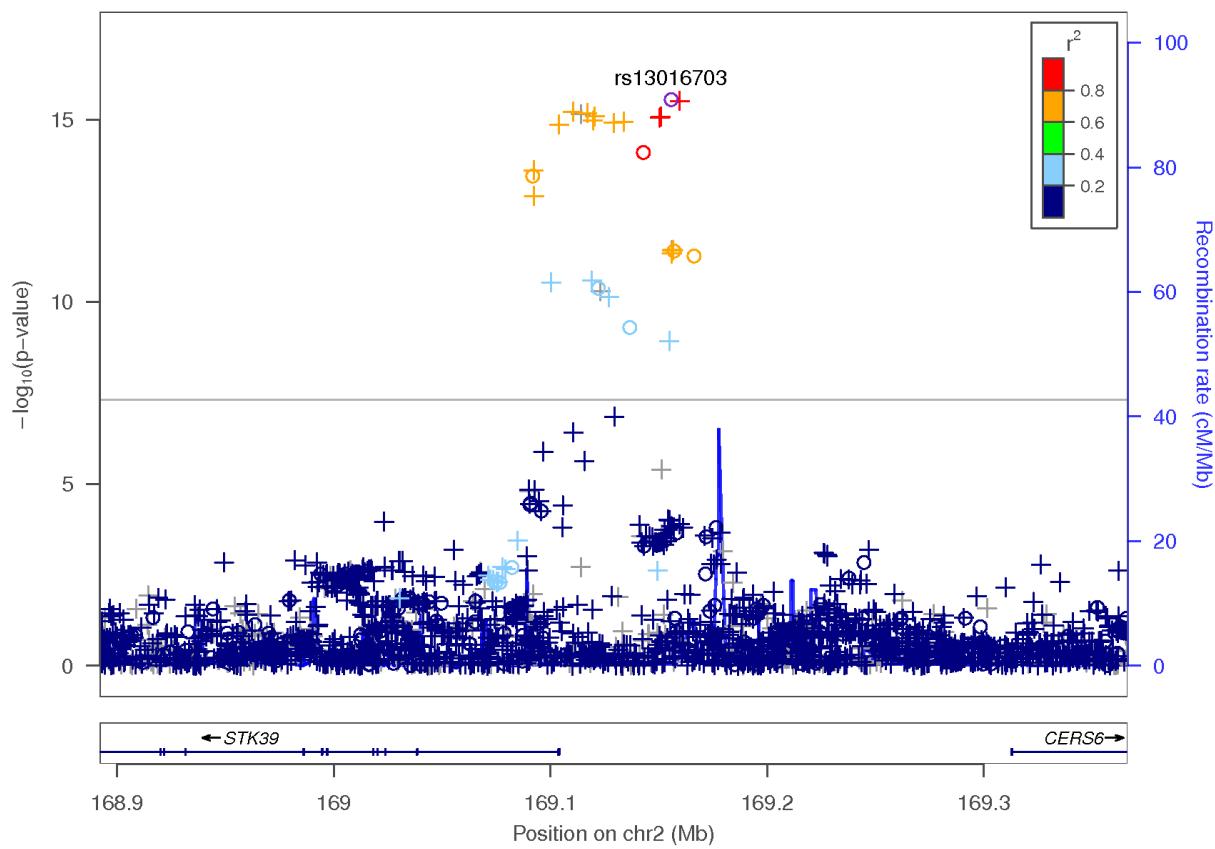
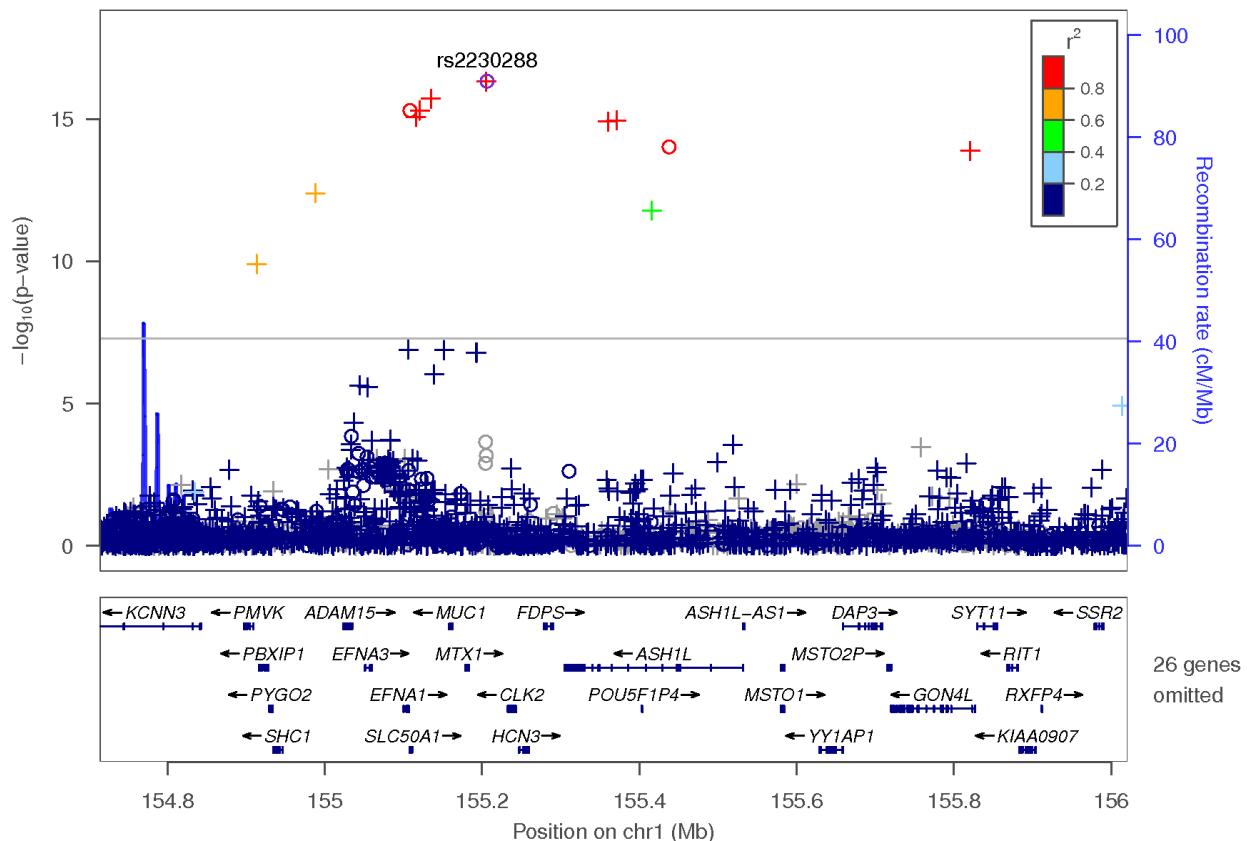
17q21.31	44191483	rs11653367	2.0×10 ⁻⁷	-69904	0.582	rs72836318	heart_metabolic_disease	8.1×10 ⁻⁷	[KANSL1]	FALSE
17q21.31	44191483	rs11653367	2.0×10 ⁻⁷	-69904	0.582	rs72836318	bmi	6.4×10 ⁻⁹	[KANSL1]	FALSE
17q21.31	44191483	rs11653367	2.0×10 ⁻⁷	-69904	0.582	rs72836318	weight	5.1×10 ⁻⁸	[KANSL1]	FALSE
17q21.31	44227393	rs2696579	6.5×10 ⁻³²	-61315	0.613	rs201408539	male_pattern_meta	8.4×10 ⁻²⁰	[KANSL1]	
17q21.31	44227393	rs2696579	6.5×10 ⁻³²	-44410	0.547	rs78381082	height	9.2×10 ⁻⁸	[KANSL1]	TRUE
17q21.31	44289832	rs2696466	4.0×10 ⁻¹⁹	-89640	0.537	rs9908639	iqb.cartwheel	3.4×10 ⁻⁷	[KANSL1]	TRUE
17q21.31	44227393	rs2696579	6.5×10 ⁻³²	55079	0.599	rs2668658	iqb.fresh_cucumbers_yummy	6.8×10 ⁻⁹	LRRC37A KANSL1--[]--	TRUE
17q21.31	44227393	rs2696579	6.5×10 ⁻³²	77806	0.765	rs2260227	bald_image	2.8×10 ⁻¹³	LRRC37A	
17q21.31	44699851	rs116956554	1.5×10 ⁻²⁶	-26900	0.681	rs147847247	nearsightedness	3.7×10 ⁻⁷	[NSF]	TRUE
17q21.31	44699851	rs116956554	1.5×10 ⁻²⁶	0	1.000	rs116956554	hair_loss	5.4×10 ⁻¹⁸	[NSF]	TRUE
17q21.31	44812188	rs199447	2.5×10 ⁻²⁶	-12142	0.661	rs7224296	community_standing	6.0×10 ⁻⁸	[NSF]	TRUE
17q21.31	44812188	rs199447	2.5×10 ⁻²⁶	13868	0.992	rs9896243	vertex	9.5×10 ⁻⁷	[NSF]	TRUE
17q21.31	44812188	rs199447	2.5×10 ⁻²⁶	36250	0.767	rs199524	eyecolor_herc2_blue	1.1×10 ⁻¹⁰	[WNT3]	TRUE
17q24.2	65892280	rs59640213	1.0×10 ⁻⁶	55360	0.658	rs12449442	weight	4.6×10 ⁻⁷	[BPTF]	FALSE
Xq24	119436726	rs5909698	4.4×10 ⁻⁵	2609	0.966	rs5957354	eyecolor_herc2_blue	2.2×10 ⁻¹⁶	[TMEM255A]	TRUE
Xq24	119436726	rs5909698	4.4×10 ⁻⁵	2609	0.966	rs5957354	eyecolor_simple	2.3×10 ⁻¹⁴	[TMEM255A]	TRUE

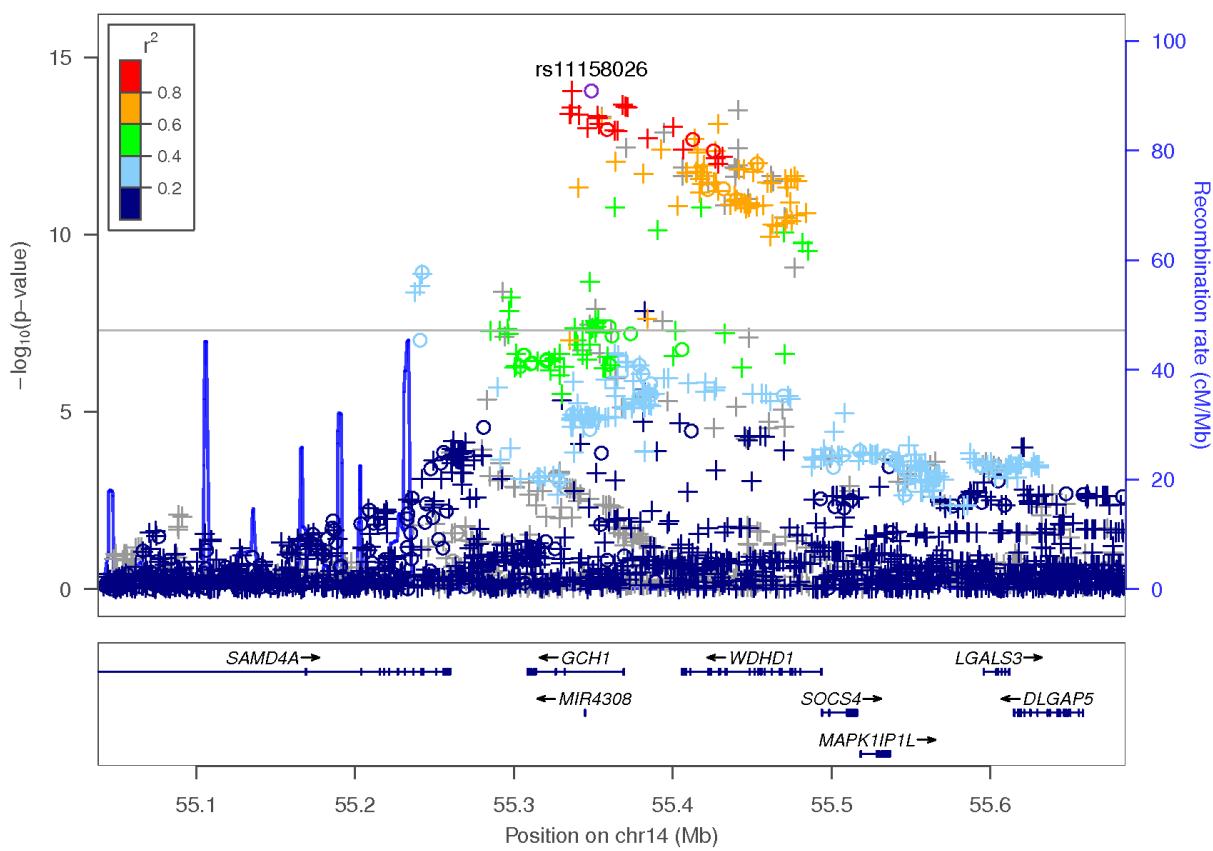
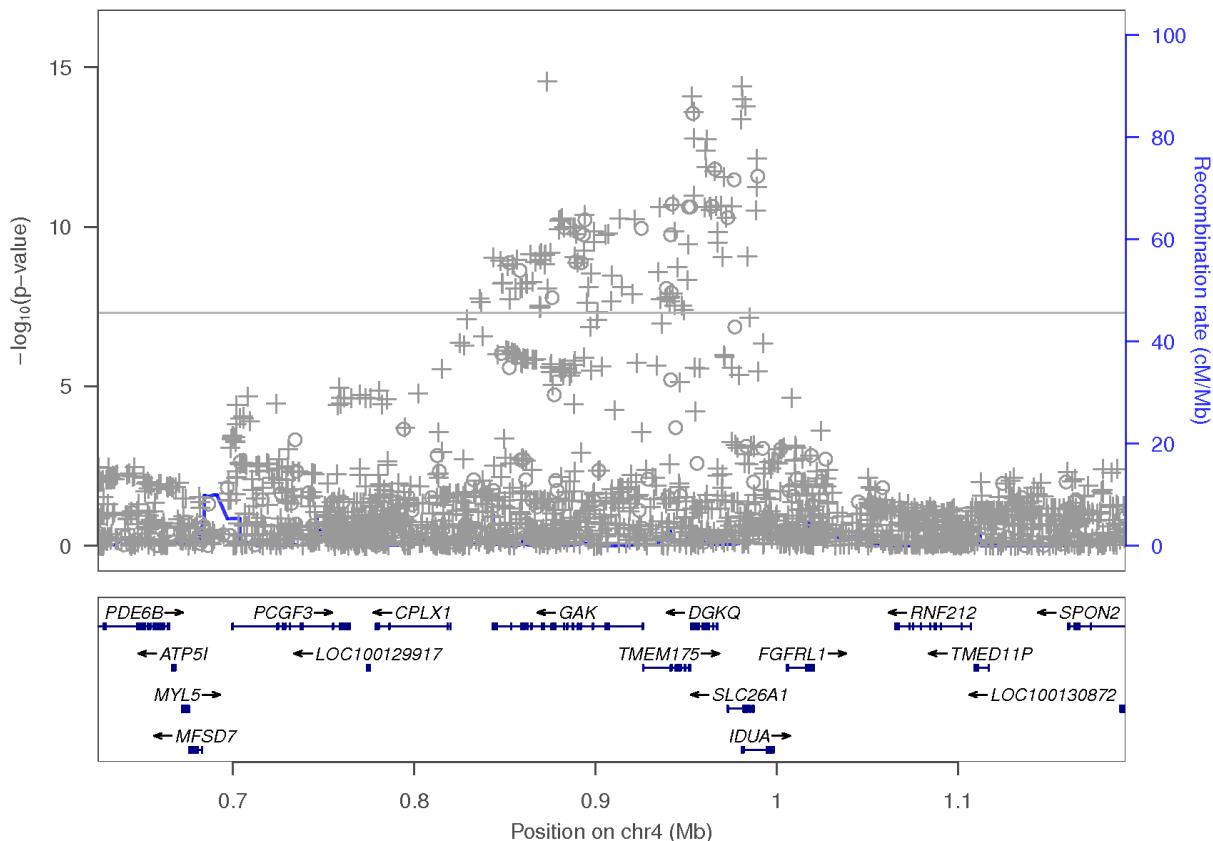
Regional Association Plots

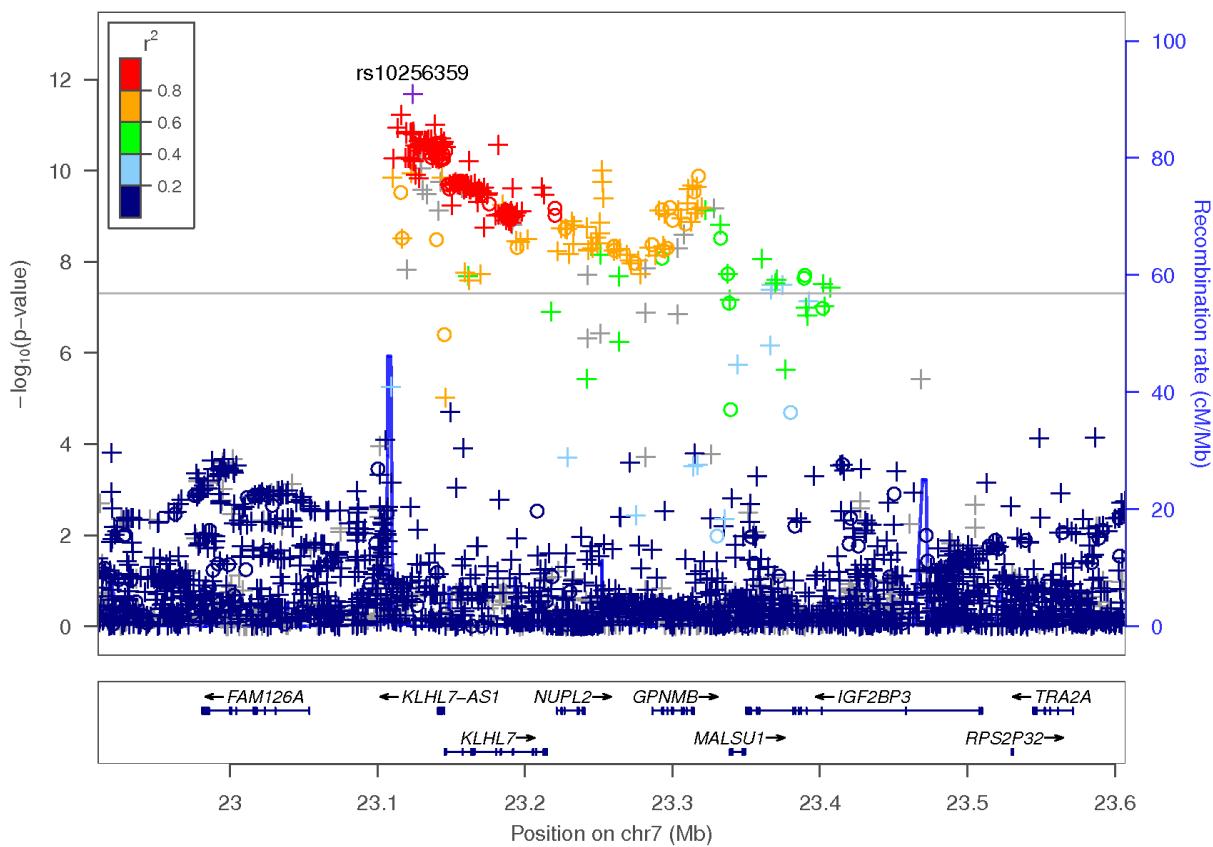
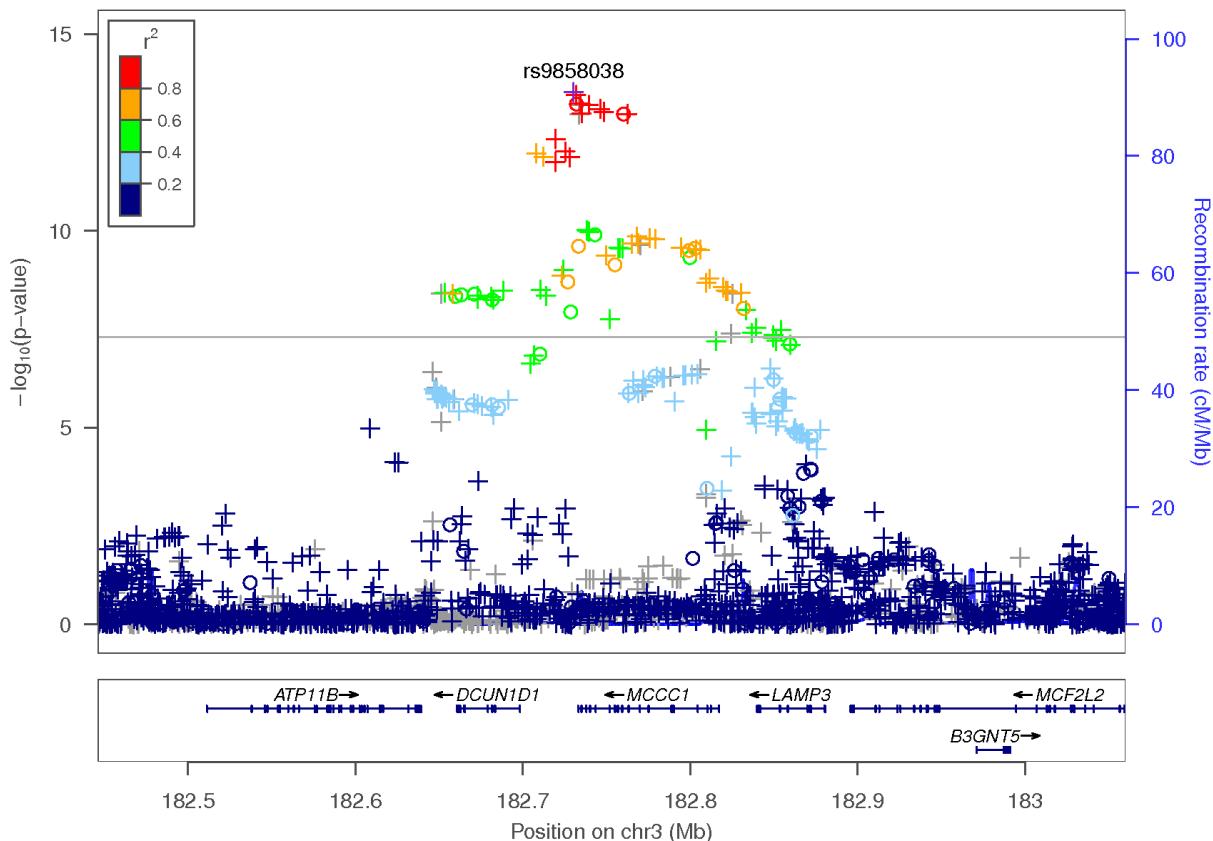


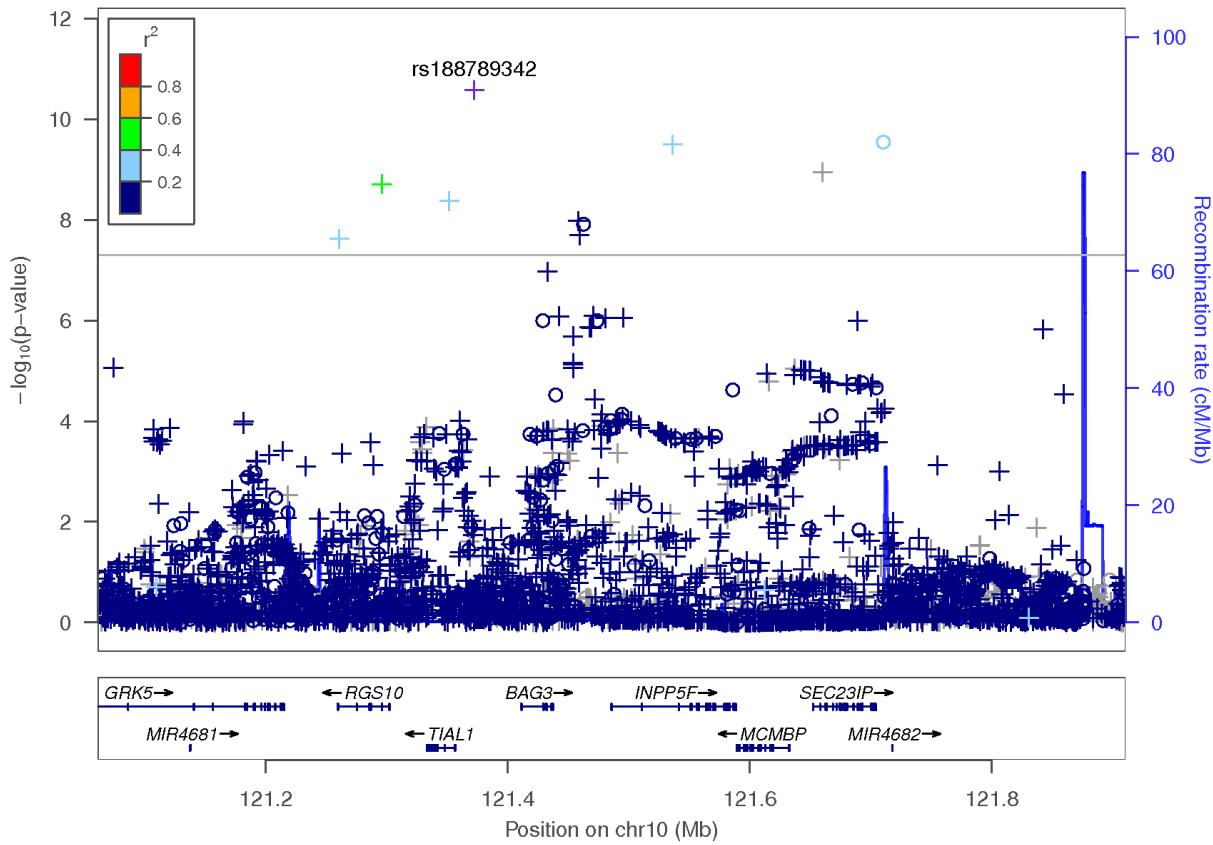
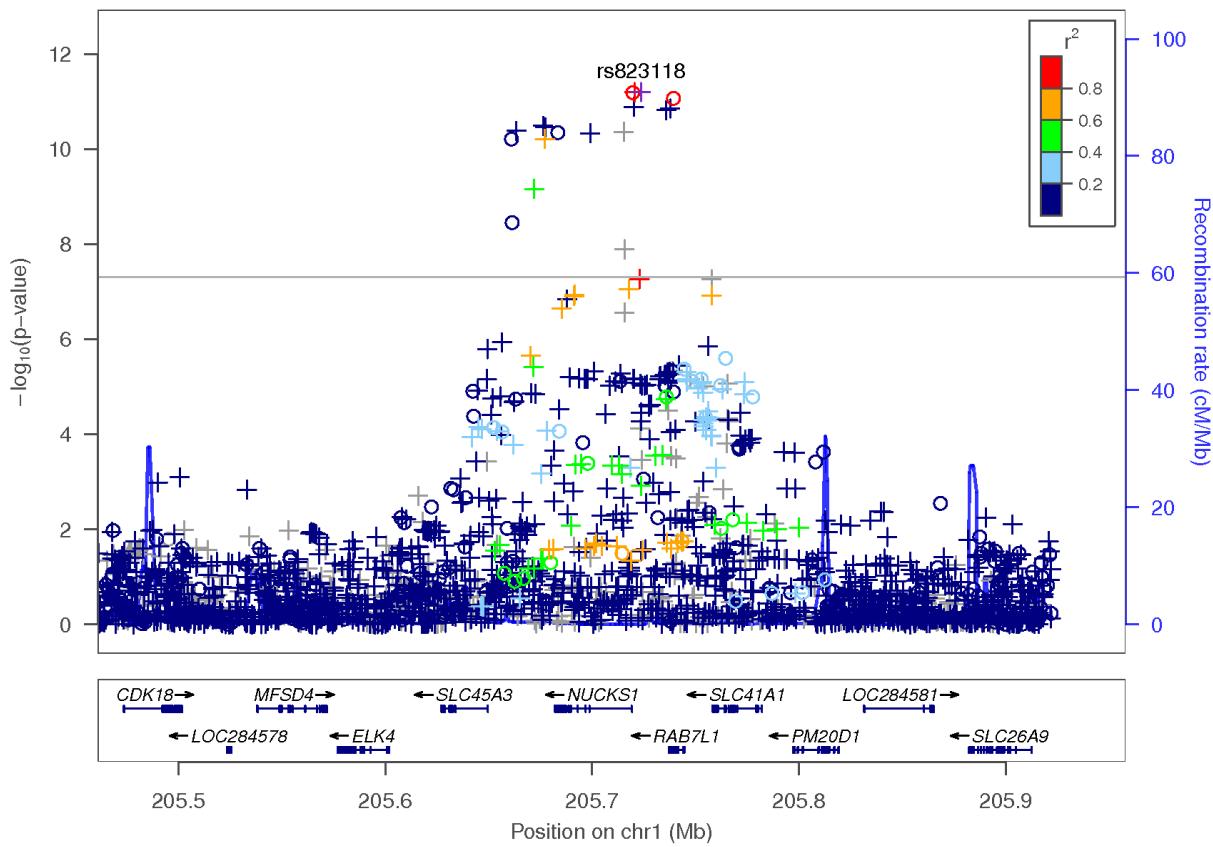


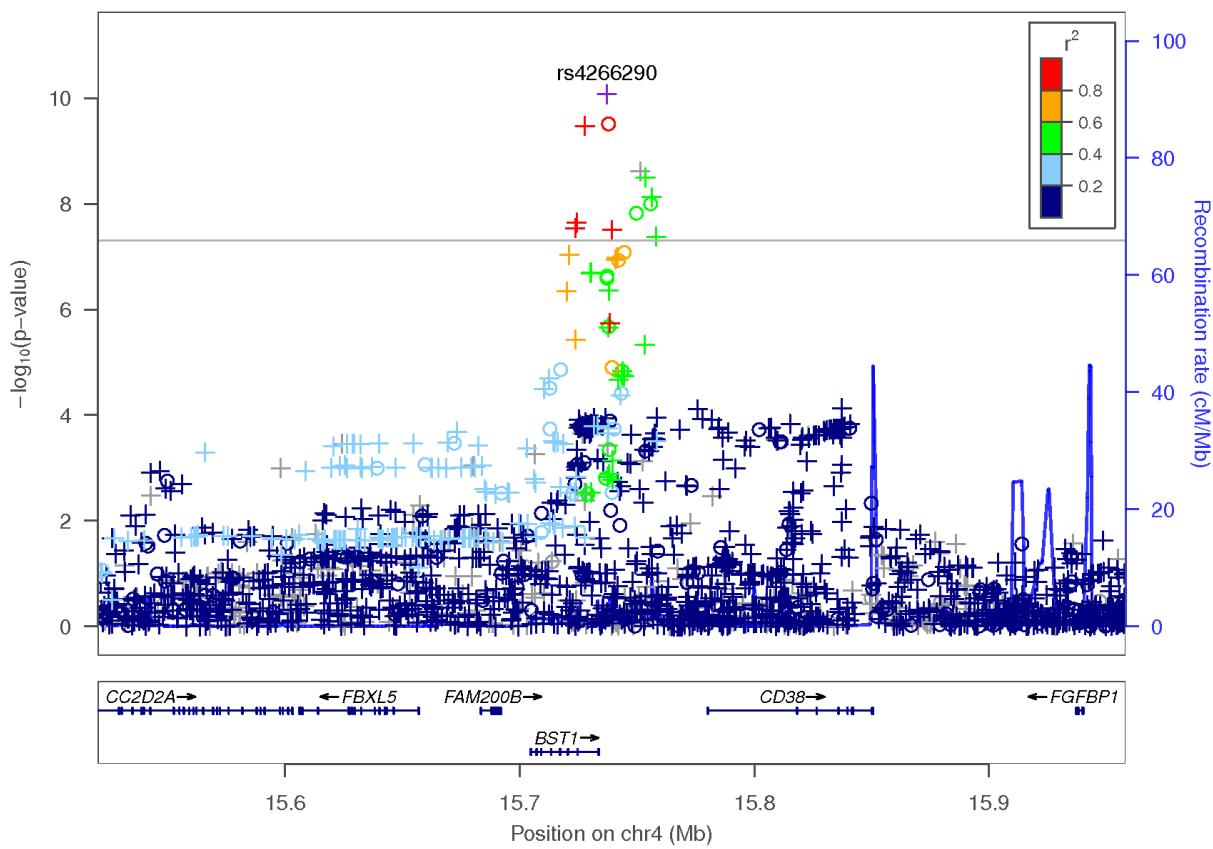
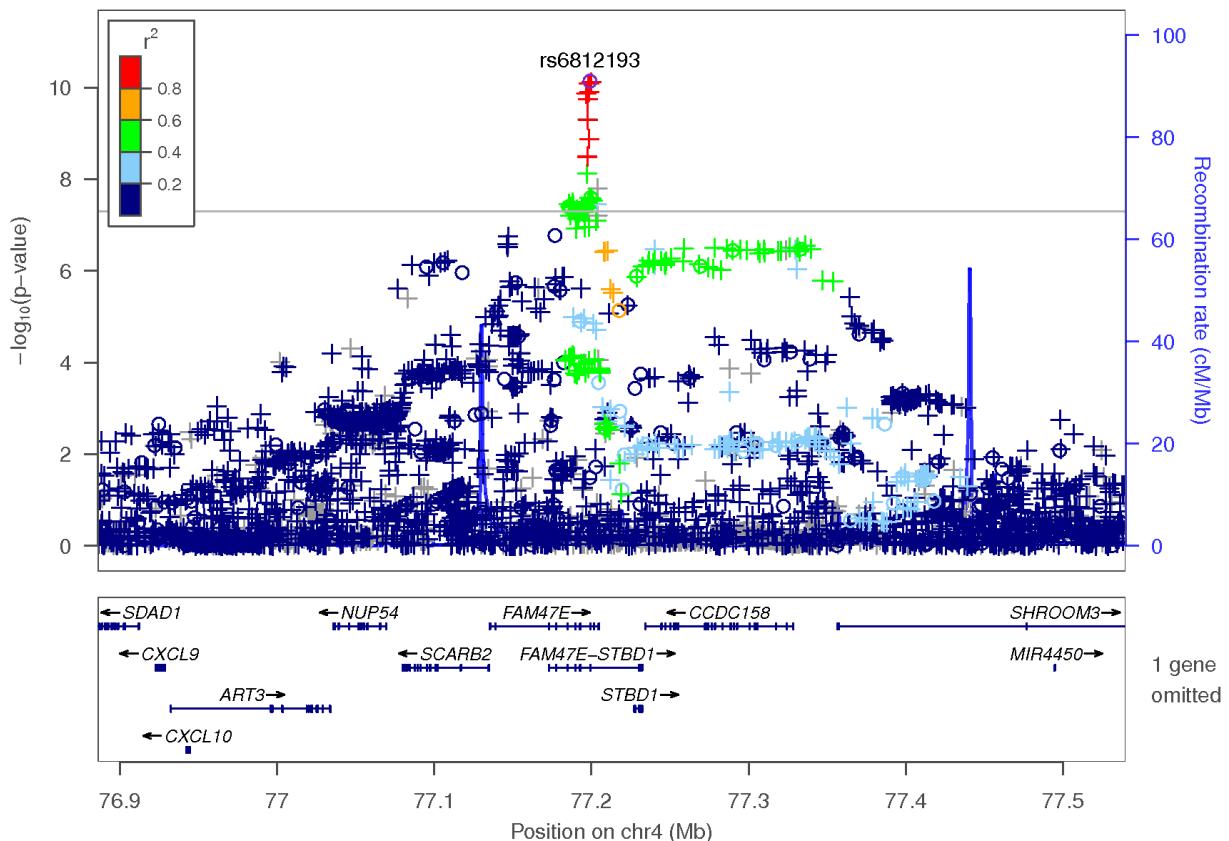


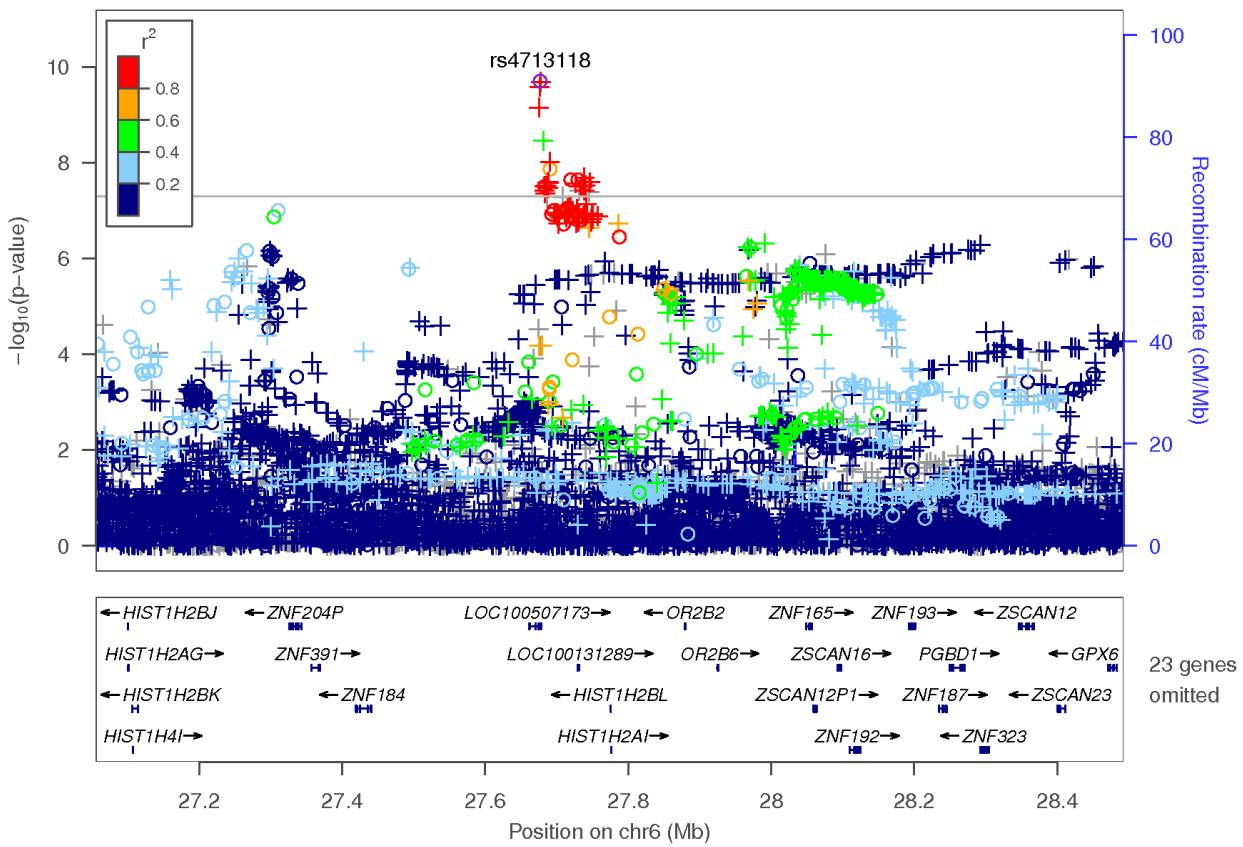
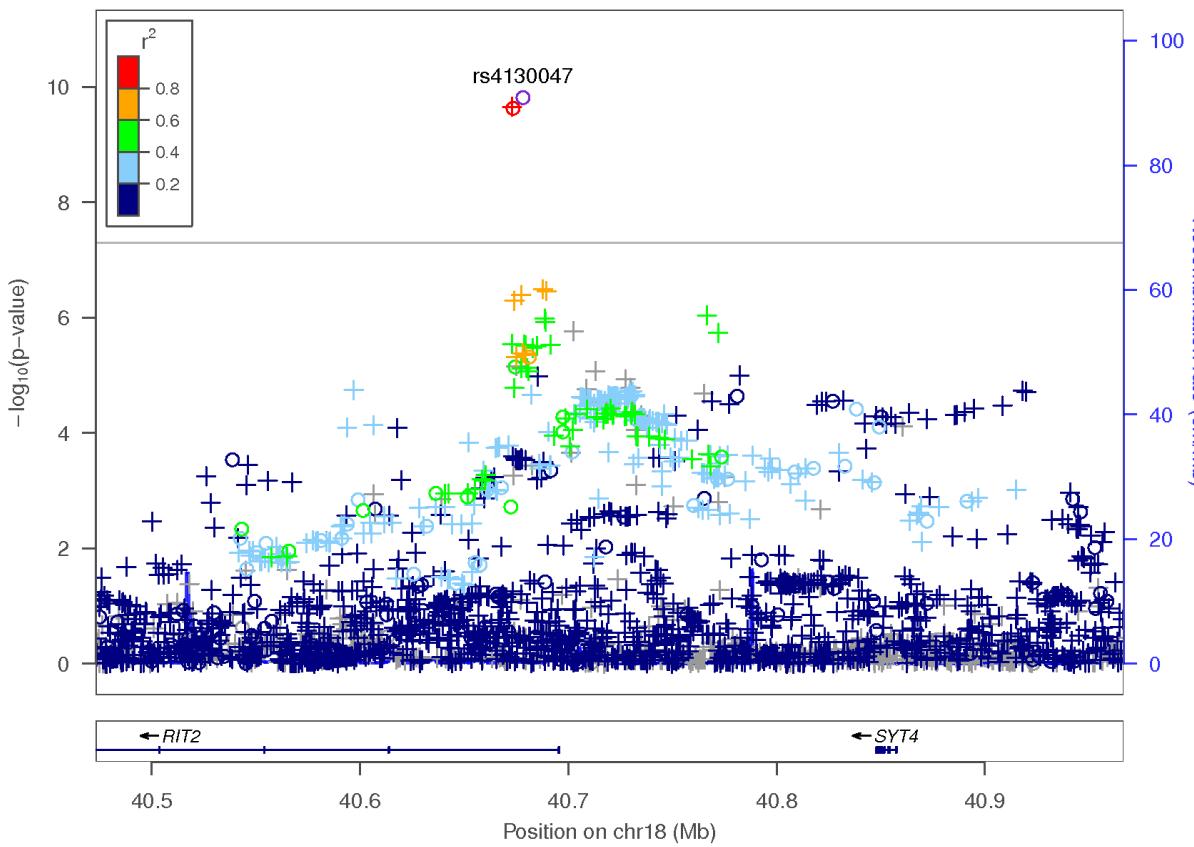


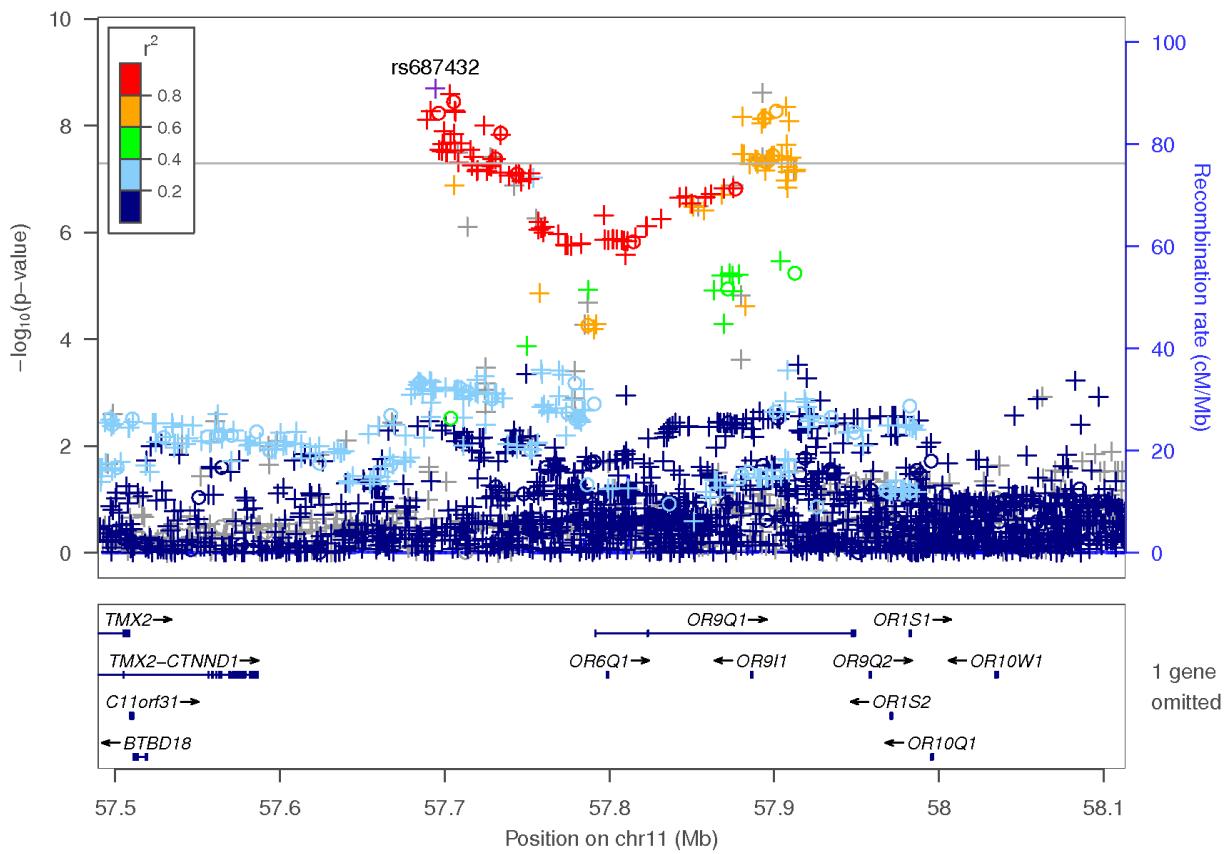
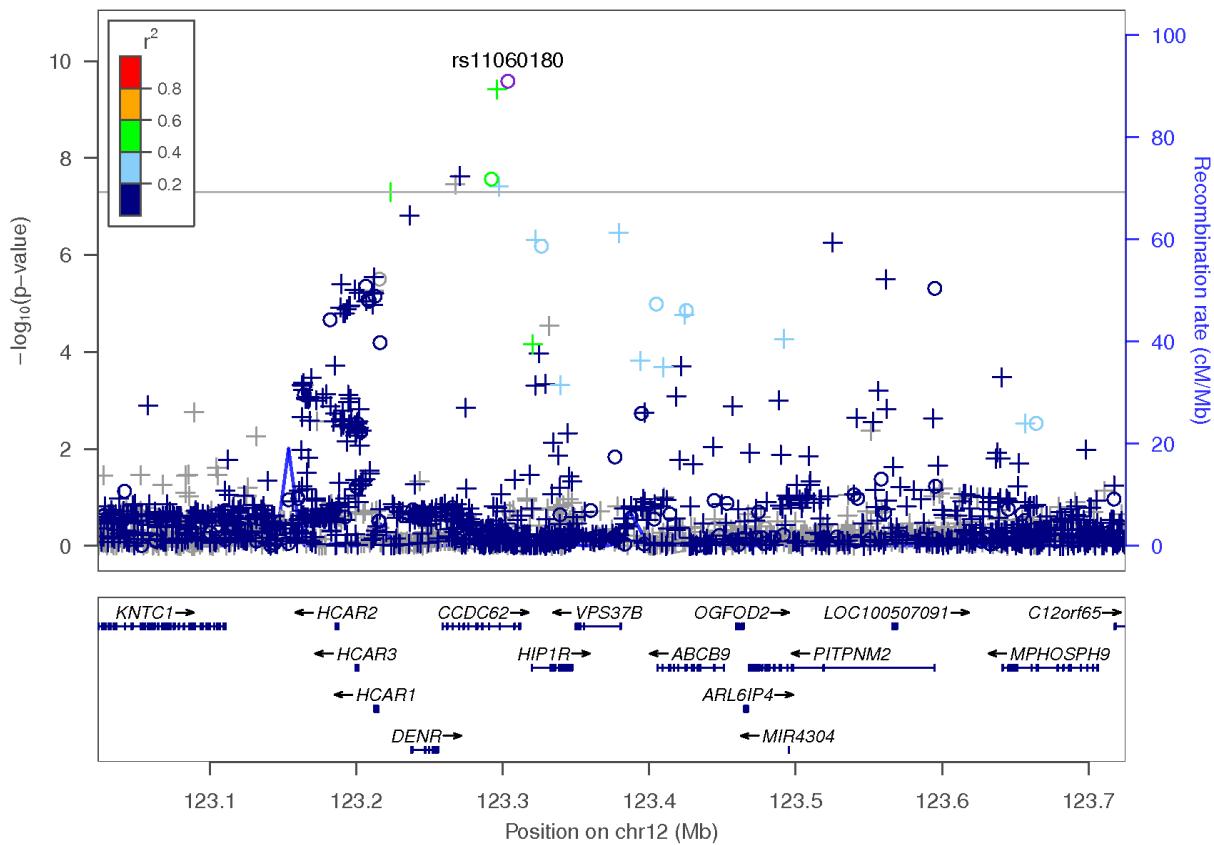


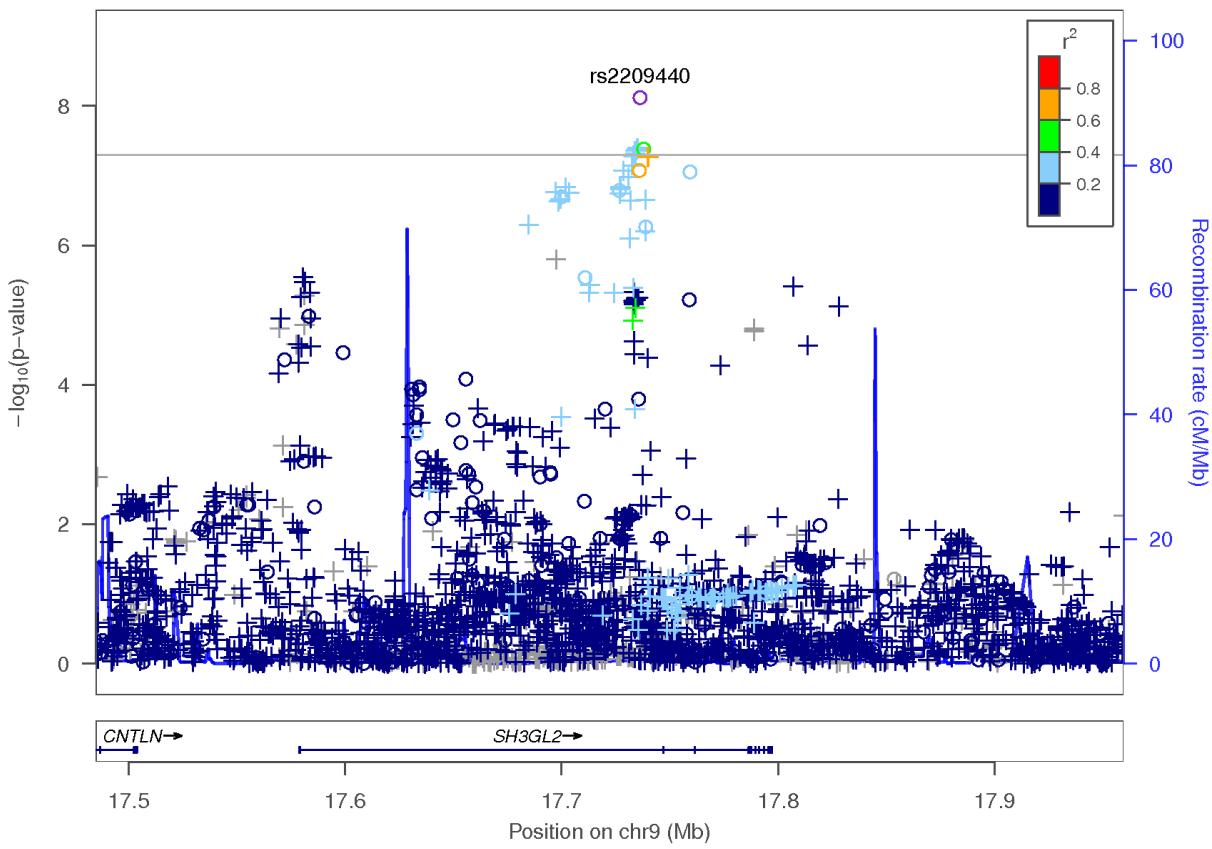
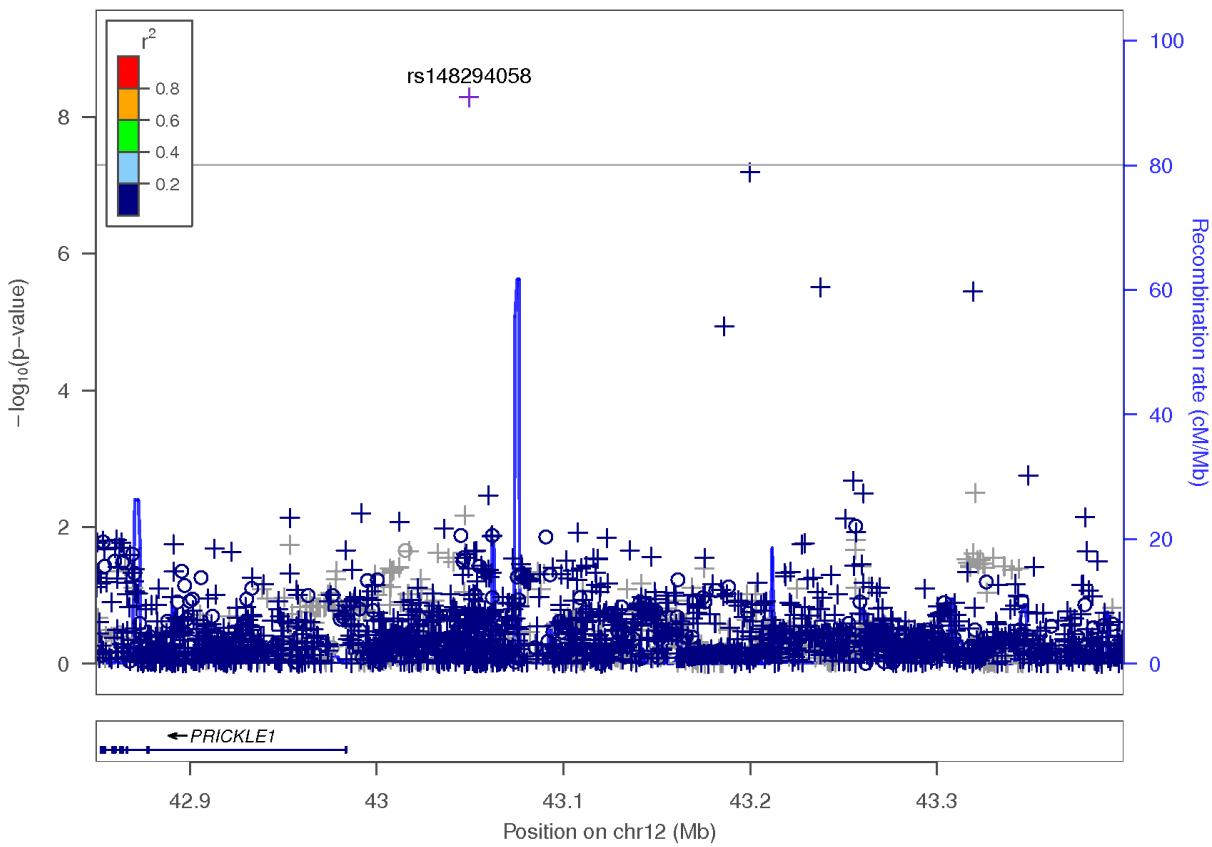


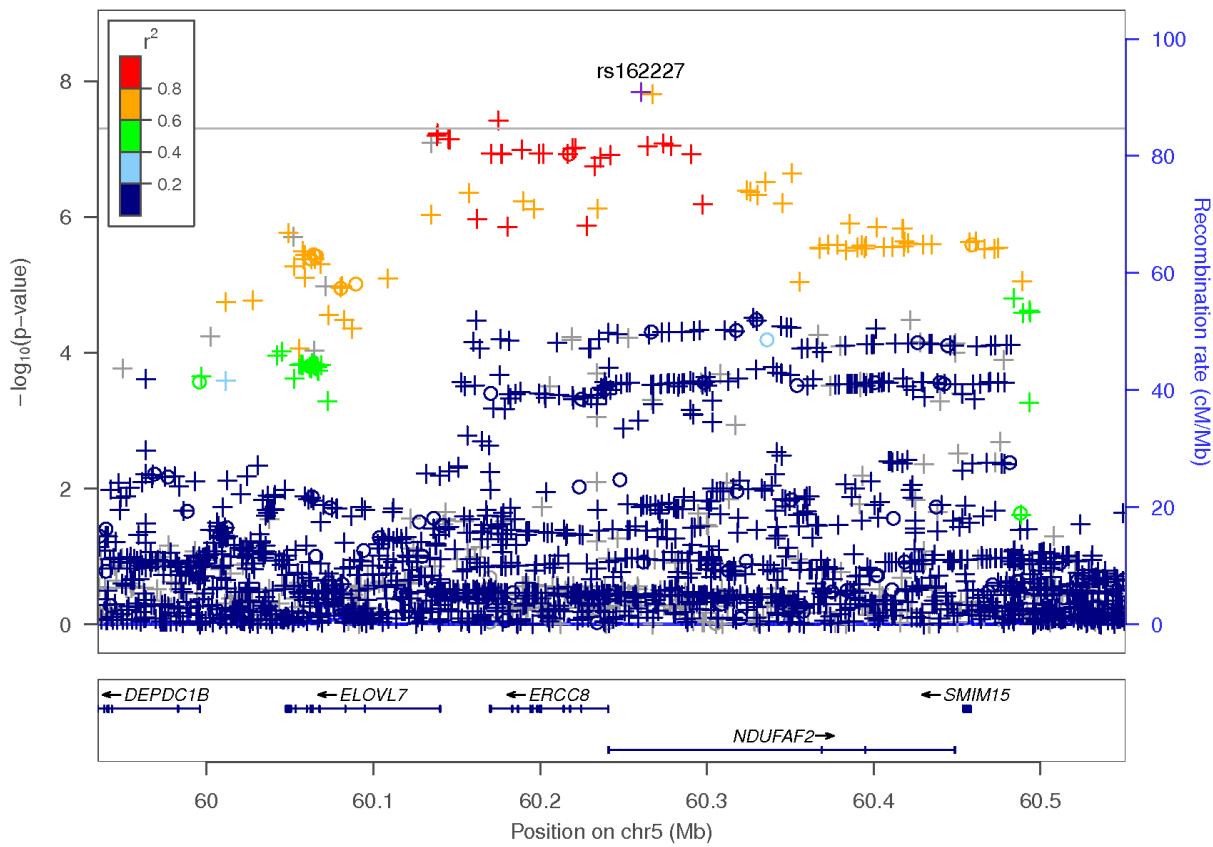
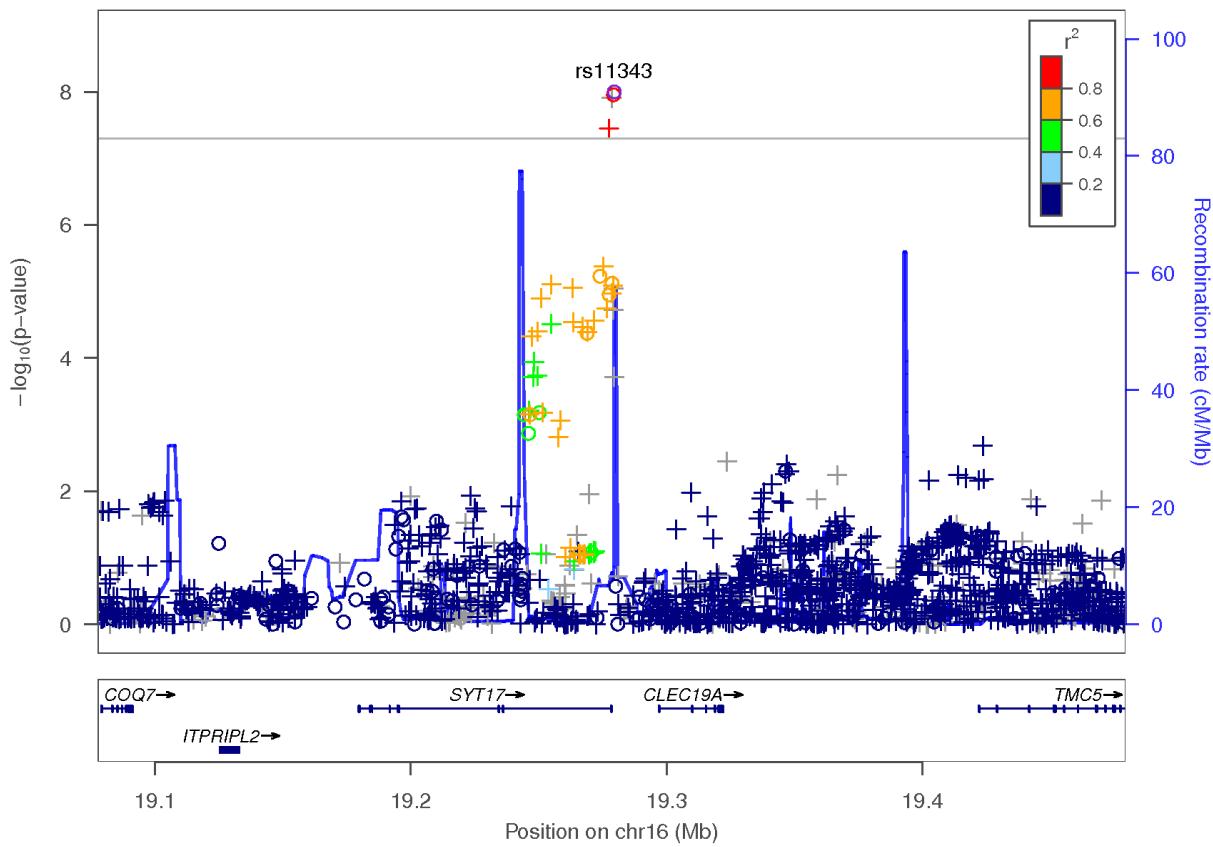


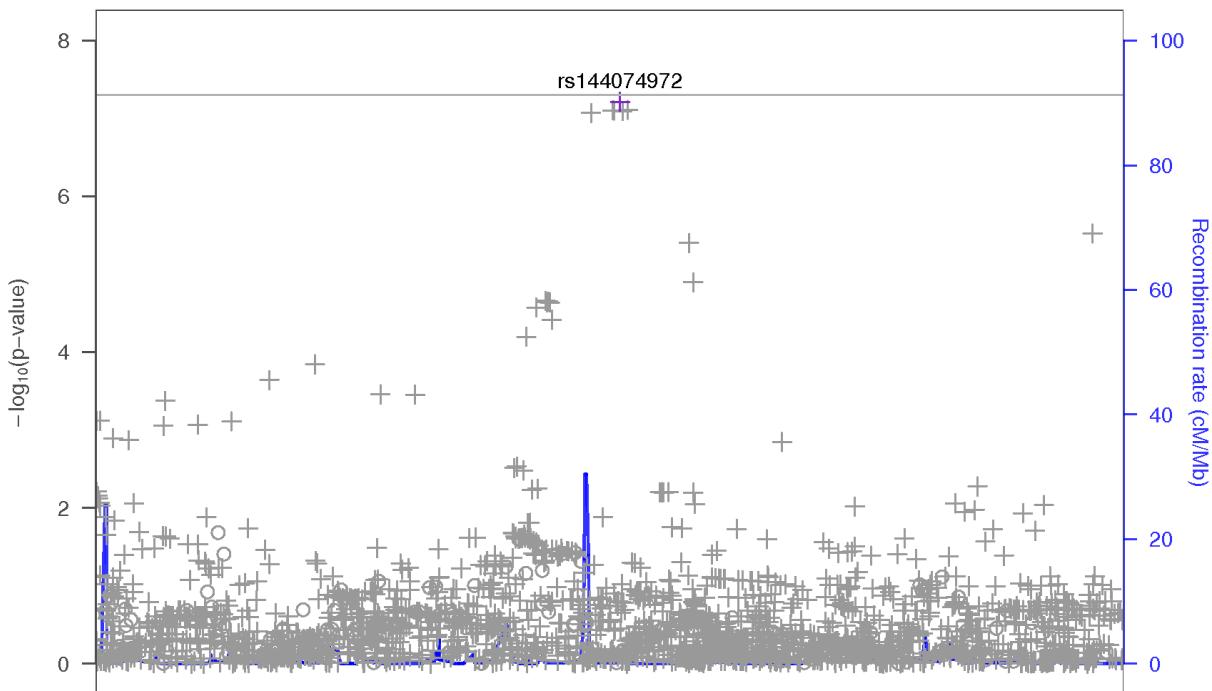




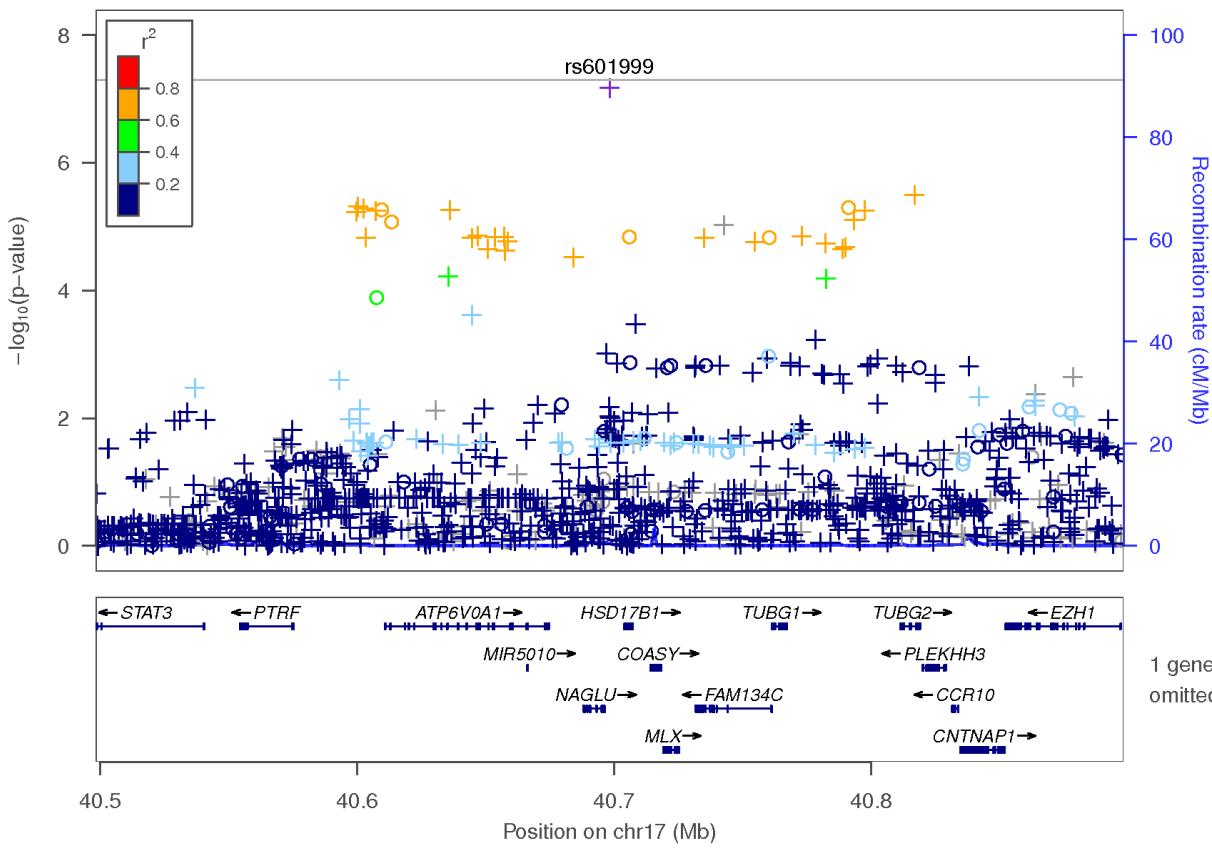








61 61.1 61.2 61.3
Position on chr8 (Mb)



40.5 40.6 40.7 40.8
Position on chr17 (Mb)

STAT3 PTRF ATP6V0A1 HSD17B1 TUBG1 TUBG2 EZH1
MIR5010 COASY NAGLU FAM134C PLEKHH3 CCR10 CNTNAP1
MLX 1 gene omitted