

Locus zoom plots

Locus#	Phenotype	Loci	SNP	Page
1	Total Surface Area	3:28007315	rs12630663	6
2	Total Surface Area	3:141721762	rs34464850	7
3	Total Surface Area	5:170778824	rs7715167	8
4	Total Surface Area	6:108926496	rs2802295	9
5	Total Surface Area	6:126964510	rs4273712	10
6	Total Surface Area	6:127052822	rs190958130	11
7	Total Surface Area	10:21830104	rs11012732	12
8	Total Surface Area	10:105012994	rs1628768	13
9	Total Surface Area	12:56470625	rs11171739	14
10	Total Surface Area	12:66327632	rs10878349	15
11	Total Surface Area	17:43904528	rs62057153	16
12	Average Thickness	2:27149258	rs6738528	17
13	Average Thickness	2:98275354	rs11692435	18
14	Average Thickness	3:39489619	rs630934	19
15	Average Thickness	4:121643726	rs13110077	20
16	Average Thickness	8:8189438	rs2980497	21
17	Average Thickness	8:110585288	rs7824177	22
18	Average Thickness	17:43927290	rs17769552	23
19	Frontal Pole Surface Area	2:188278203	rs17464221	24
20	Frontal Pole Surface Area	16:86088307	rs113050298	25
21	Lateral Orbitofrontal Surface Area	1:119799099	rs7529542	26
22	Lateral Orbitofrontal Surface Area	6:126760994	rs2184968	27
23	Lateral Orbitofrontal Surface Area	7:19626622	rs4721802	28
24	Lateral Orbitofrontal Surface Area	8:8975103	rs1822951	29
25	Lateral Orbitofrontal Surface Area	12:65905126	rs79487293	30
26	Lateral Orbitofrontal Surface Area	14:54724054	rs2358483	31
27	Lateral Orbitofrontal Surface Area	21:34298974	rs12626790	32
28	Rostral Anterior Cingulate Surface Area	9:101633220	rs7874287	33
29	Rostral Anterior Cingulate Surface Area	15:55983271	rs1986012	34
30	Rostral Anterior Cingulate Surface Area	17:46584657	rs2202895	35
31	Rostral Anterior Cingulate Thickness	14:73319928	rs4899448	36
32	Rostral Anterior Cingulate Thickness	19:51621561	rs62115964	37
33	Caudal Anterior Cingulate Surface Area	2:162822709	rs13021985	38
34	Caudal Anterior Cingulate Surface Area	5:82843609	rs7728751	39
35	Caudal Anterior Cingulate Surface Area	16:60827388	rs80241863	40
36	Superior Frontal Surface Area	1:64235727	rs4915928	41
37	Superior Frontal Surface Area	6:126023555	rs4492207	42
38	Superior Frontal Surface Area	9:83480392	rs76696867	43
39	Superior Frontal Surface Area	12:79951566	rs4842266	44
40	Superior Frontal Thickness	16:2886967	rs9929219	45
41	Rostral Middle Frontal Surface Area	1:17311882	rs6682671	46
42	Rostral Middle Frontal Surface Area	2:188379320	rs35612915	47
43	Rostral Middle Frontal Surface Area	3:193536820	rs1165645	48
44	Rostral Middle Frontal Surface Area	5:94072933	rs7731360	49
45	Rostral Middle Frontal Surface Area	8:120596023	rs10283100	50
46	Rostral Middle Frontal Surface Area	11:65562257	rs2276133	51
47	Rostral Middle Frontal Surface Area	13:100825526	rs4772268	52
48	Rostral Middle Frontal Surface Area	14:99698931	rs1257415	53
49	Rostral Middle Frontal Surface Area	15:39634222	rs1080066	54
50	Rostral Middle Frontal Surface Area	16:65014909	rs40115	55
51	Rostral Middle Frontal Thickness	1:247527168	rs6671321	56
52	Rostral Middle Frontal Thickness	3:55067916	rs5008703	57
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54	Pars Orbitalis Surface Area	2:65971597	rs2287283	59
55	Pars Orbitalis Surface Area	3:88380871	rs6551313	60
56	Pars Orbitalis Surface Area	3:98722159	rs2262601	61
57	Pars Orbitalis Surface Area	11:92348702	rs61901866	62
58	Pars Orbitalis Surface Area	14:54777745	rs7147119	63
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60	Pars Triangularis Surface Area	3:147106319	rs2279829	65

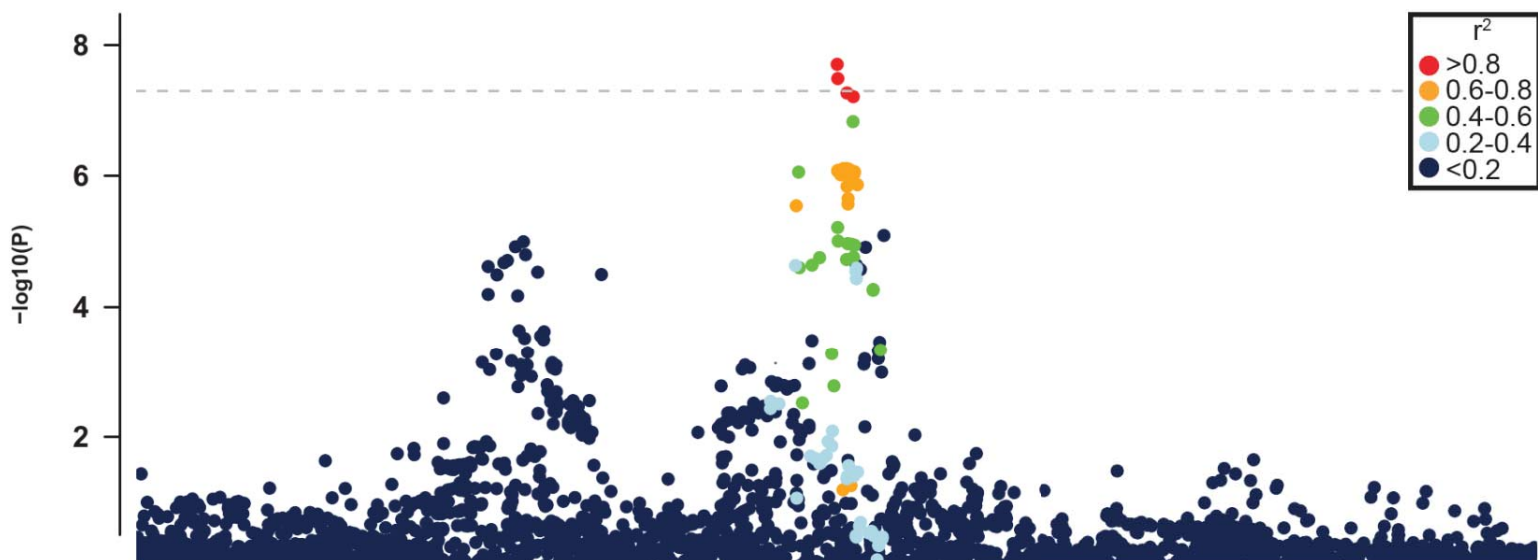
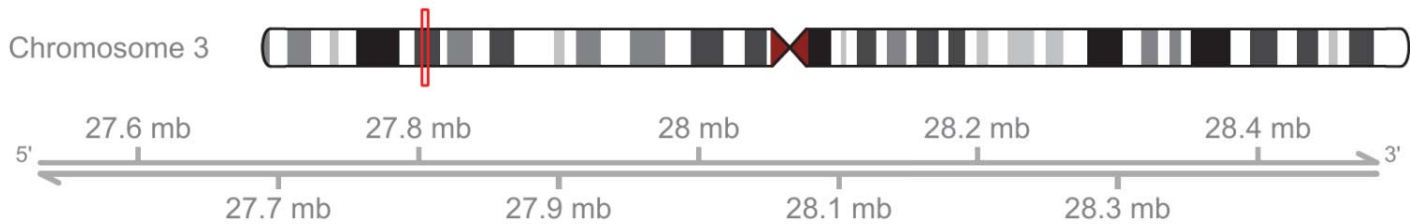
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67	Pars Opercularis Surface Area	15:39633904	rs2033939	72
68	Pars Opercularis Surface Area	17:13054525	rs12938190	73
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75	Caudal Middle Frontal Surface Area	16:52538337	rs7184835	80
76	Caudal Middle Frontal Surface Area	18:52918717	rs1893430	81
77	Caudal Middle Frontal Thickness	5:171928642	rs62385486	82
78	Paracentral Surface Area	10:33539487	rs2269084	83
79	Paracentral Surface Area	12:27792335	rs7303526	84
80	Paracentral Surface Area	12:106476805	rs12146713	85
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88	Precentral Thickness	7:45582776	rs4724413	93
89	Postcentral Surface Area	3:147106319	rs2279829	94
90	Postcentral Surface Area	4:127537734	rs313135	95
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97	Postcentral Thickness	15:39631771	rs11639391	102
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107	Precuneus Thickness	1:47984143	rs12118339	112
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122	Superior Parietal Thickness	7:120721087	rs73215353	127

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127	Supramarginal Surface Area	14:59069053	rs221326	132
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130	Supramarginal Surface Area	15:39639898	rs4924345	135
131	Supramarginal Surface Area	15:39720600	rs78502100	136
132	Inferior Parietal Surface Area	1:88423397	rs1413536	137
133	Inferior Parietal Surface Area	2:4554304	rs639016	138
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141	Inferior Parietal Surface Area	20:52432451	rs4437022	146
142	Posterior Cingulate Surface Area	1:87630392	rs11161942	147
143	Posterior Cingulate Surface Area	2:37175444	rs11695609	148
144	Posterior Cingulate Surface Area	5:66155020	rs72761270	149
145	Posterior Cingulate Surface Area	10:126486443	rs12764880	150
146	Isthmus Cingulate Surface Area	2:37150793	rs3770776	151
147	Isthmus Cingulate Surface Area	12:78241285	rs1123680	152
148	Isthmus Cingulate Thickness	10:126551900	rs2559509	153
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150	Insula Surface Area	10:118752250	rs1122688	155
151	Insula Surface Area	11:111056158	rs58066679	156
152	Insula Surface Area	17:28275095	rs6505147	157
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154	Entorhinal Surface Area	5:81882571	rs4147321	159
155	Entorhinal Surface Area	14:59860705	rs7141150	160
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157	Parahippocampal Surface Area	5:82350191	rs27493	162
158	Parahippocampal Surface Area	6:126964510	rs4273712	163
159	Parahippocampal Surface Area	11:92319654	rs1792354	164
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164	Fusiform Surface Area	14:59827306	rs17834032	169
165	Fusiform Surface Area	17:44865439	rs2074404	170
166	Temporal Pole Surface Area	4:103112470	rs6855246	171
167	Inferior Temporal Surface Area	2:37899310	rs9309013	172
168	Inferior Temporal Surface Area	19:13165884	rs73006837	173
169	Middle Temporal Surface Area	2:237708487	rs1344762	174
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173	Middle Temporal Surface Area	19:13109955	rs73006822	178
174	Middle Temporal Thickness	14:59074136	rs160459	179
175	Superior Temporal Surface Area	2:48277490	rs7601767	180
176	Superior Temporal Surface Area	2:149994571	rs13011264	181
177	Superior Temporal Surface Area	8:8605632	rs4841029	182
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179	Superior Temporal Thickness	1:47980916	rs6658111	184
180	Superior Temporal Thickness	3:48753149	rs9848203	185
181	Banks of the Superior Temporal Sulcus Surface Area	9:126603989	rs7862092	186
182	Banks of the Superior Temporal Sulcus Surface Area	14:59074878	rs160458	187

Locus#	Phenotype	Loci	SNP	Page
183	Banks of the Superior Temporal Sulcus Surface Area	19:13109955	rs73006822	188
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185	Transverse Temporal Surface Area	2:150037201	rs11684511	190
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187	Transverse Temporal Surface Area	5:131341541	rs7714191	192
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192	Transverse Temporal Surface Area	22:46760481	rs9615351	197
193	Transverse Temporal Thickness	4:35449515	rs57125540	198
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235	Pericalcarine Thickness	14:59605116	rs117461235	240
236	Cuneus Surface Area	2:204473751	rs7585601	241
237	Cuneus Surface Area	3:104647183	rs4895120	242
238	Cuneus Surface Area	6:126613946	rs76470478	243
239	Cuneus Surface Area	7:18877408	rs957960	244
240	Cuneus Surface Area	11:12071855	rs10765918	245
241	Cuneus Surface Area	14:59064739	rs170239	246
242	Cuneus Surface Area	14:59625997	rs73313052	247

Locus#	Phenotype	Loci	SNP	Page
243	Cuneus Surface Area	17:63912574	rs11652557	248
244	Cuneus Thickness	14:59610076	rs78580207	249
245	Lateral Occipital Surface Area	2:45170153	rs4953152	250
246	Lateral Occipital Surface Area	3:104678133	rs9863836	251
247	Lateral Occipital Surface Area	3:119001407	rs9813454	252
248	Lateral Occipital Surface Area	3:177295215	rs552305	253
249	Lateral Occipital Surface Area	6:127096181	rs9401907	254
250	Lateral Occipital Surface Area	9:98257842	rs28504650	255
251	Lateral Occipital Surface Area	14:59628609	rs74826997	256

Total Surface Area SA: rs12630663



← SLC4A7

EOMES

CMC1

AZI2

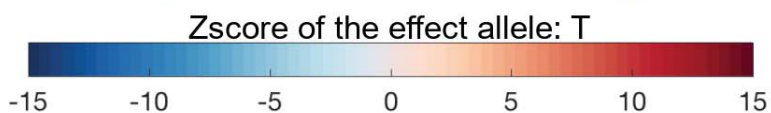
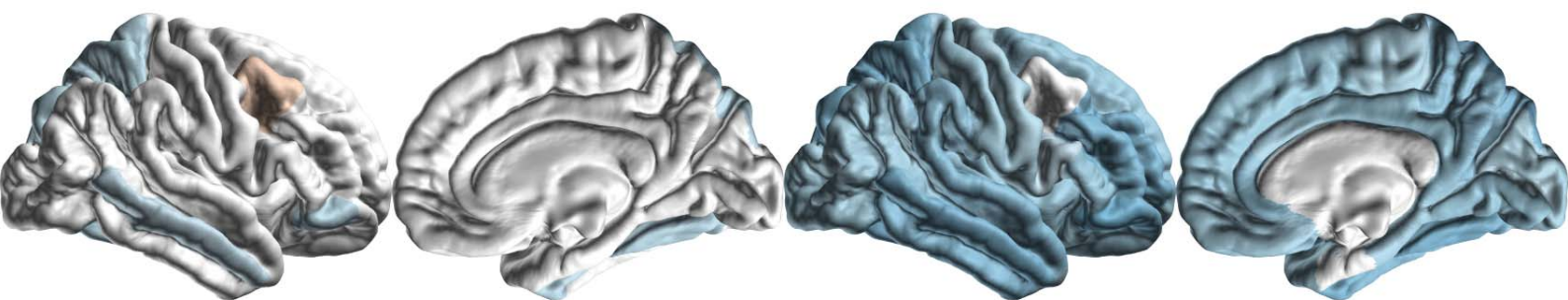
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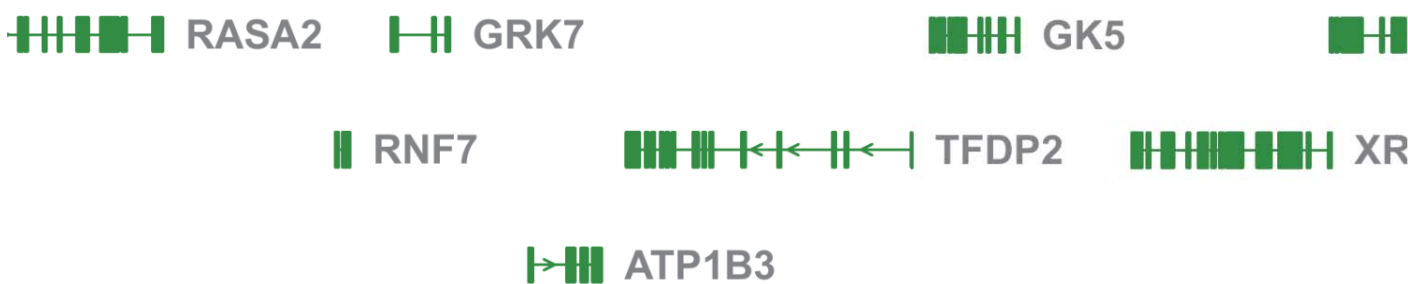
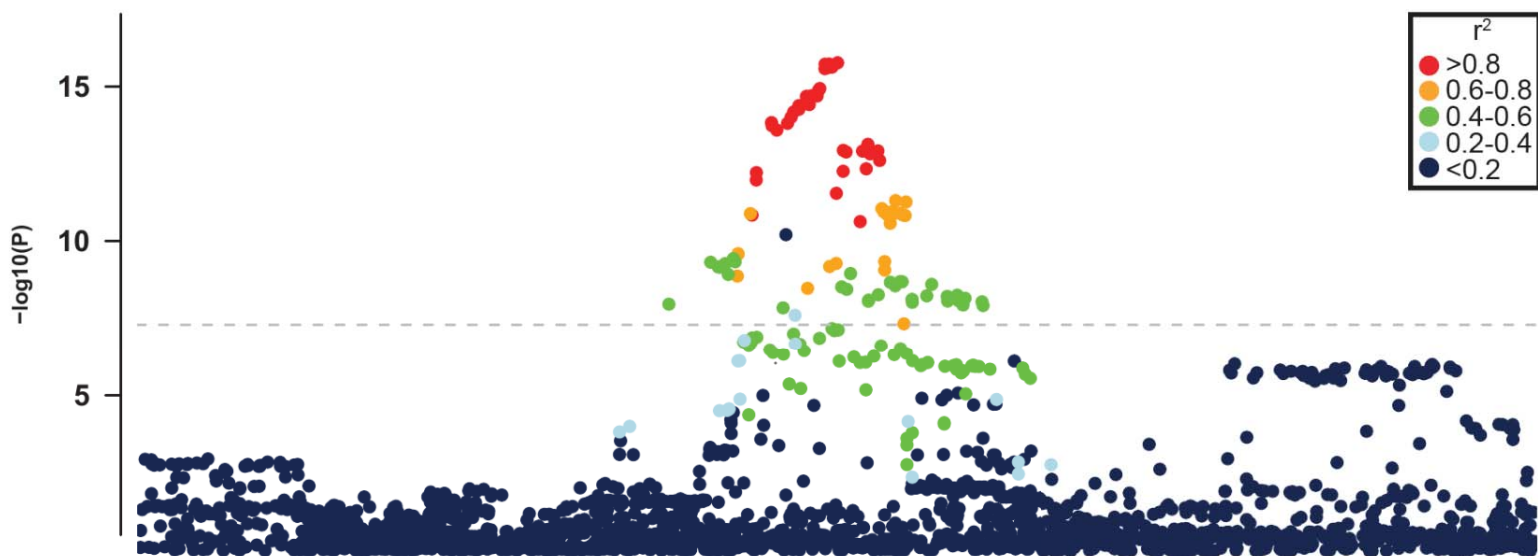
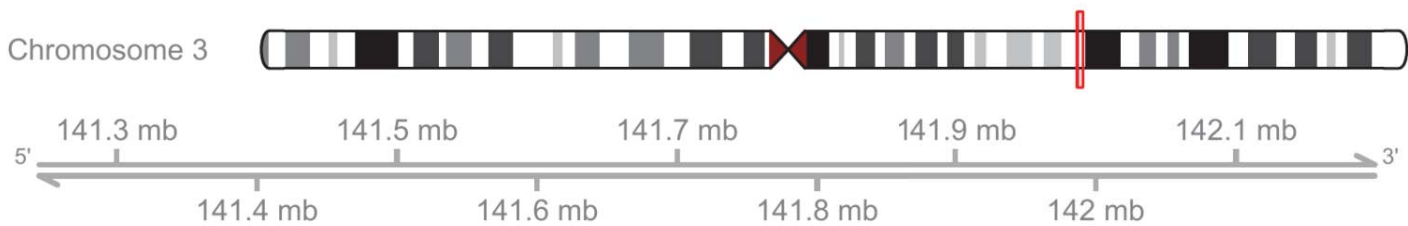
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Corrected for total SA

Uncorrected

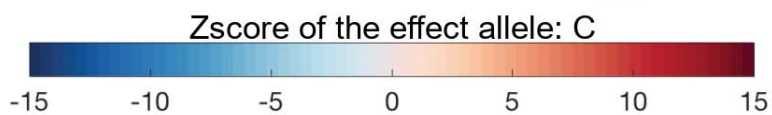
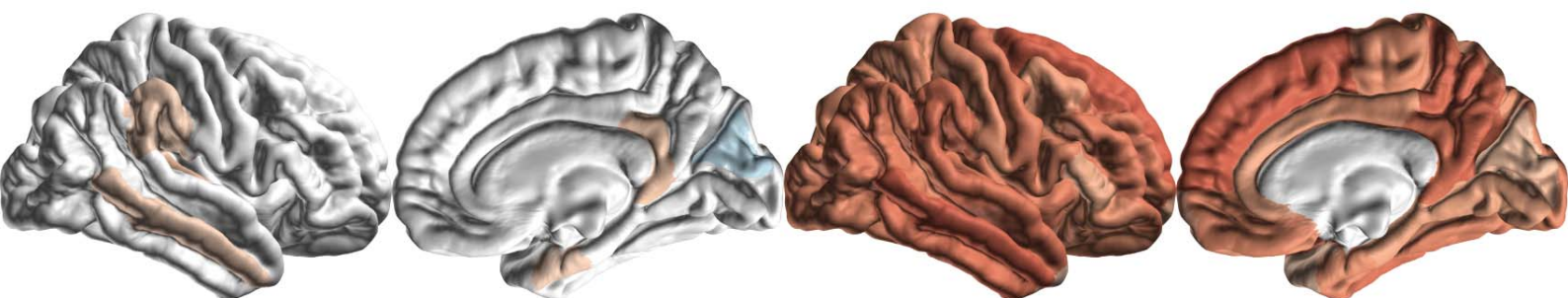


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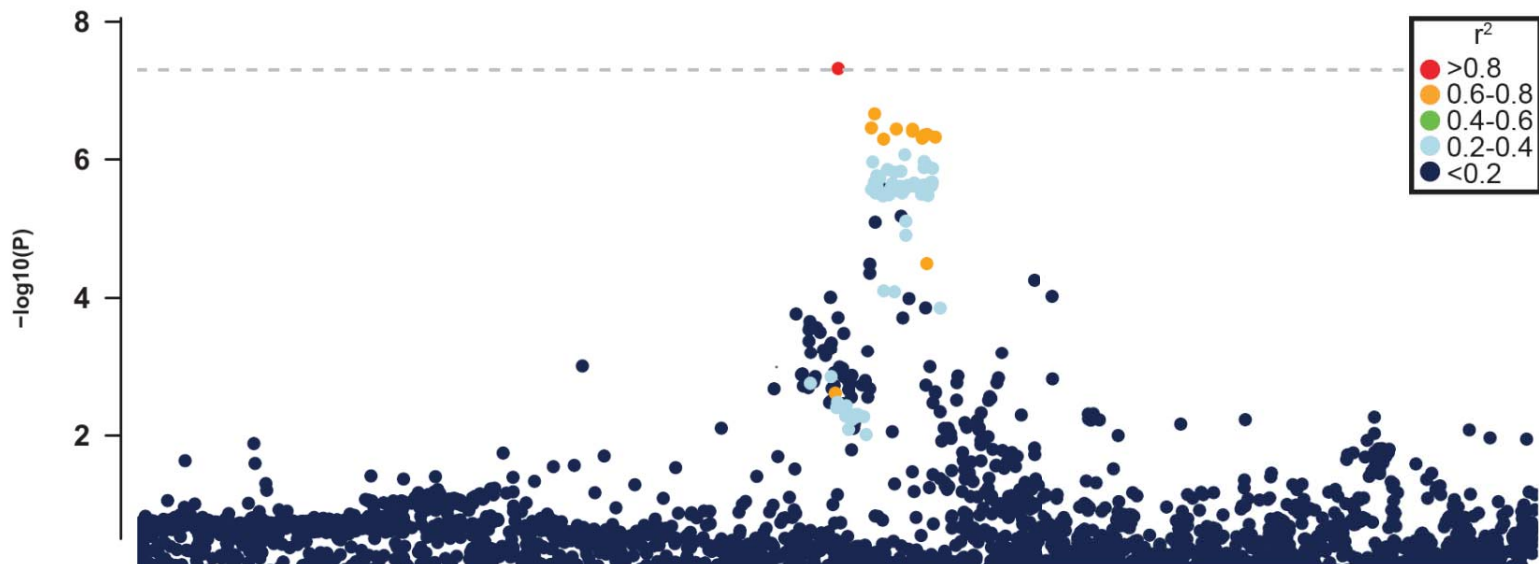
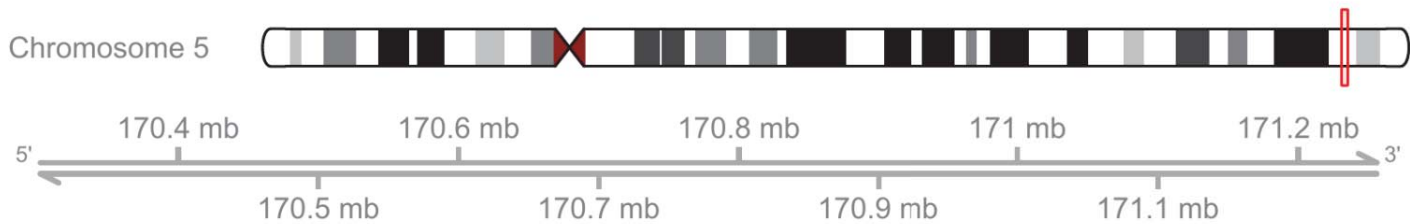


Corrected for total SA

Uncorrected



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RANBP17

TLX3 FGF18

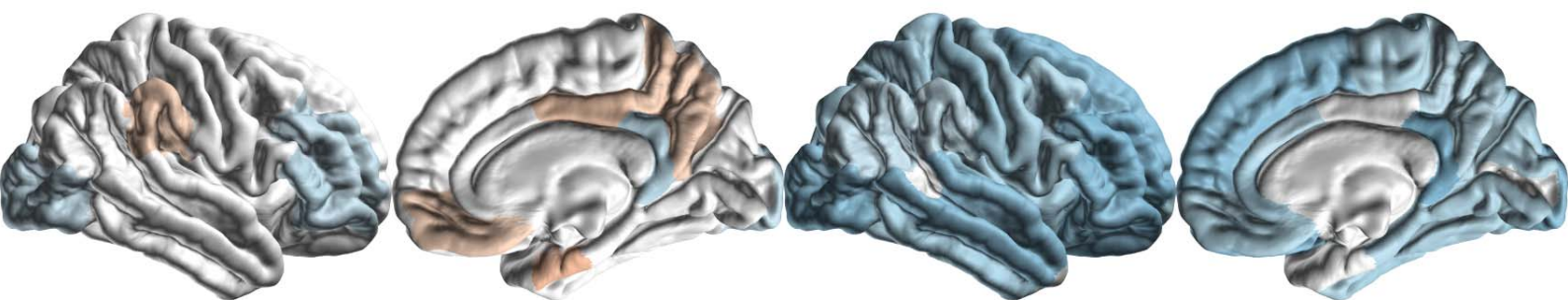
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CP>GZ

Corrected for total SA

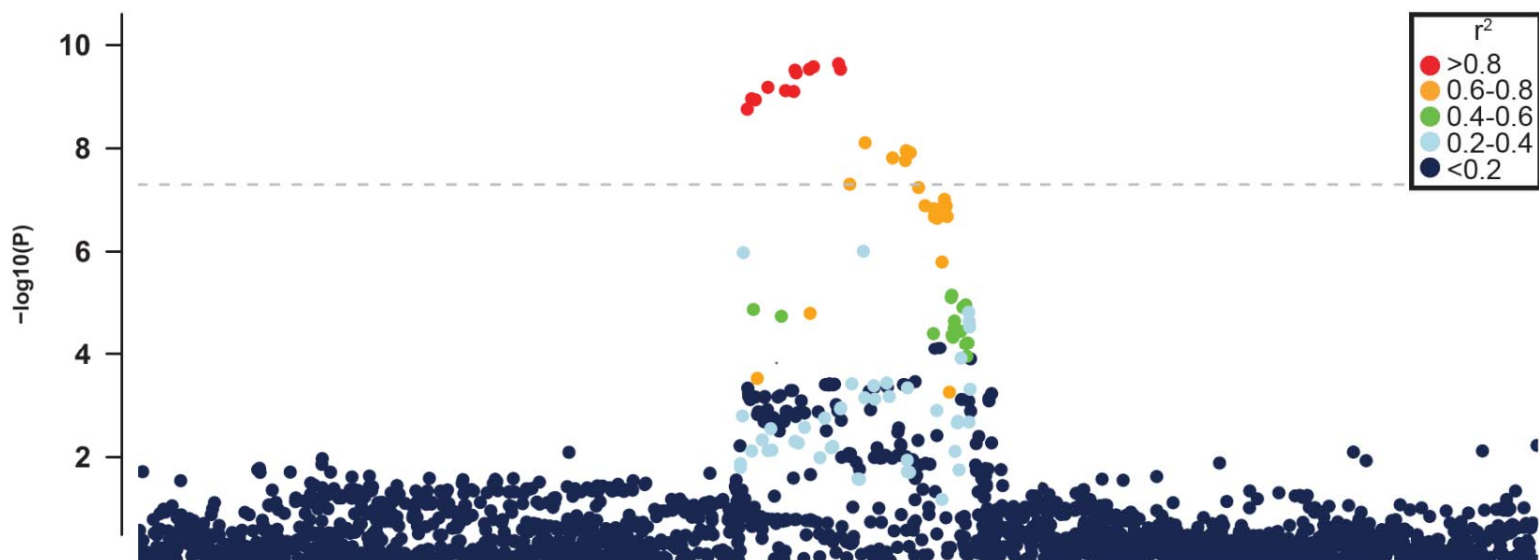
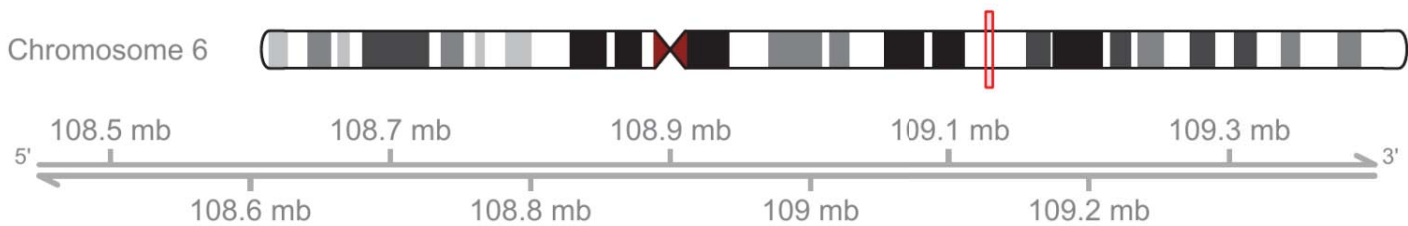
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Zscore of the effect allele: T

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Total Surface Area SA: rs2802295



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ARMC2

SNX3

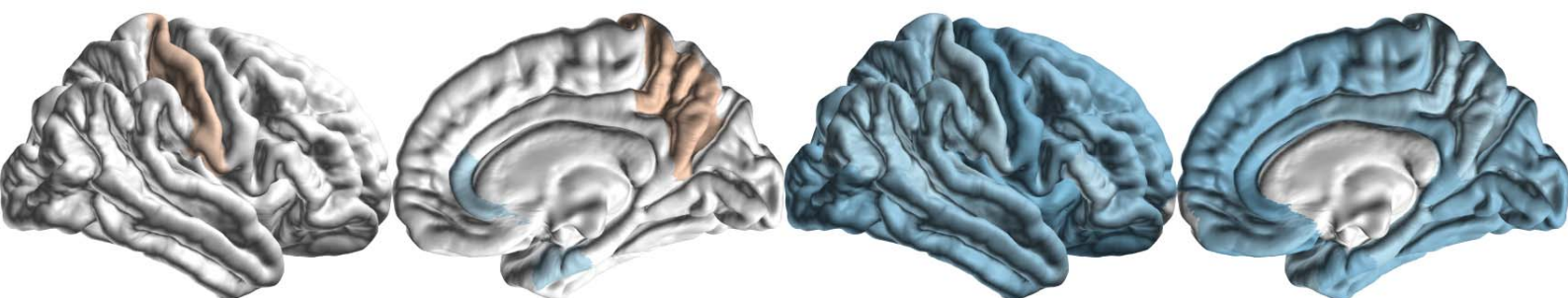
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GZ>CP

CP>GZ

Corrected for total SA

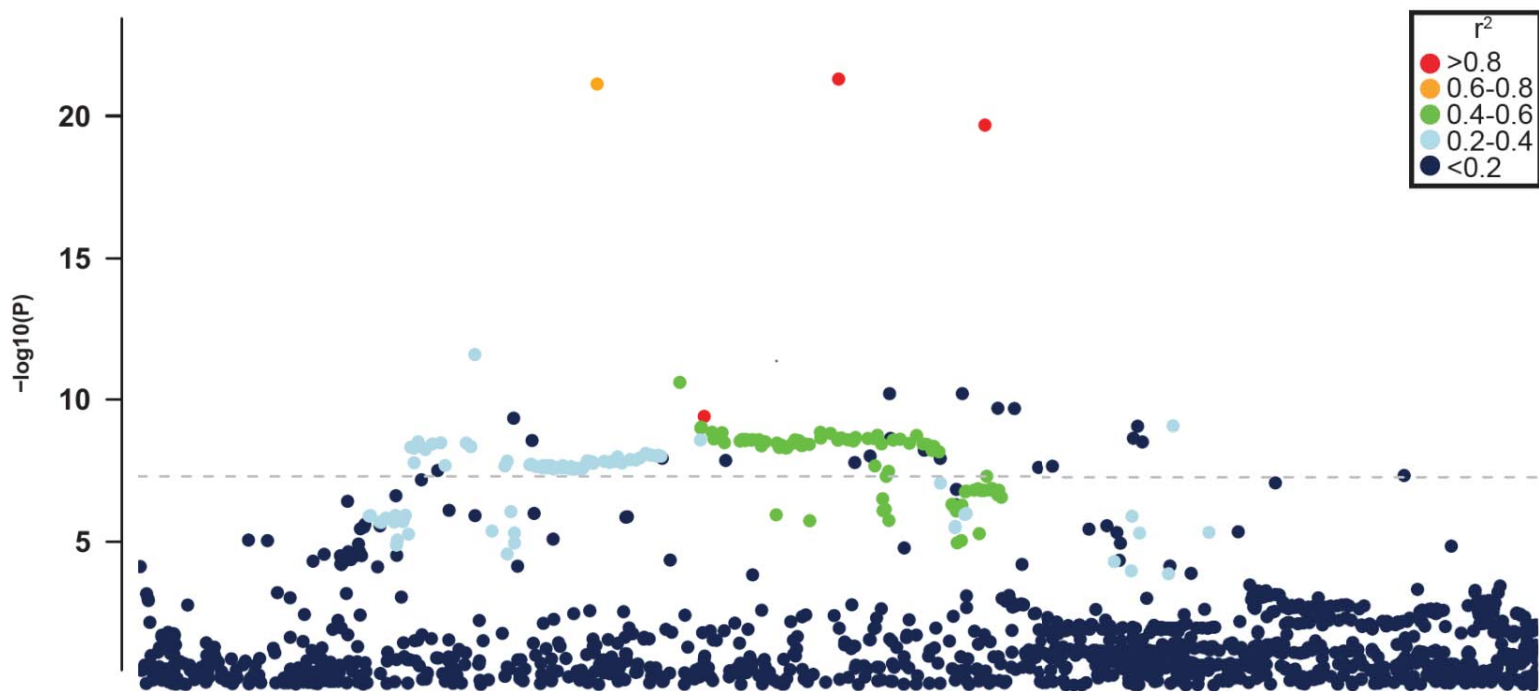
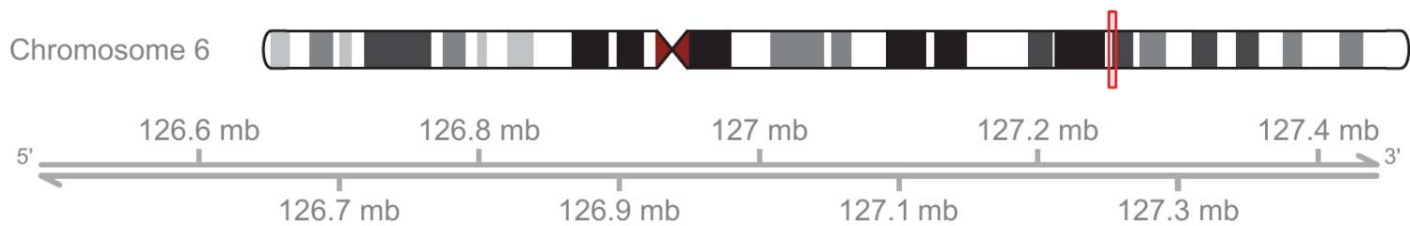
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Zscore of the effect allele: A



Total Surface Area SA: rs4273712



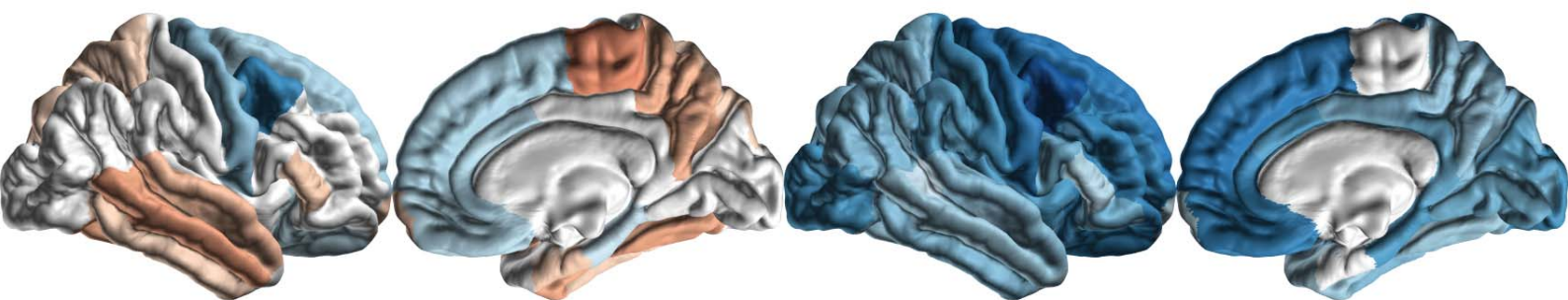
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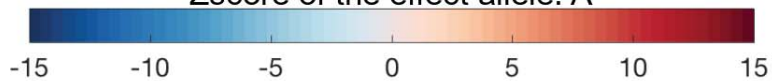
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Corrected for total SA

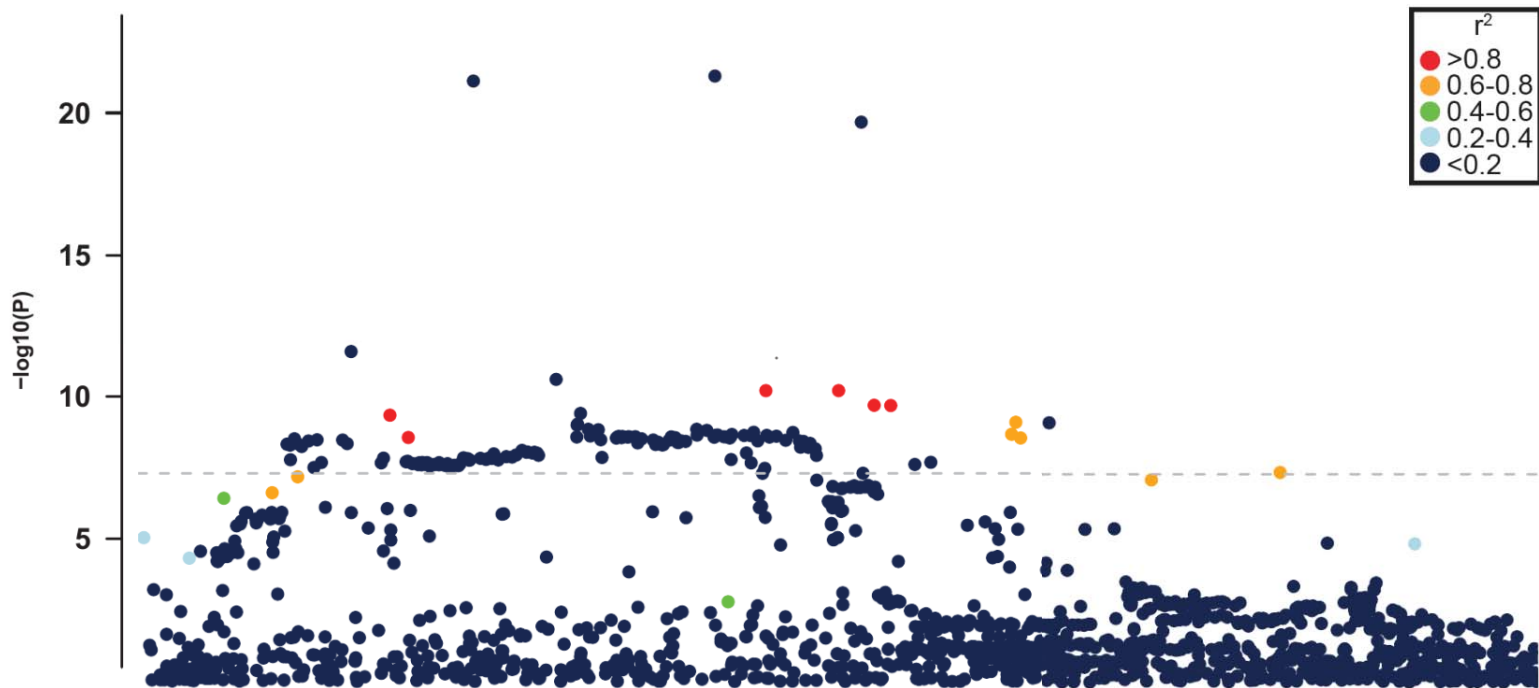
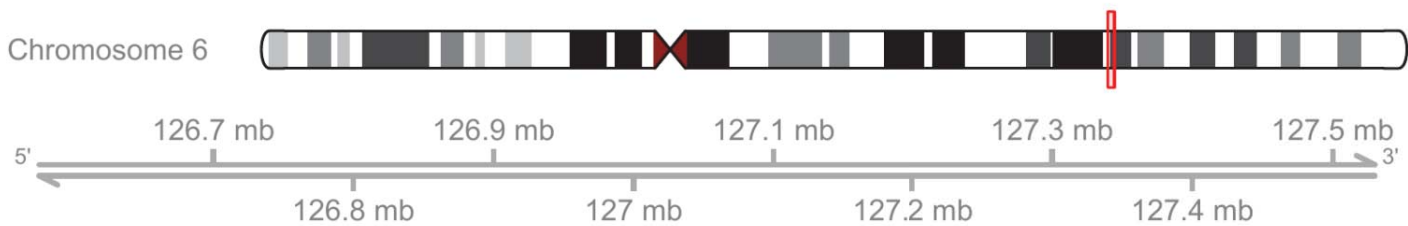
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Zscore of the effect allele: A



Total Surface Area SA: rs190958130



CENPW

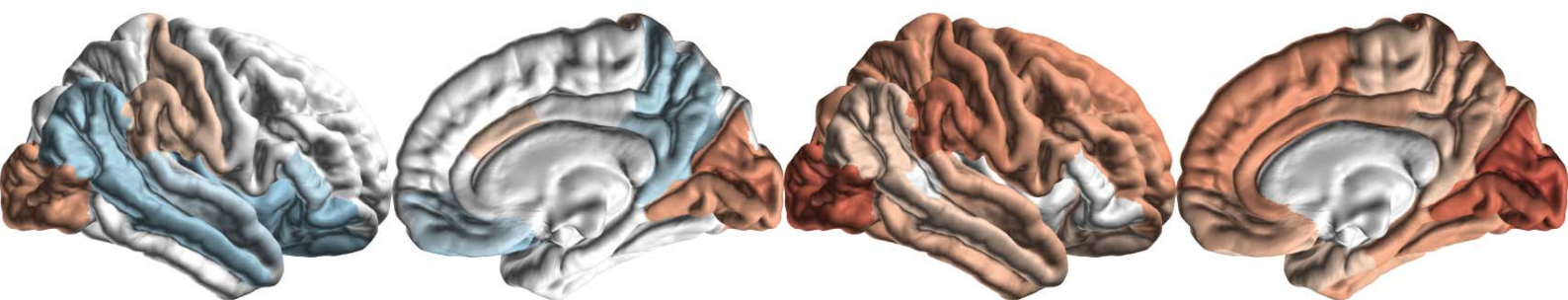
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GZ>CP

CP>GZ

Corrected for total SA

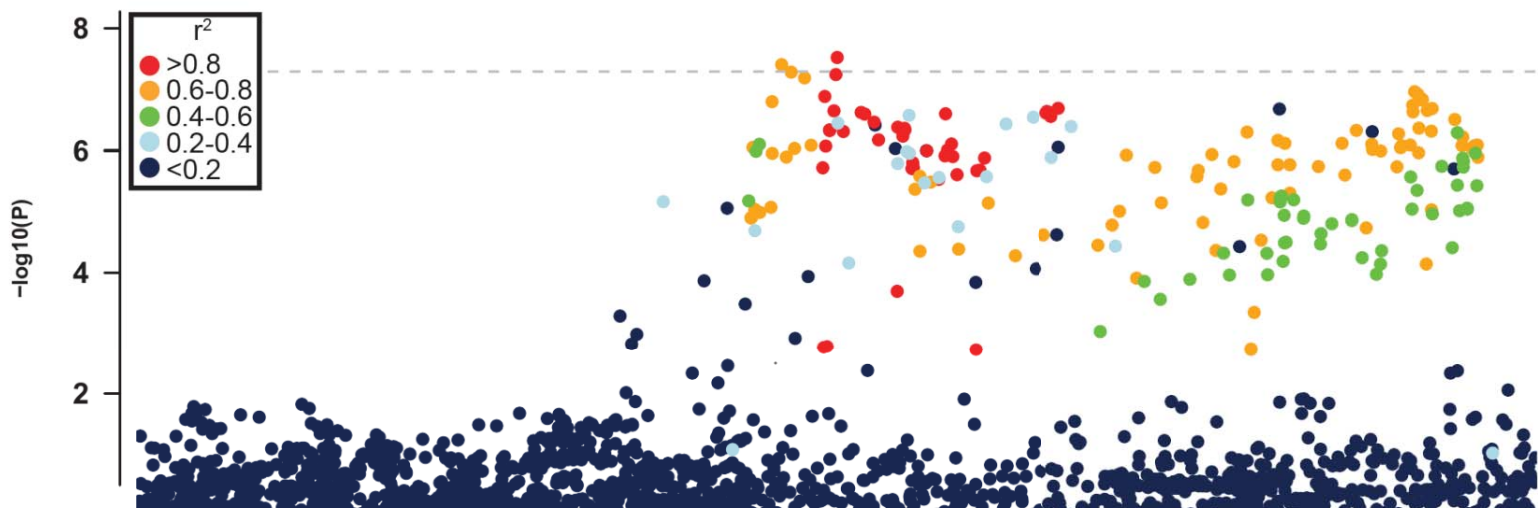
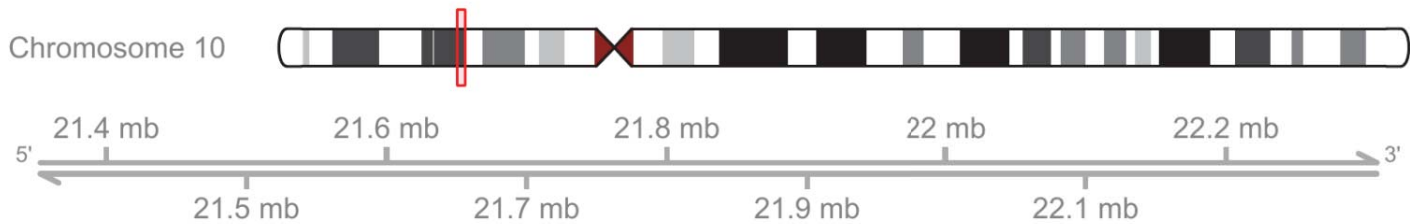
Uncorrected



Zscore of the effect allele: C

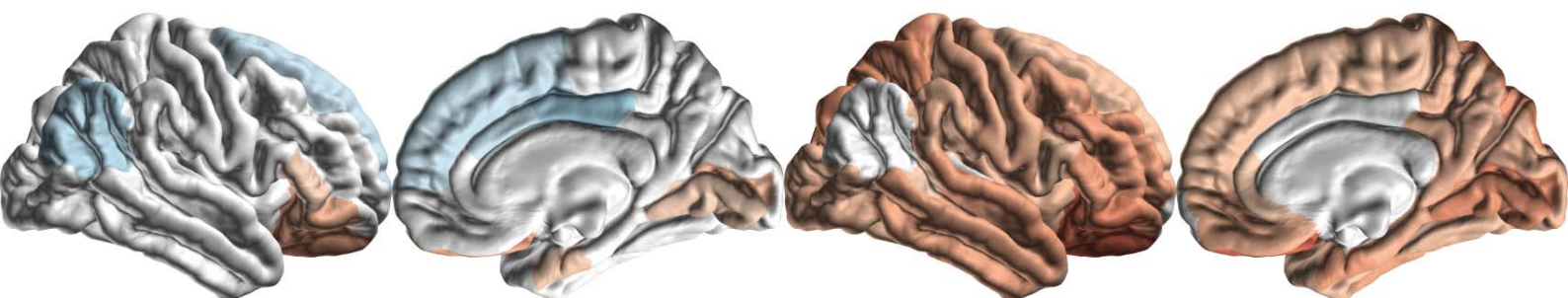
-15 -10 -5 0 5 10 15

Total Surface Area SA: rs11012732



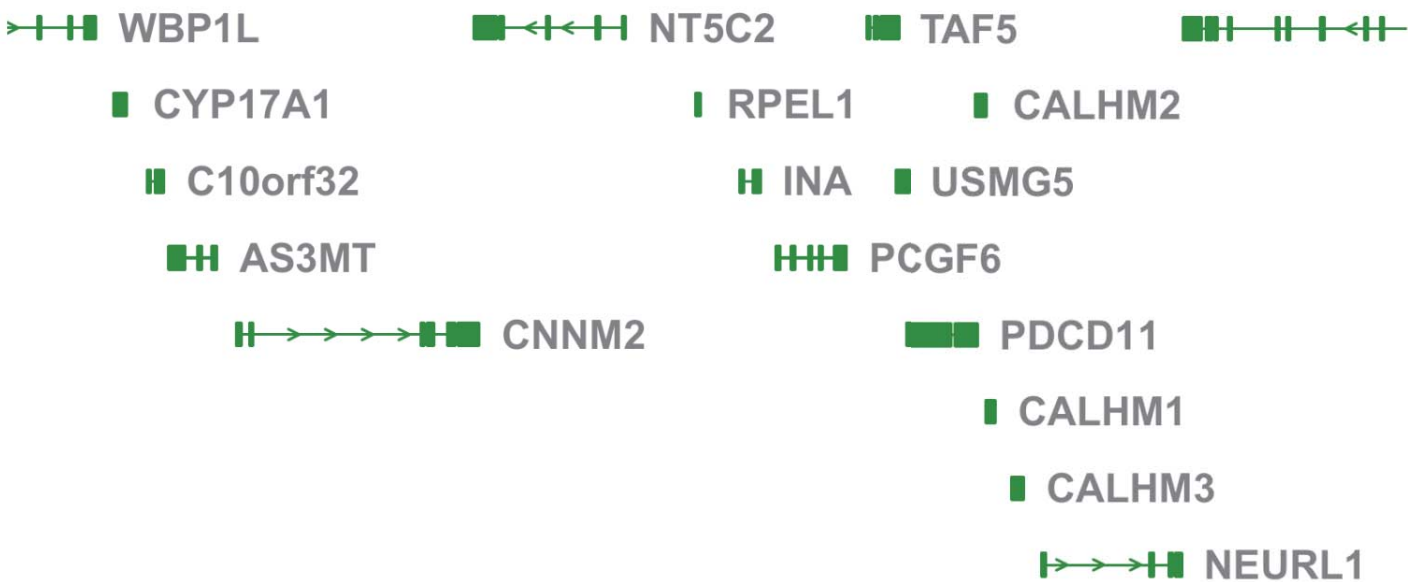
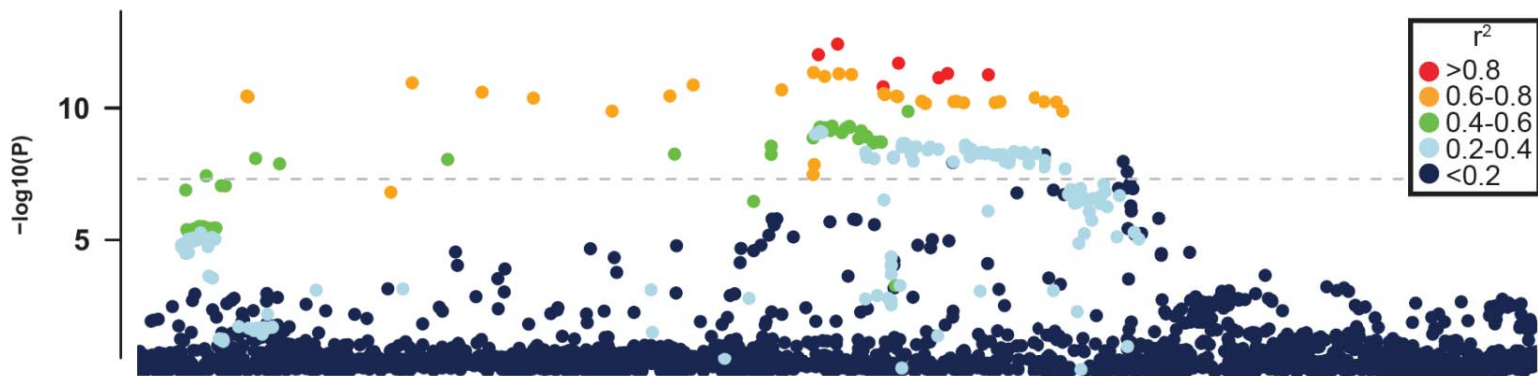
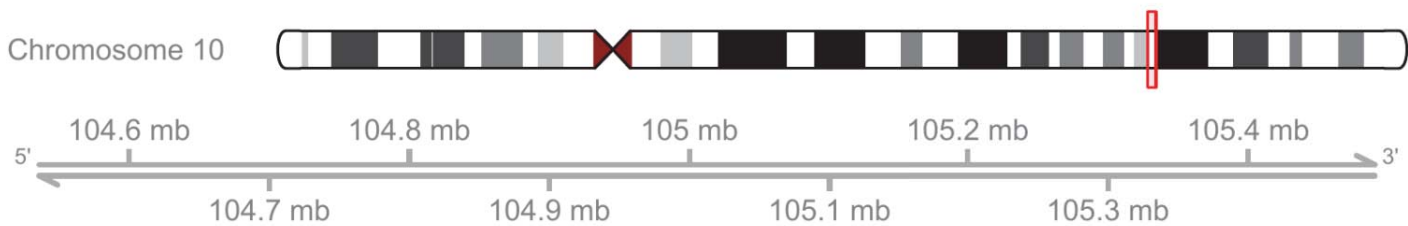
Corrected for total SA

Uncorrected



-15 -10 -5 0 5 10 15

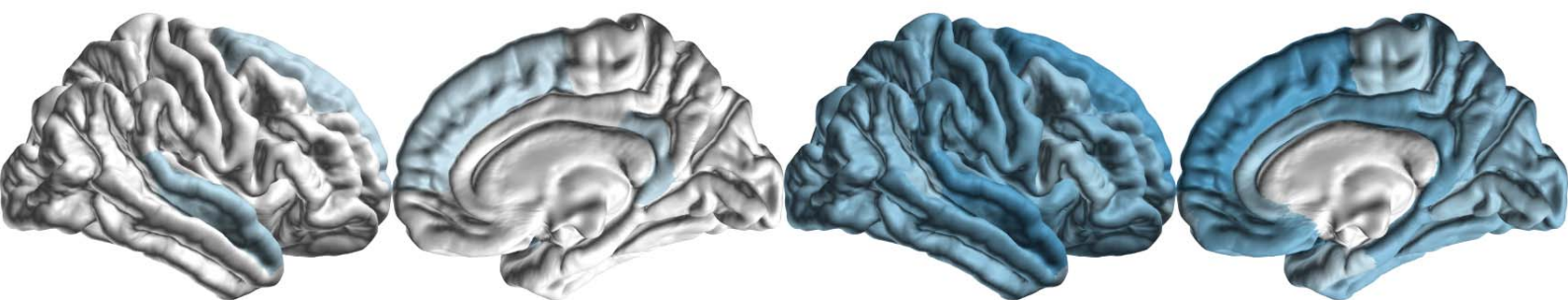
Total Surface Area SA: rs1628768



CP>GZ GZ>CP

Corrected for total SA

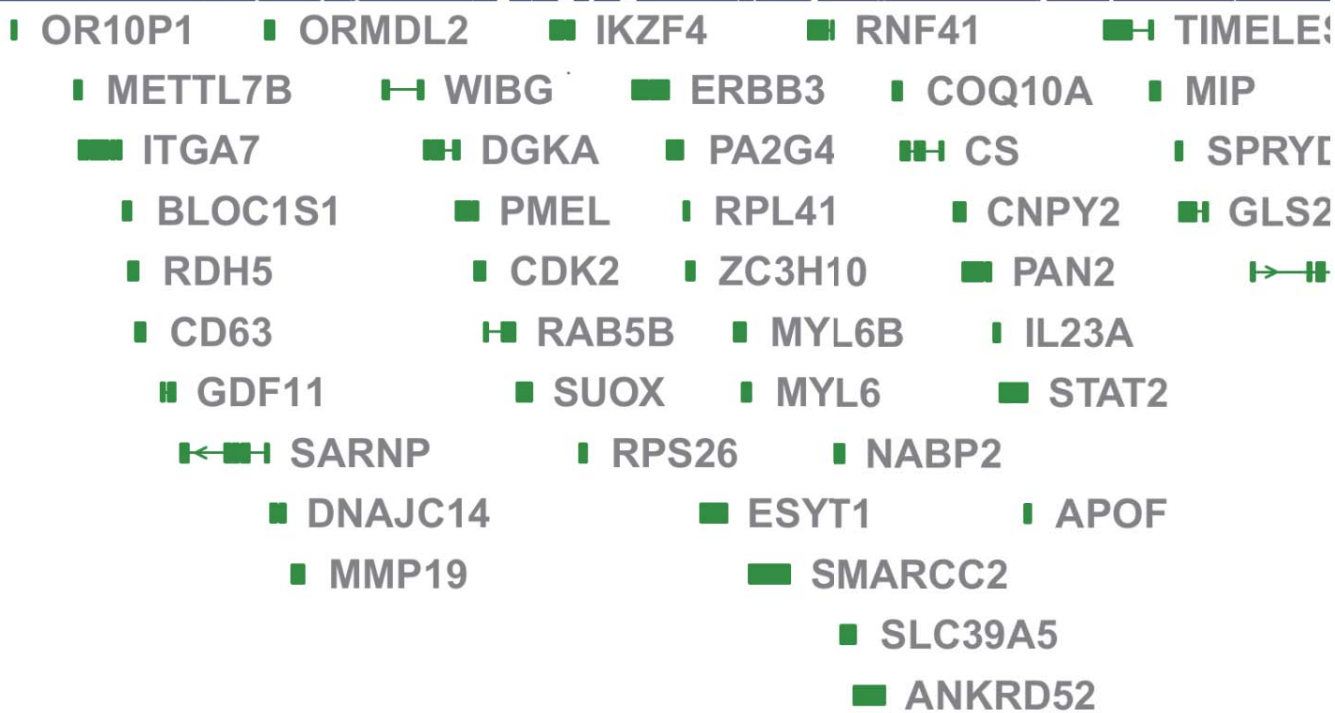
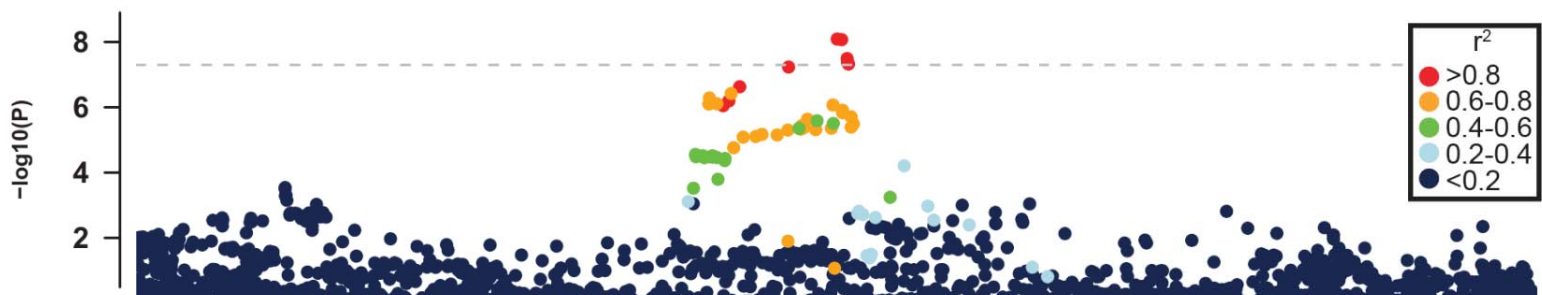
Uncorrected



Zscore of the effect allele: T

-15 -10 -5 0 5 10 15

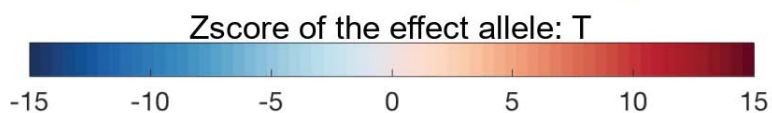
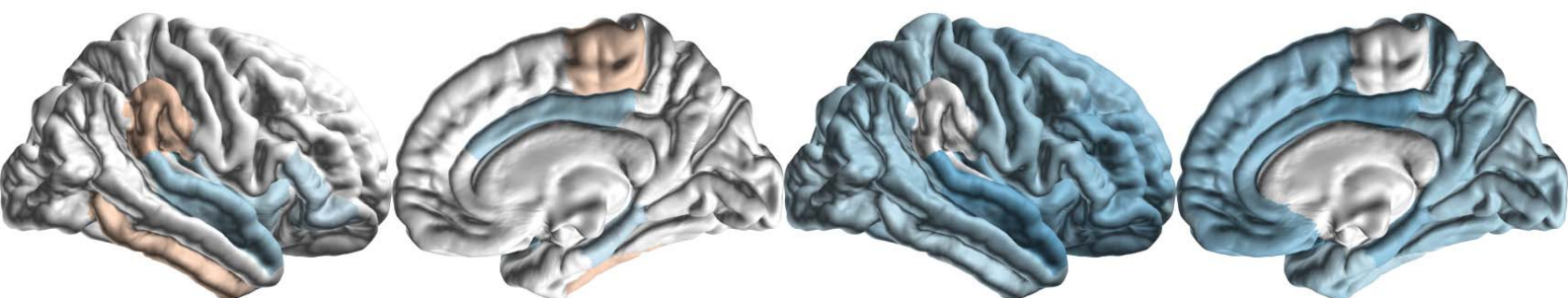
Total Surface Area SA: rs11171739



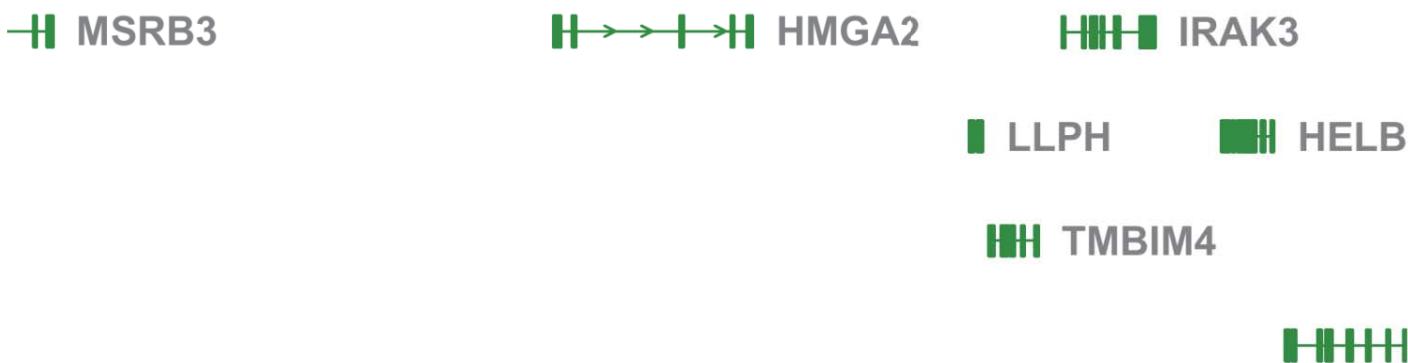
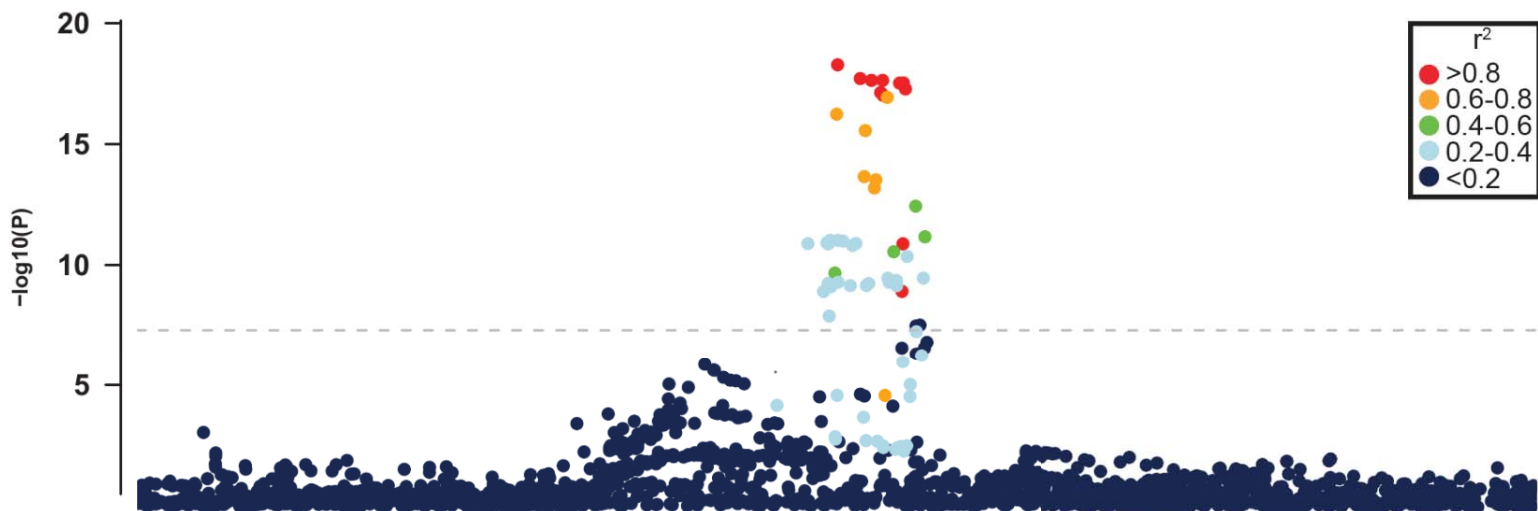
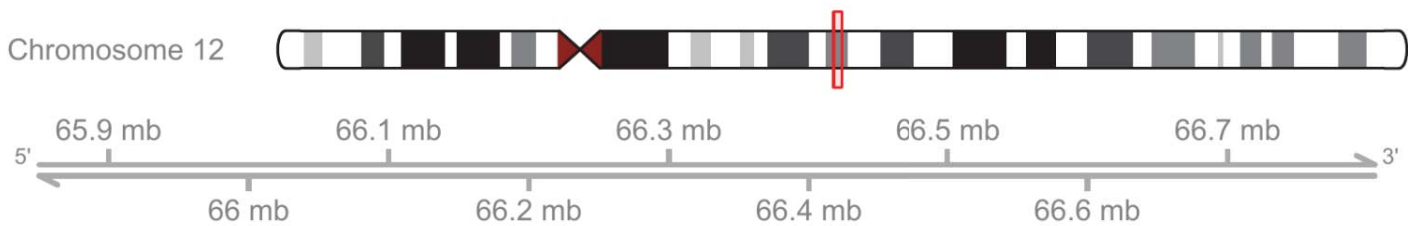
CP>GZZ>CP

Corrected for total SA

Uncorrected

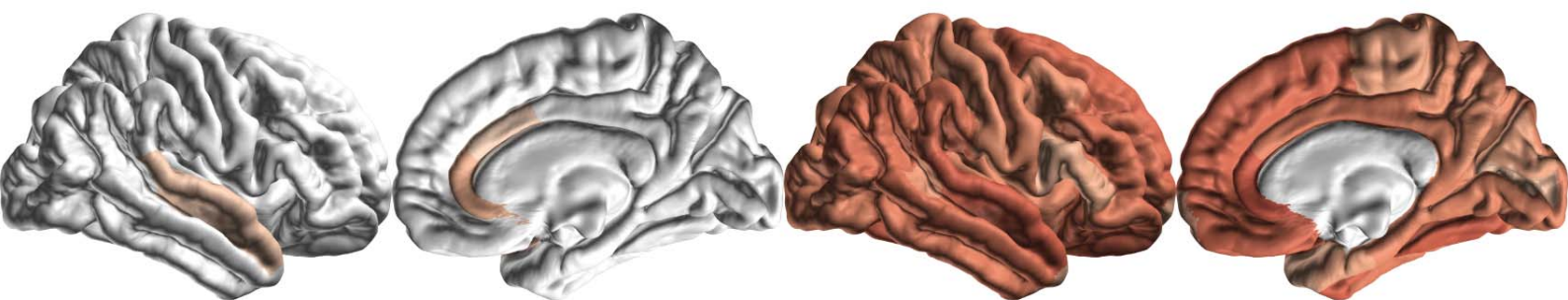


Total Surface Area SA: rs10878349



Corrected for total SA

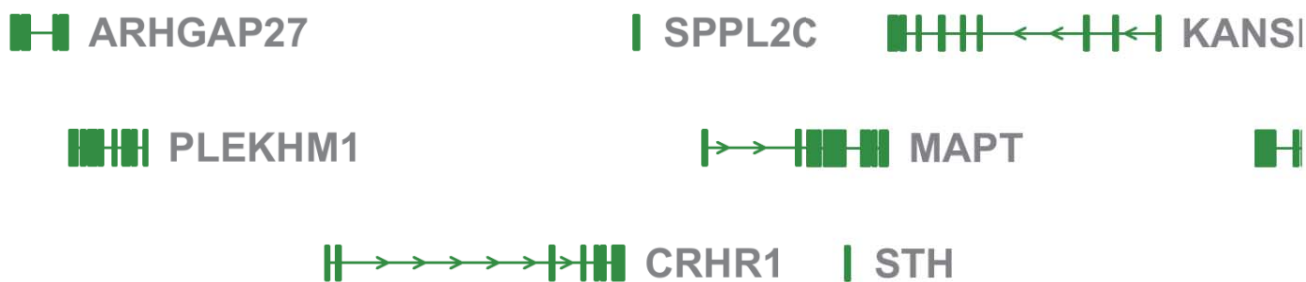
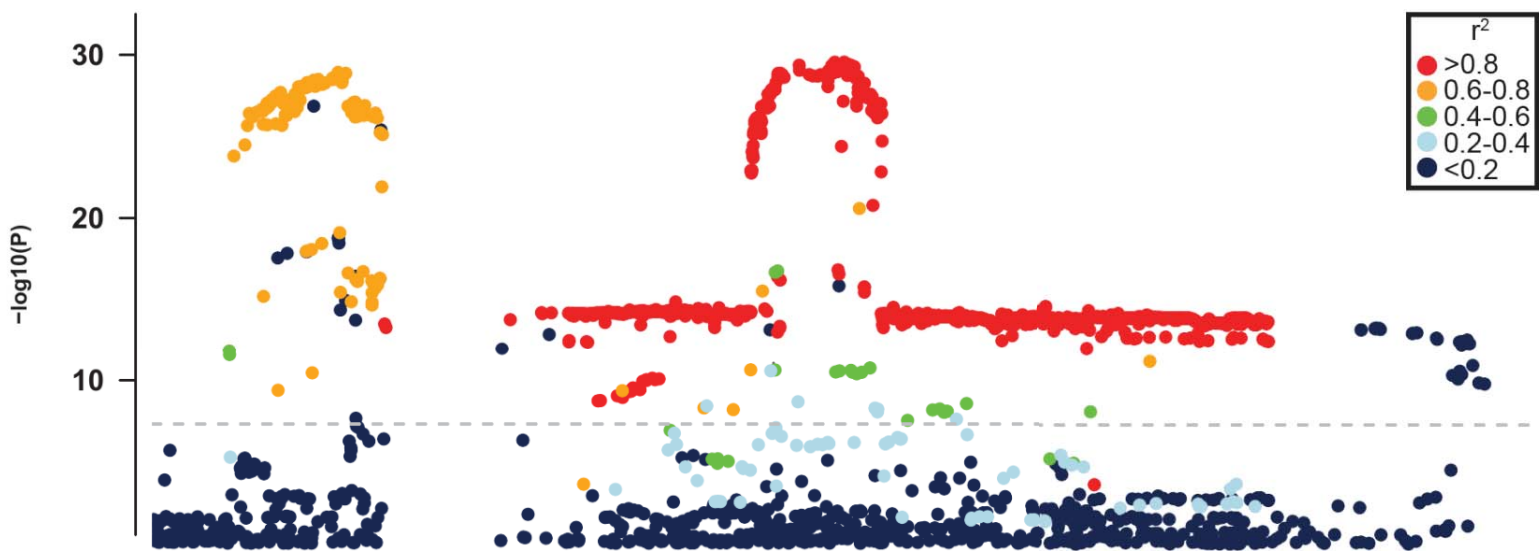
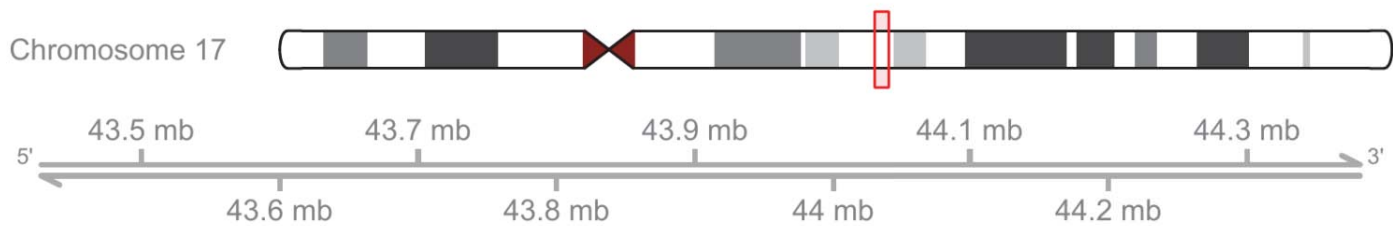
Uncorrected



Zscore of the effect allele: A

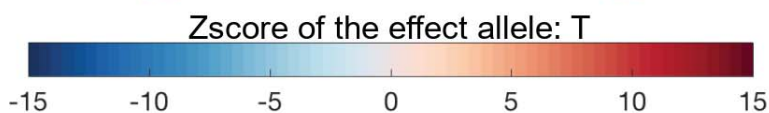
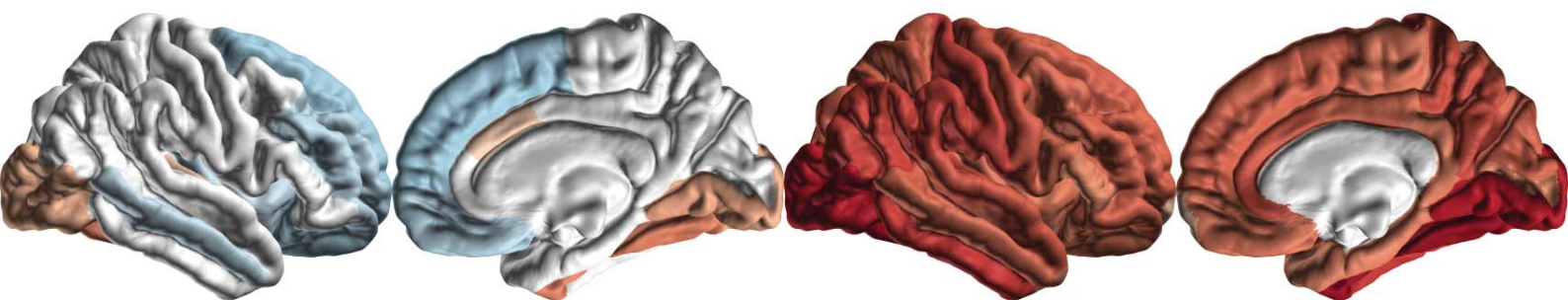


Total Surface Area SA: rs62057153

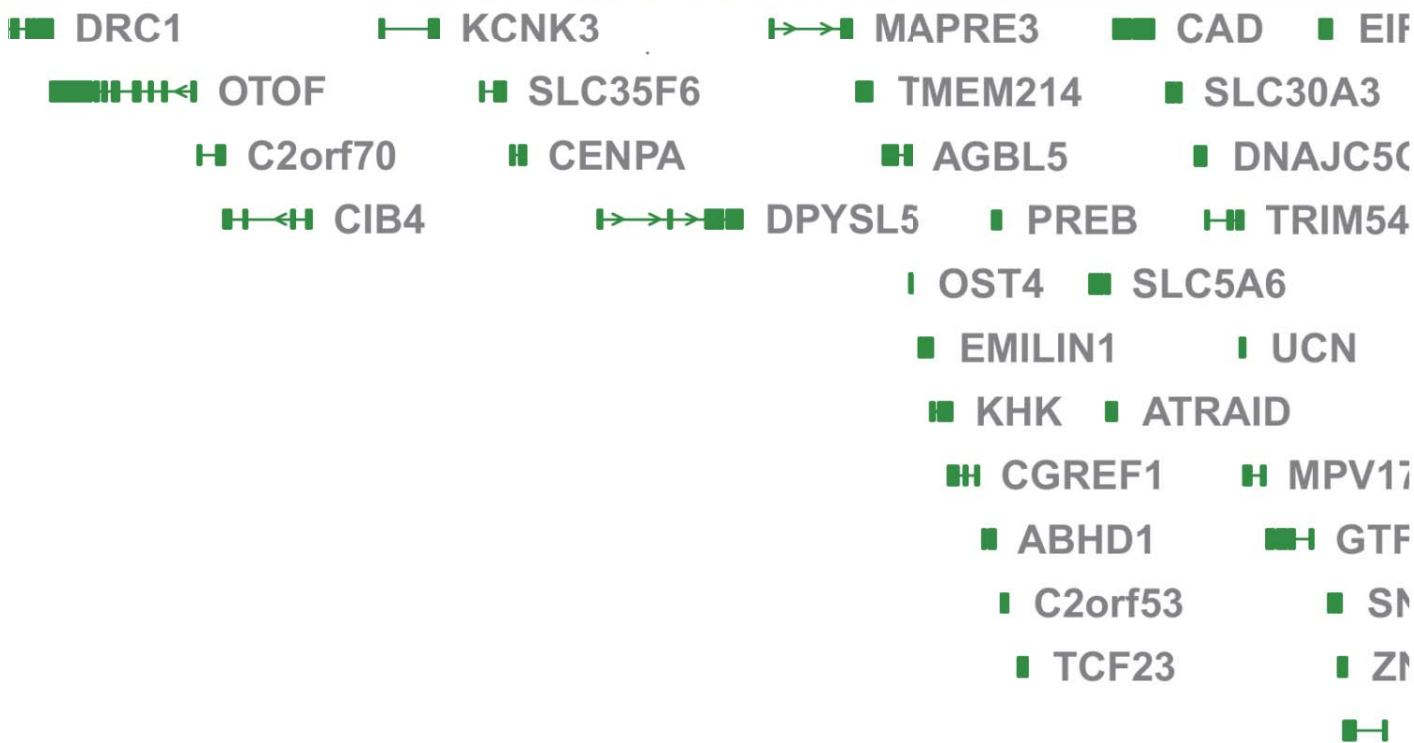
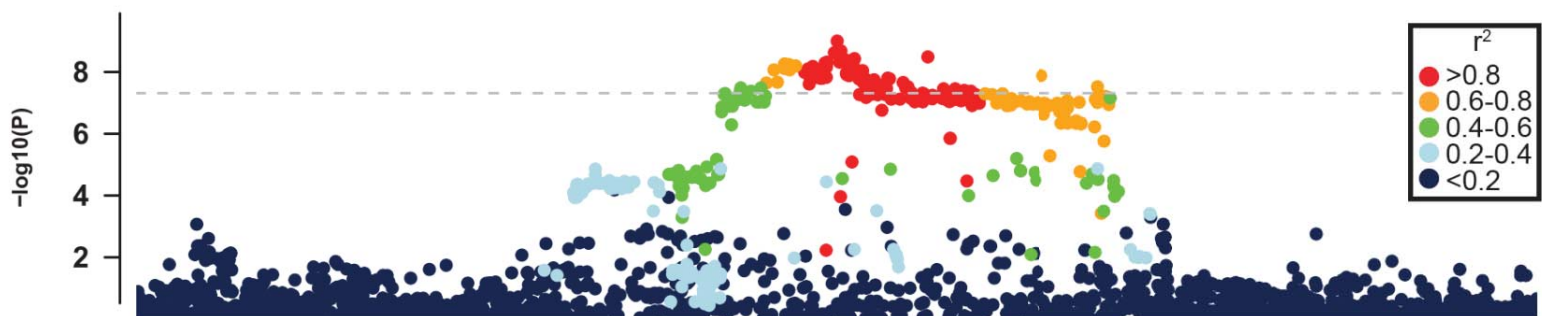
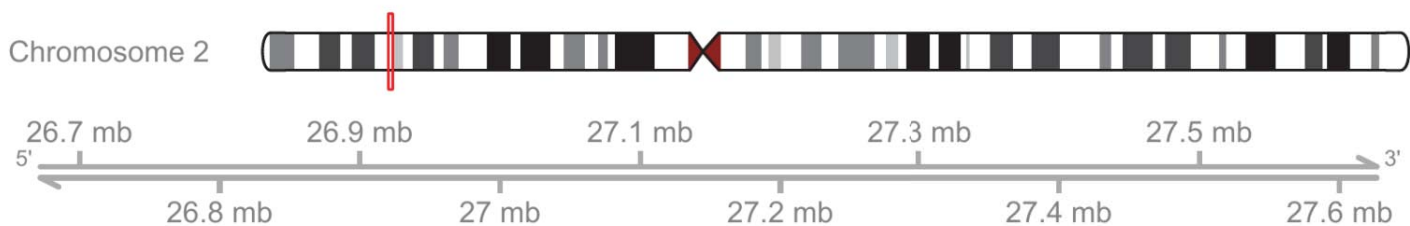


Corrected for total SA

Uncorrected



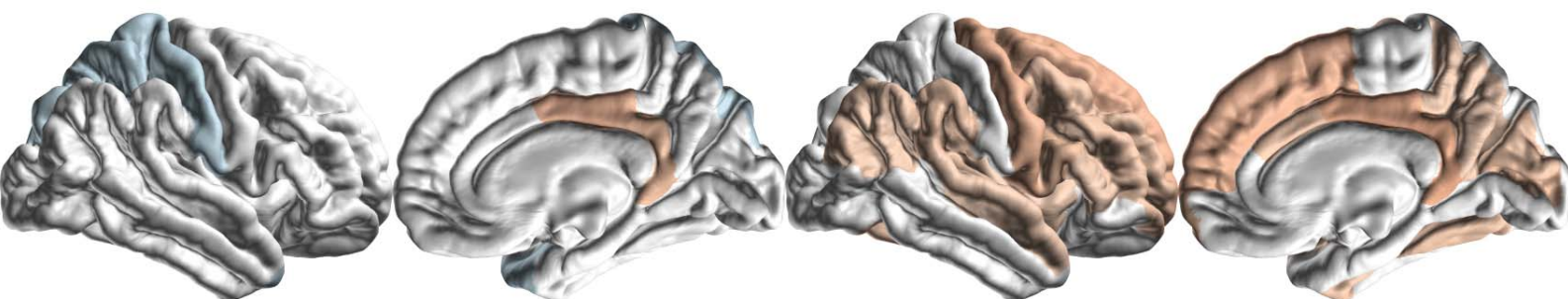
Average Thickness TH: rs6738528



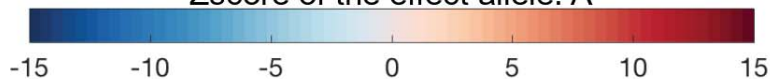
dIPFC

Corrected for average TH

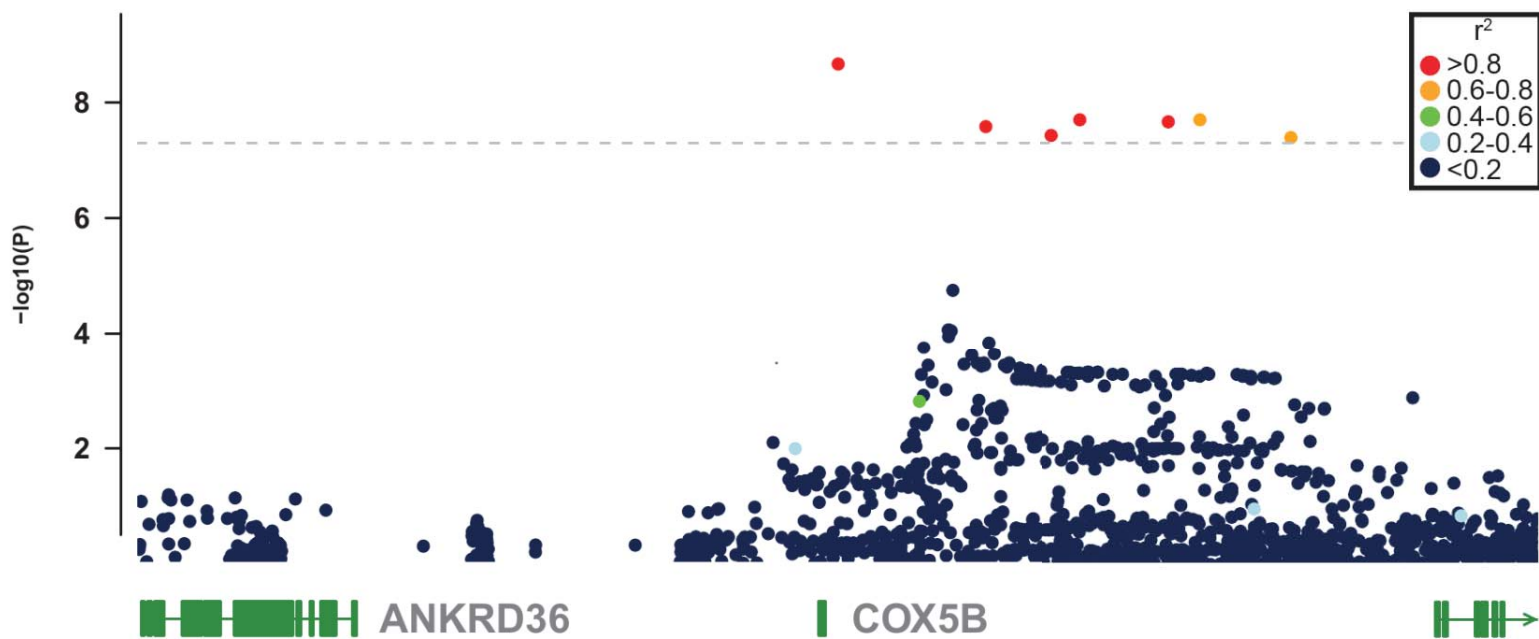
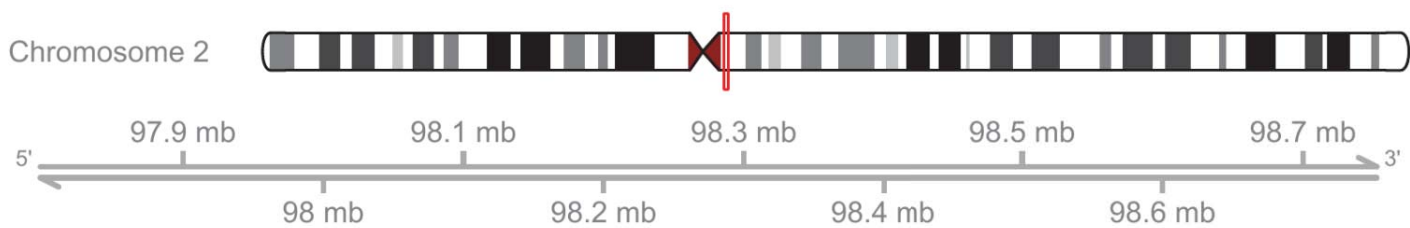
Uncorrected



Zscore of the effect allele: A



Average Thickness TH: rs11692435



ACTR1B

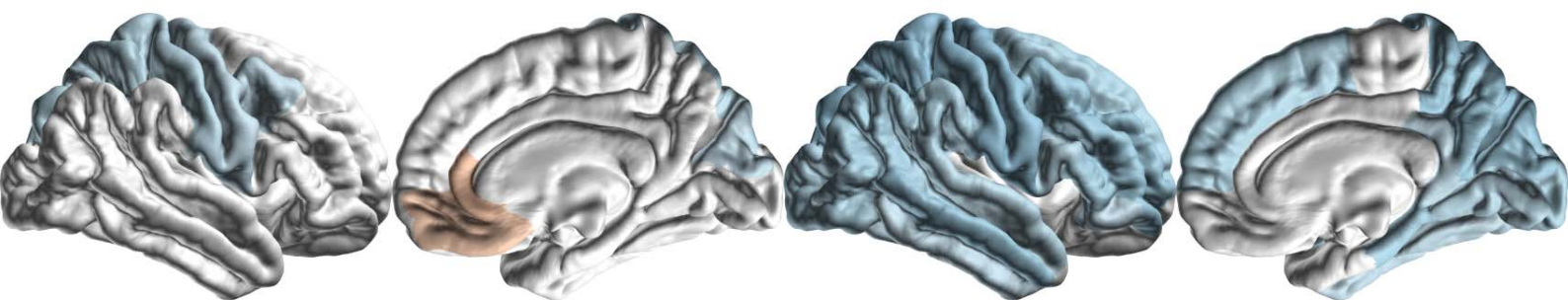
ZAP70

TMEM131

dIPFC

Corrected for average TH

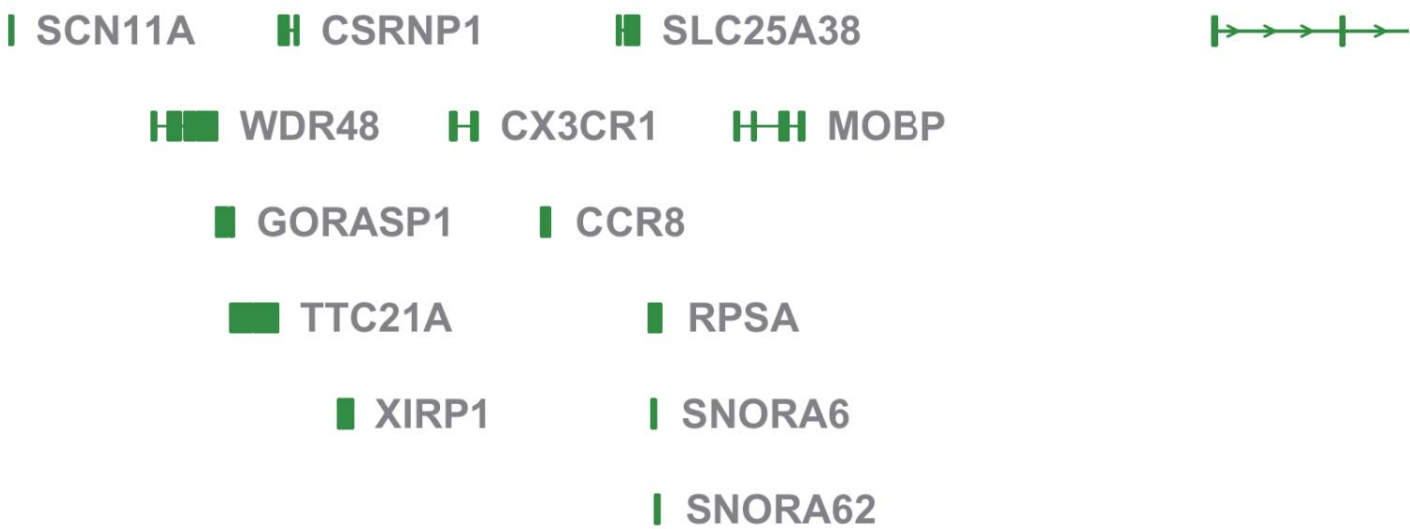
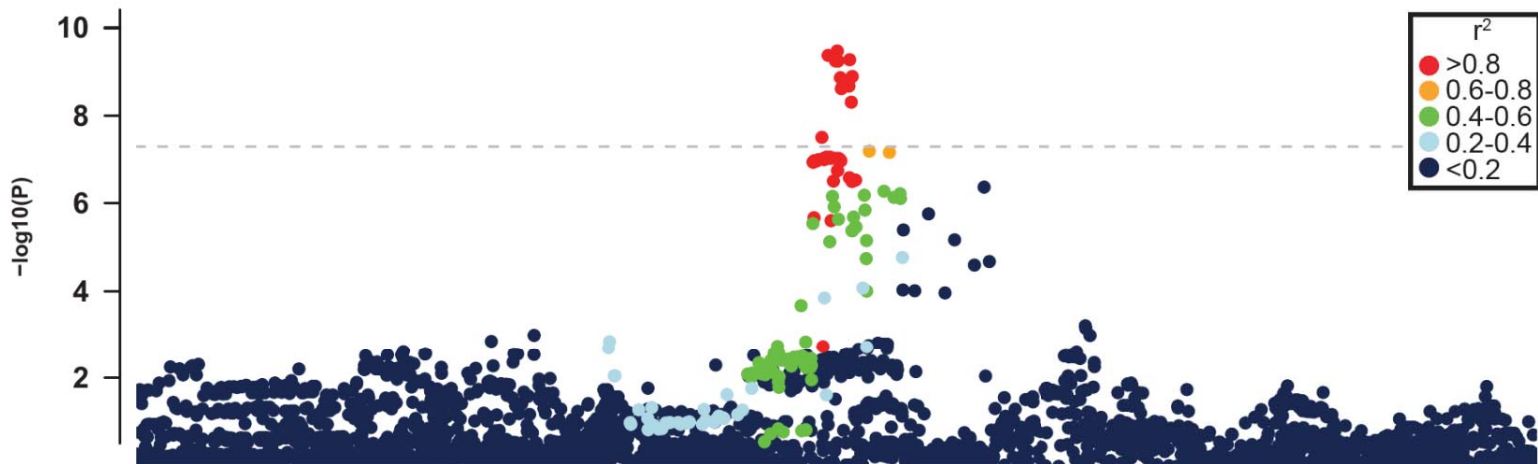
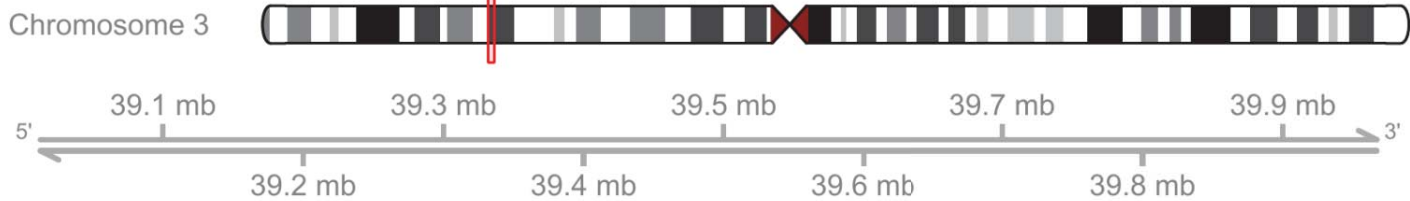
Uncorrected



Zscore of the effect allele: A

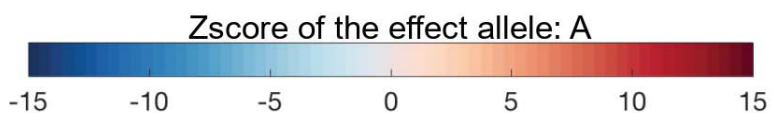
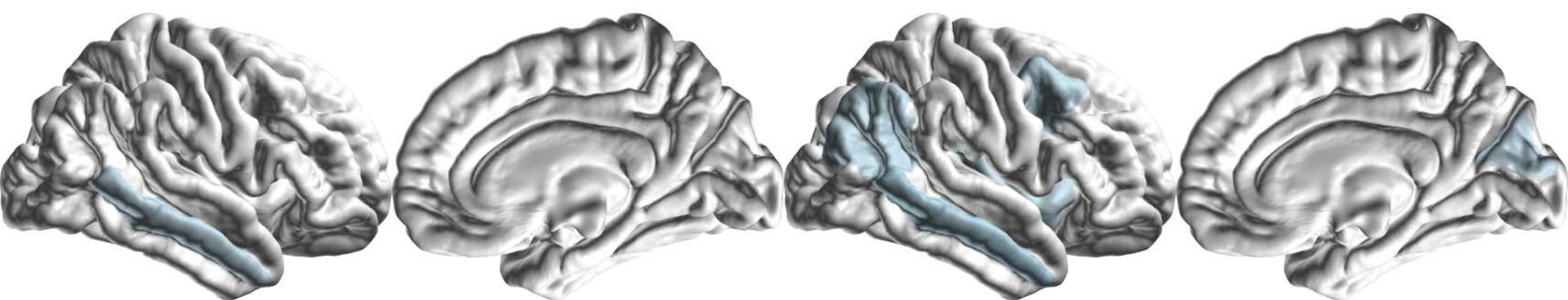


Average Thickness TH: rs630934

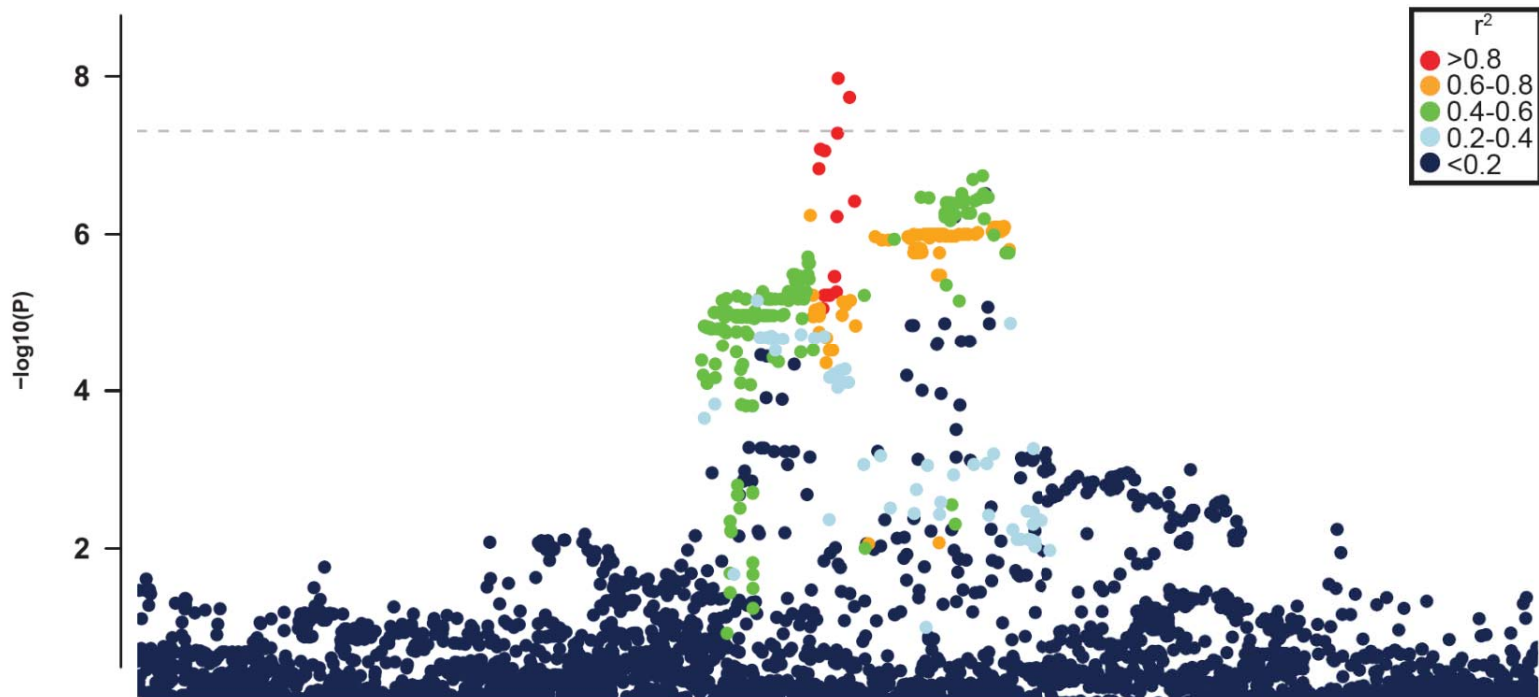
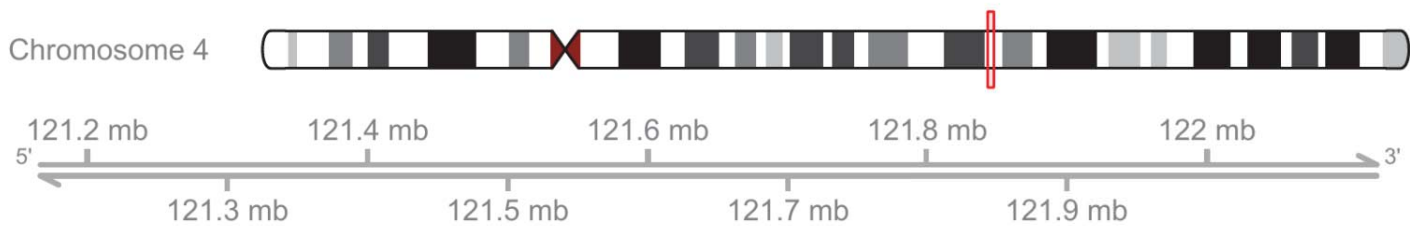


Corrected for average TH

Uncorrected



Average Thickness TH: rs13110077



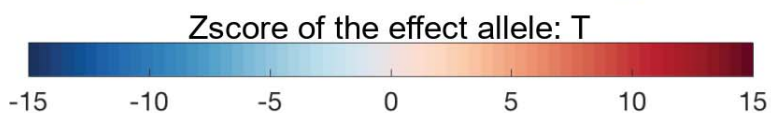
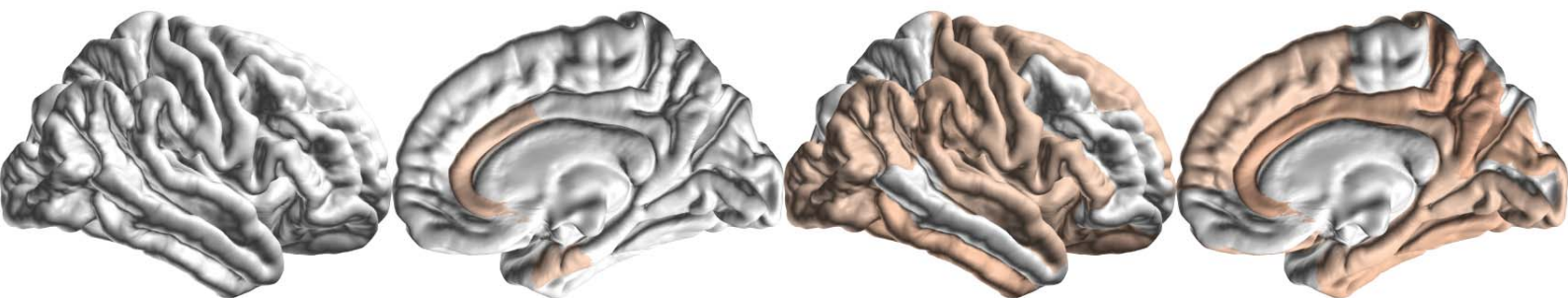
PRDM5 NDNF

NDNF

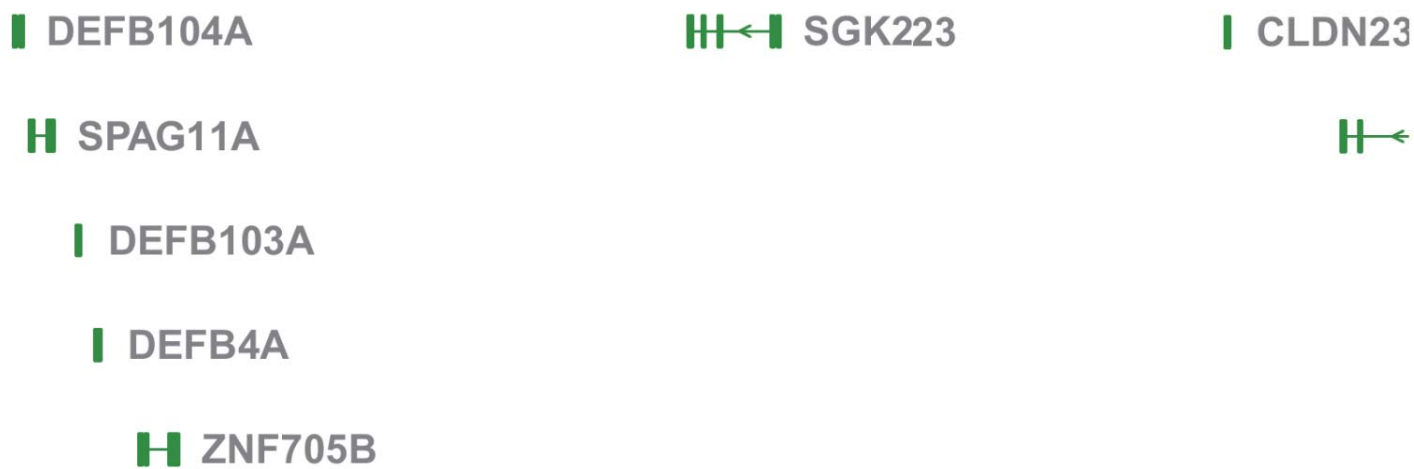
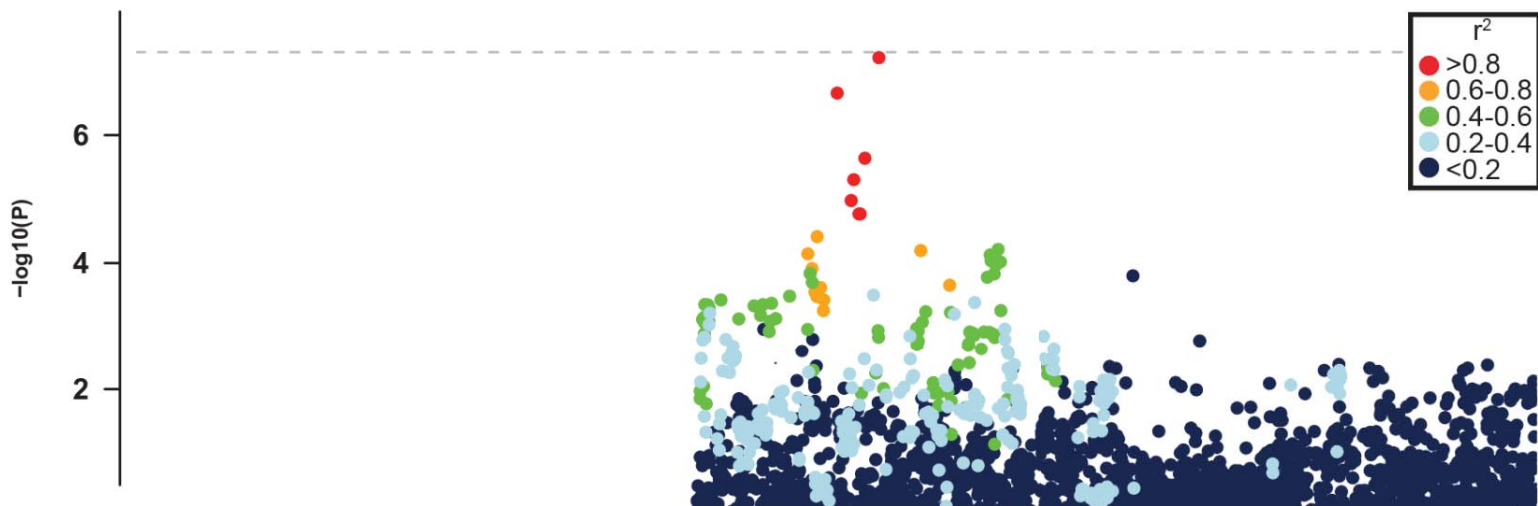
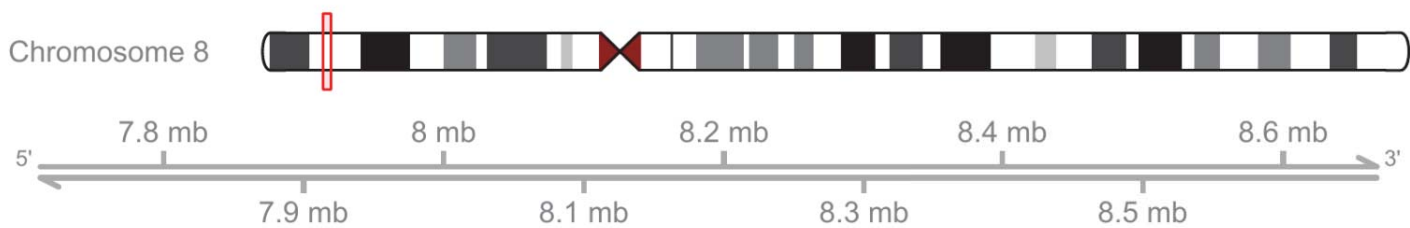
dIPFC

Corrected for average TH

Uncorrected



Average Thickness TH: rs2980497

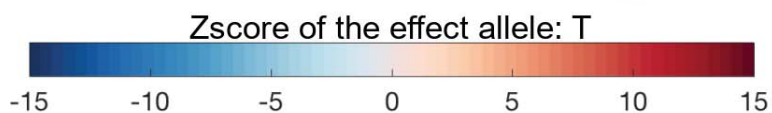
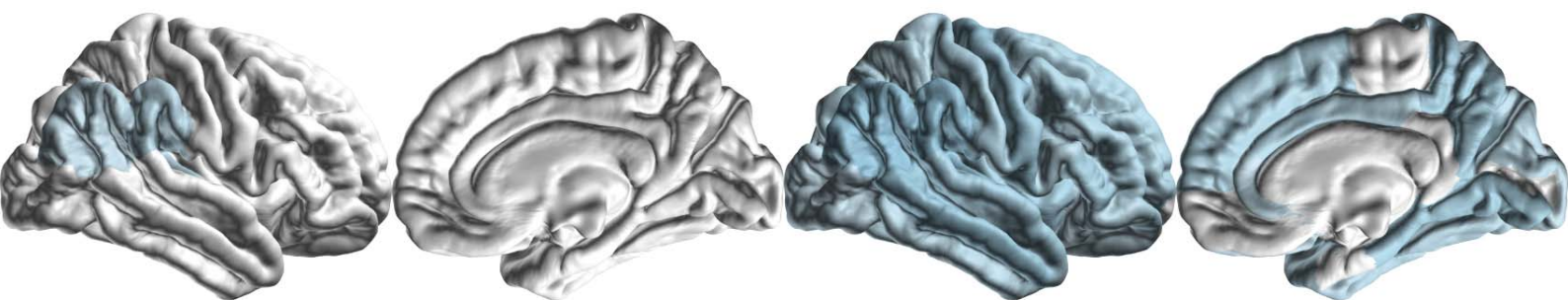


dIPFC

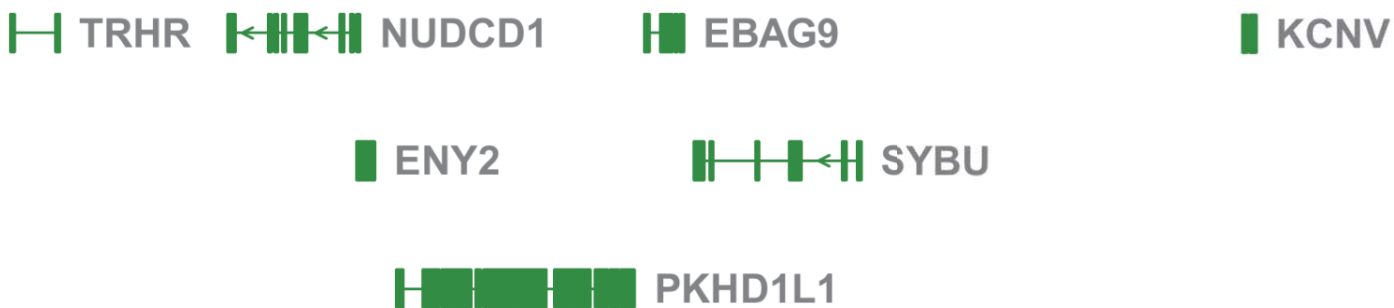
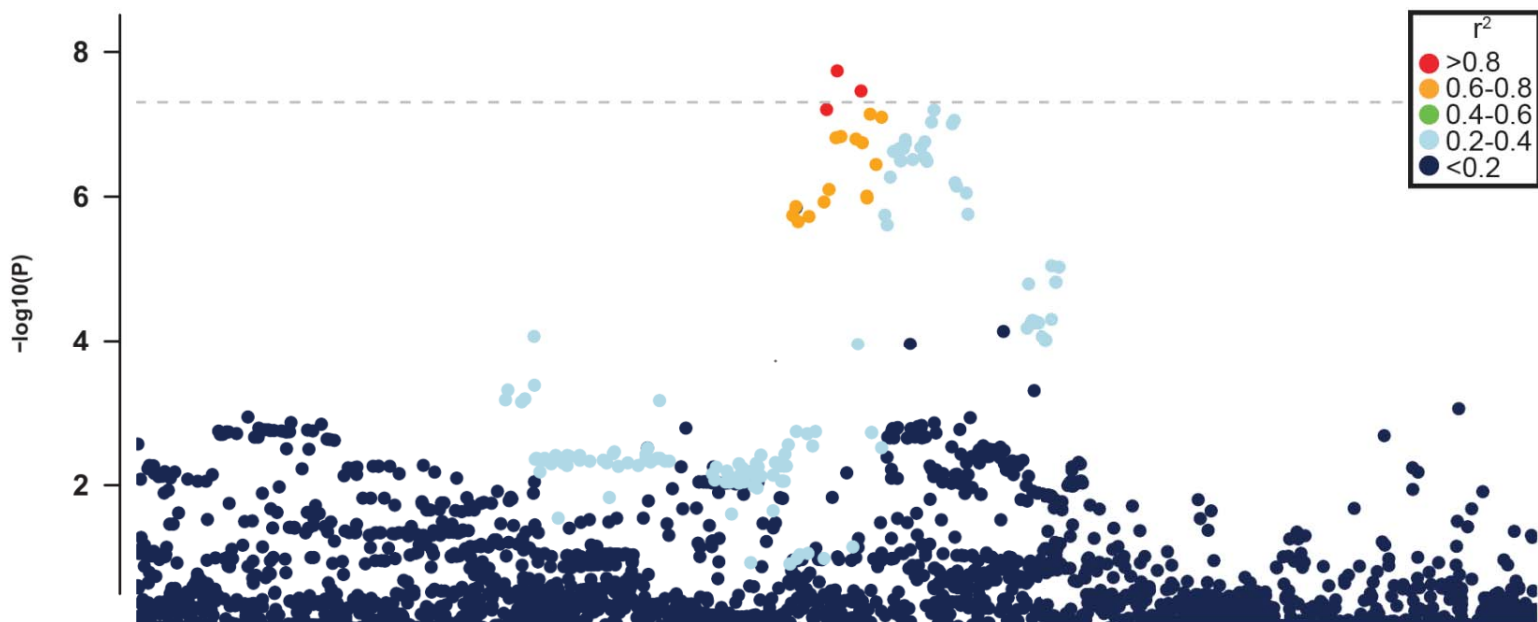
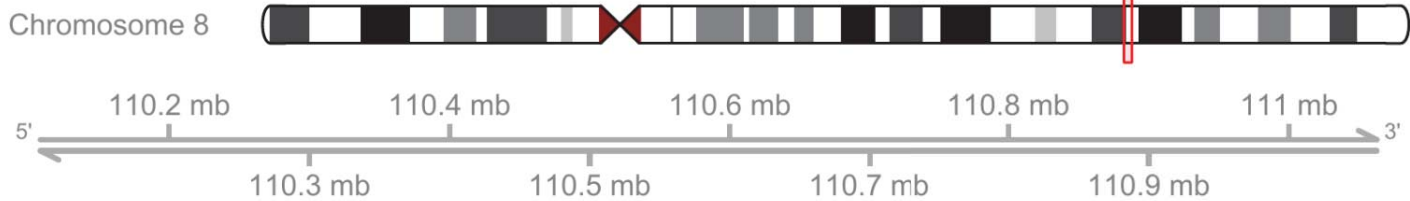


Corrected for average TH

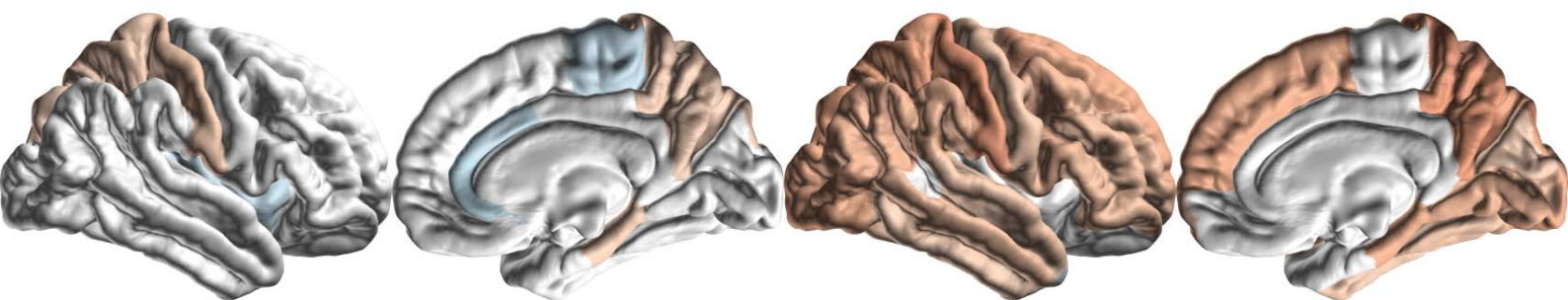
Uncorrected



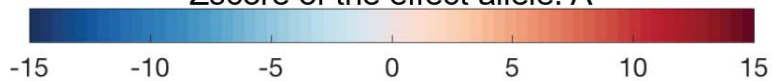
Average Thickness TH: rs7824177



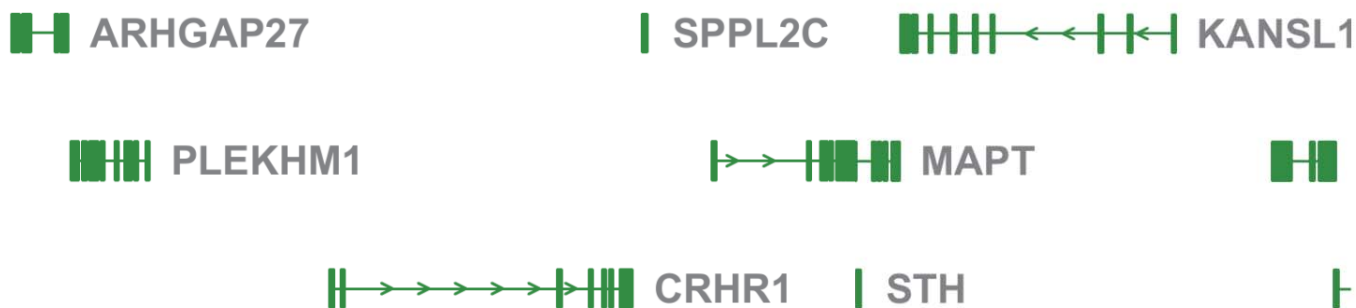
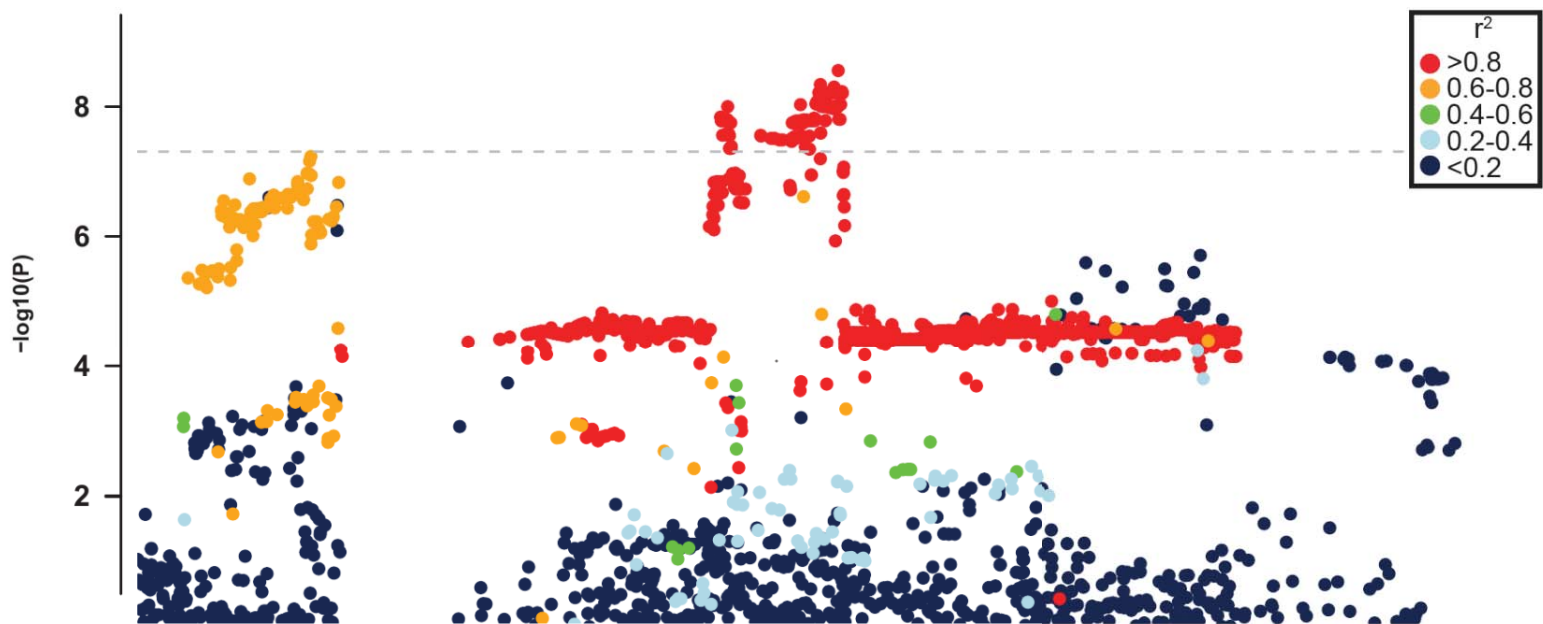
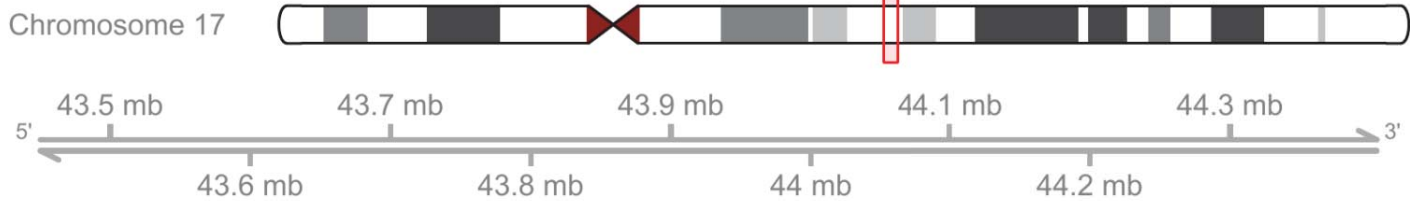
dIPFC



Zscore of the effect allele: A

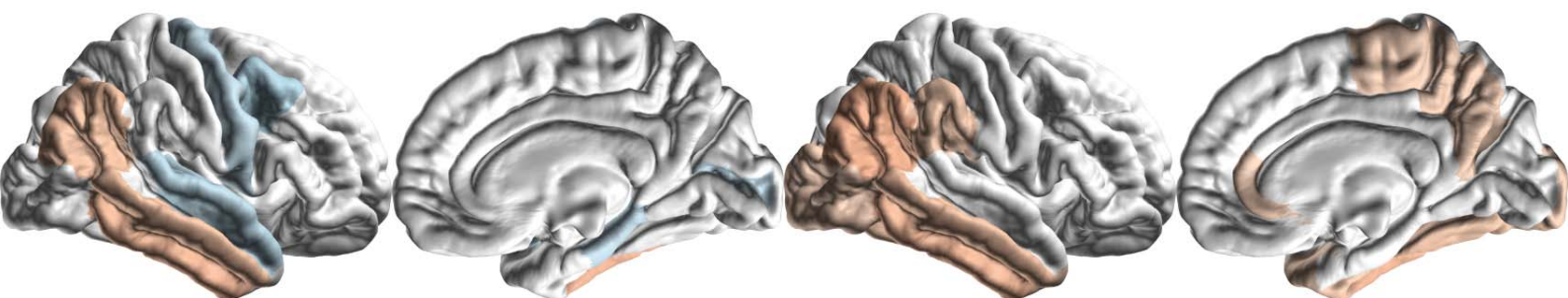


Average Thickness TH: rs17769552



Corrected for average TH

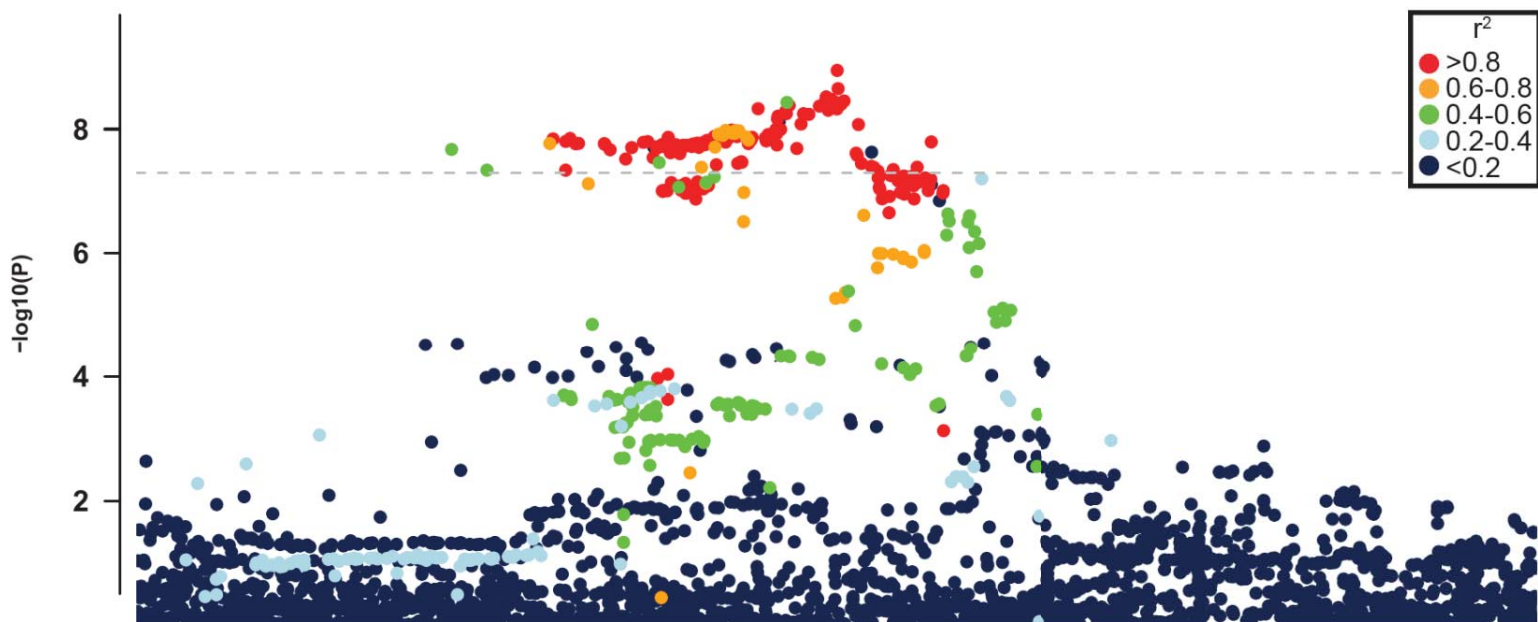
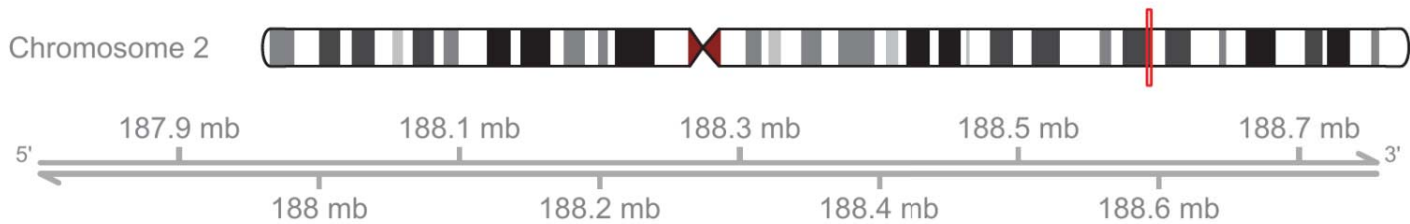
Uncorrected



Zscore of the effect allele: A



Frontal Pole SA: rs17464221



 CALCRL

 TFPI

GZ>CP

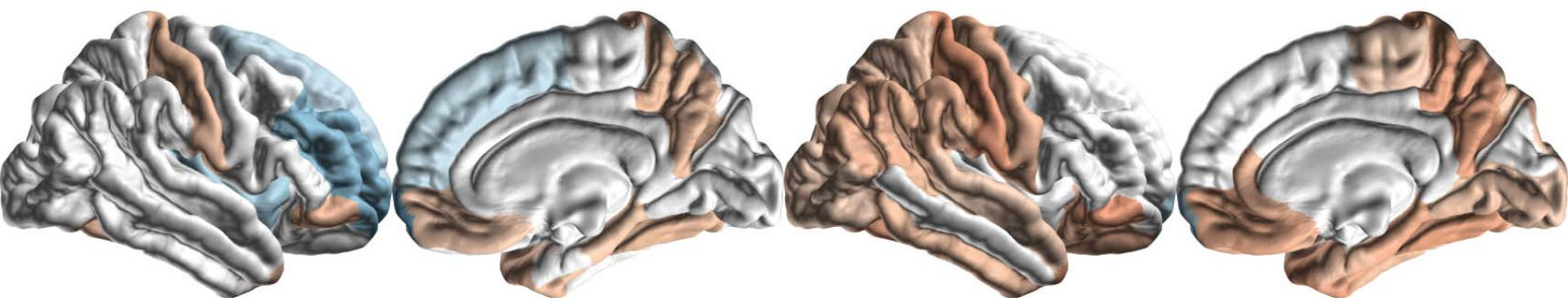


CP>GZ

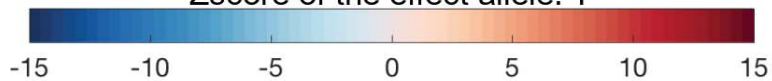


Corrected for total SA

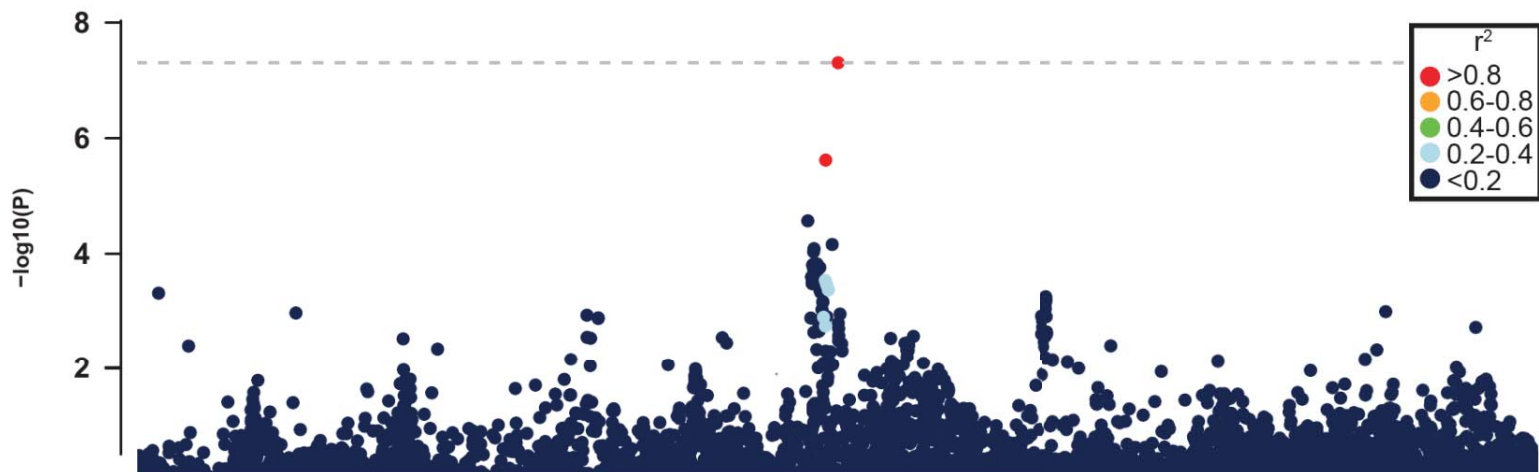
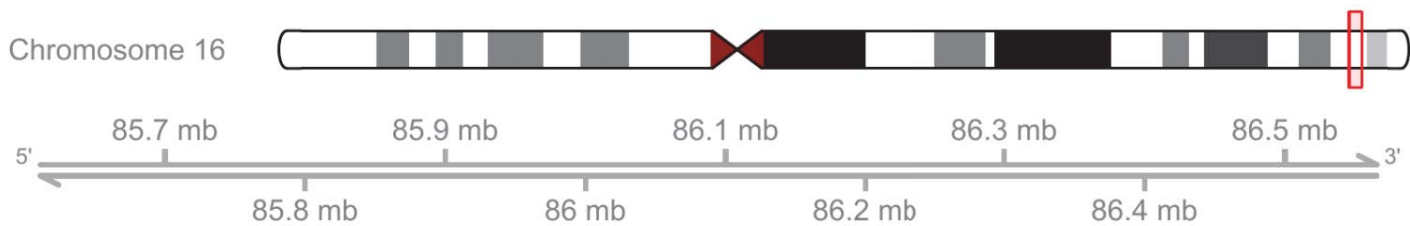
Uncorrected



Zscore of the effect allele: T

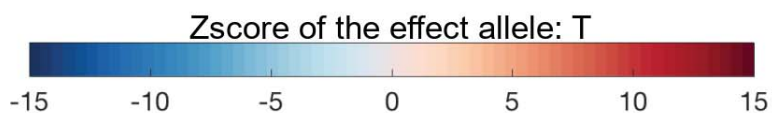
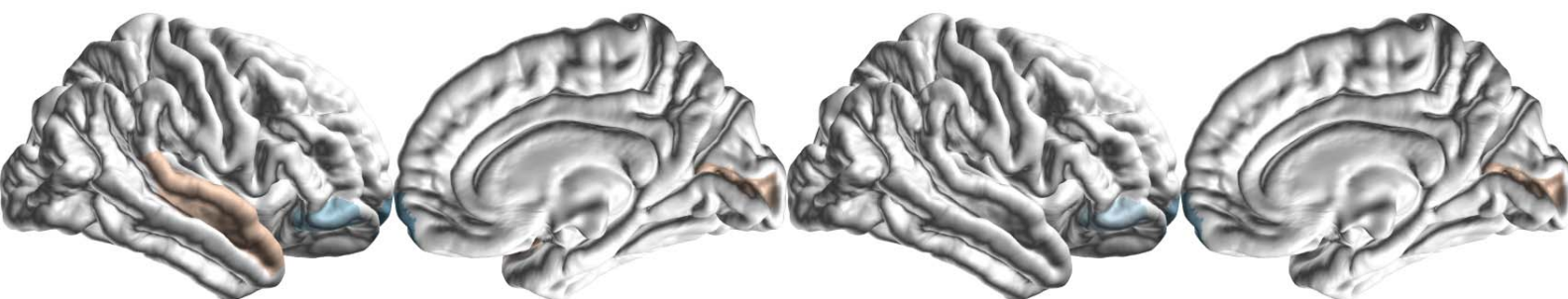


Frontal Pole SA: rs113050298

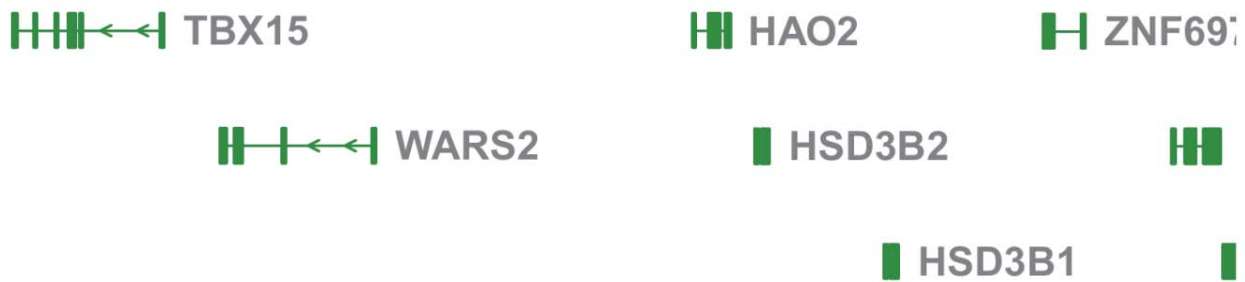
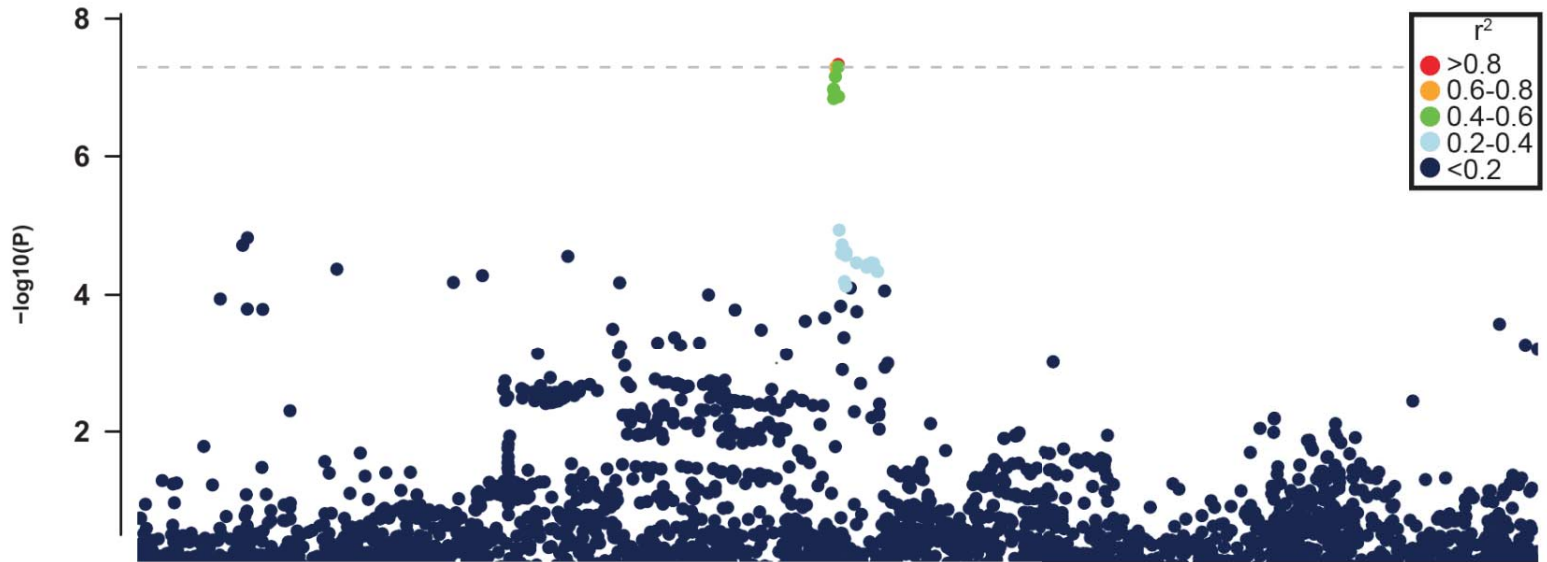
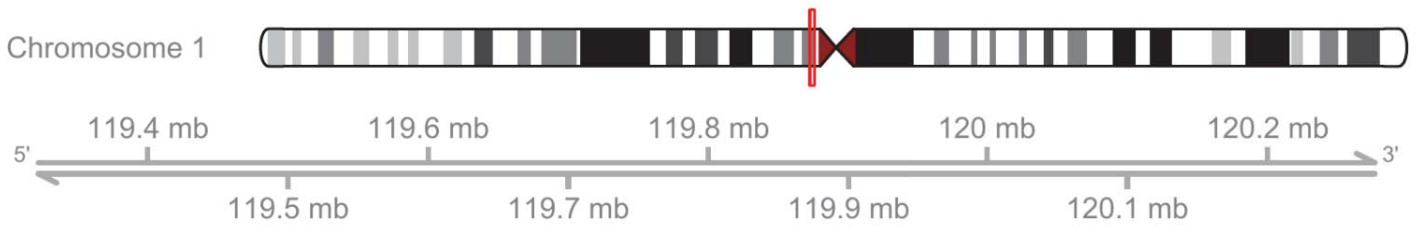


Corrected for total SA

Uncorrected



Lateral Orbitofrontal SA: rs7529542

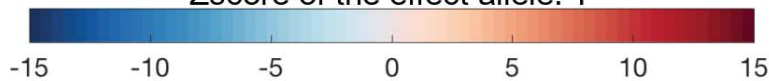


Corrected for total SA

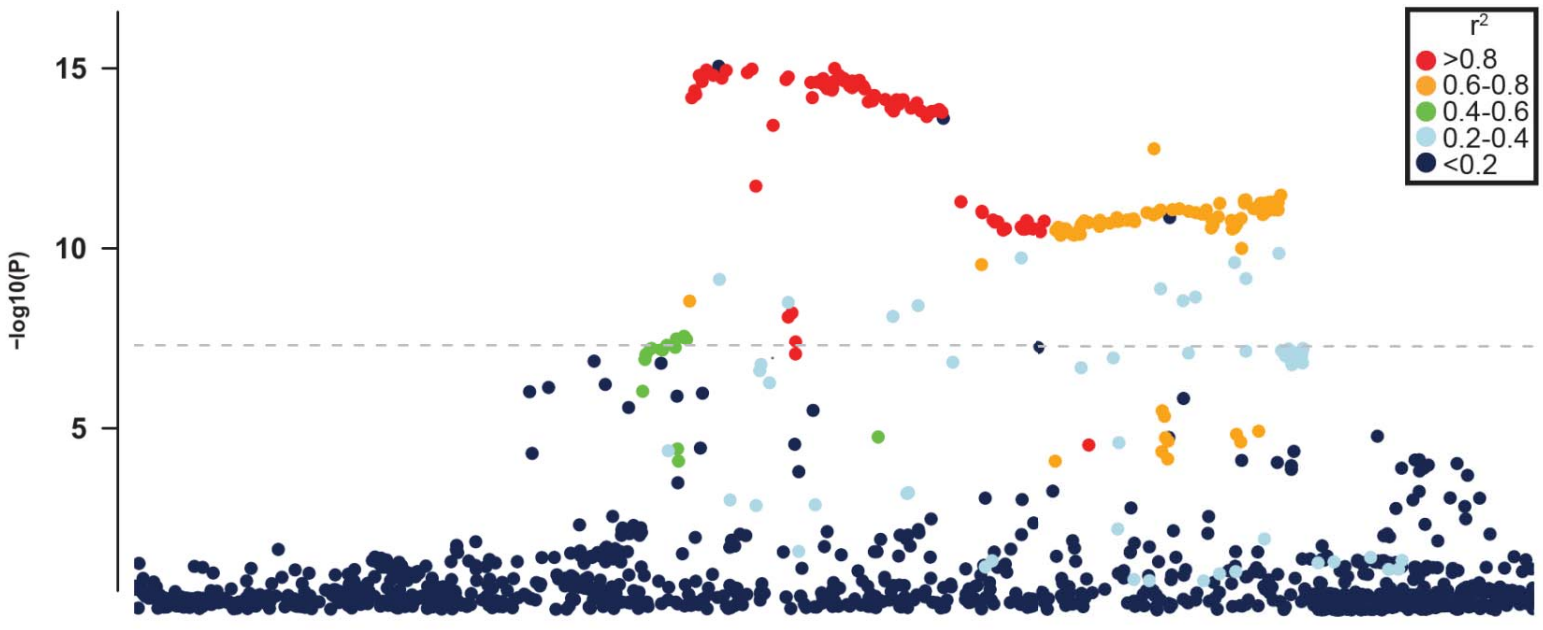
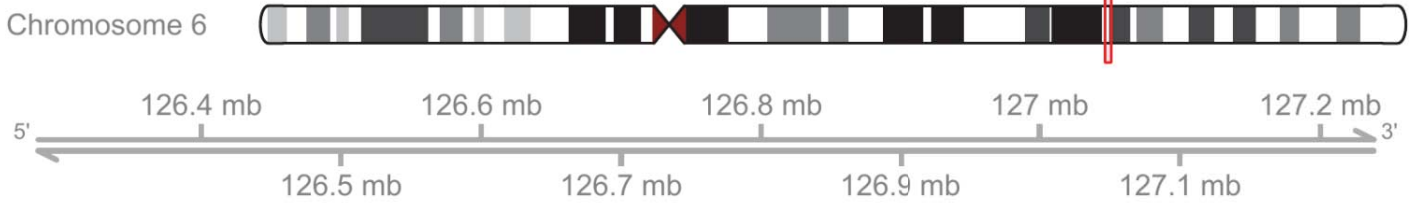
Uncorrected



Zscore of the effect allele: T



Lateral Orbitofrontal SA: rs2184968



HINT3

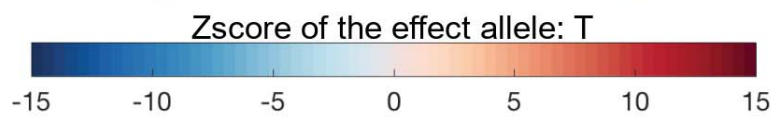
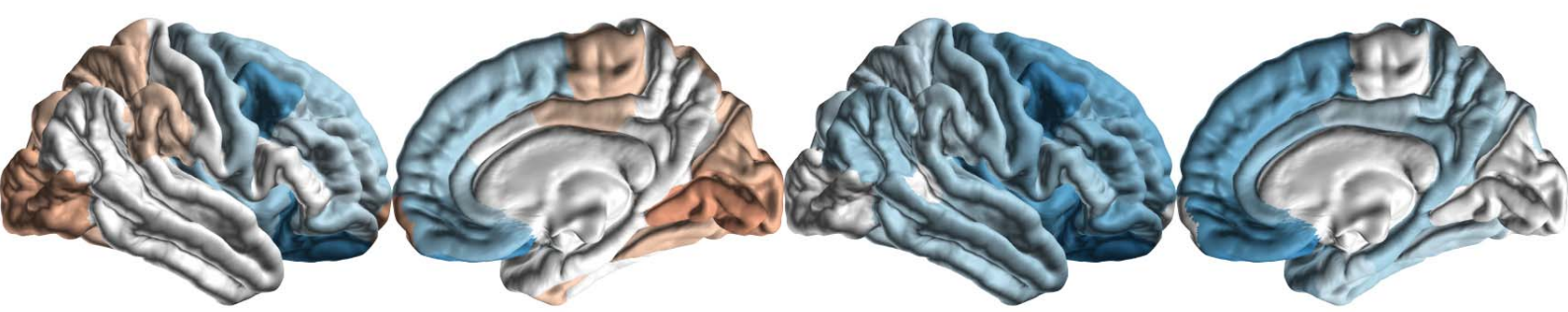
CENPW

TRMT11

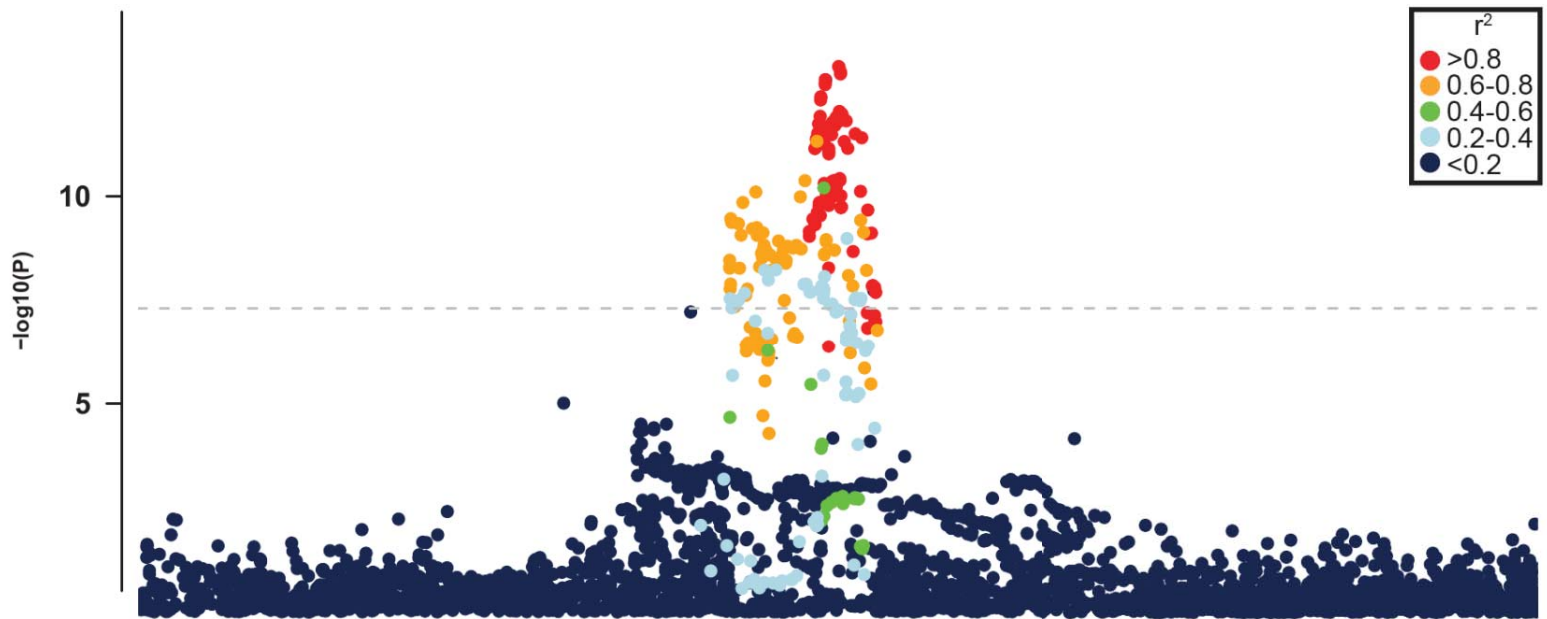
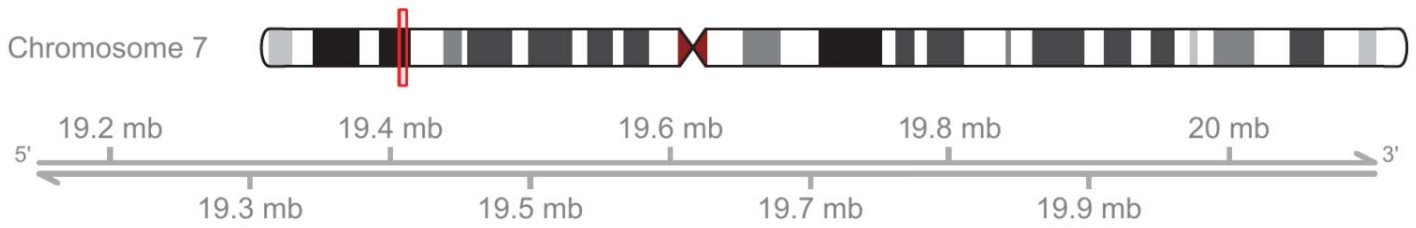


Corrected for total SA

Uncorrected



Lateral Orbitofrontal SA: rs4721802



TWIST1

TWISTNB

FERD3L

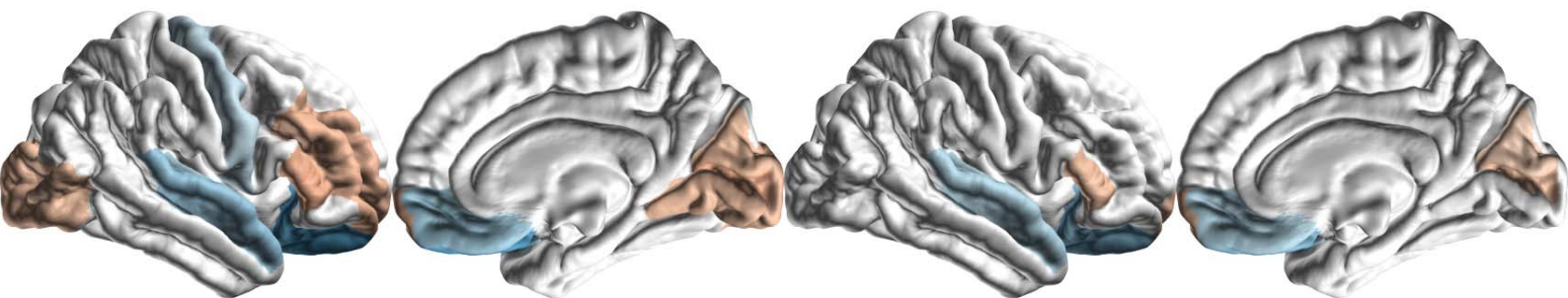
TMEM196

GZ>CP

CP>GZ

Corrected for total SA

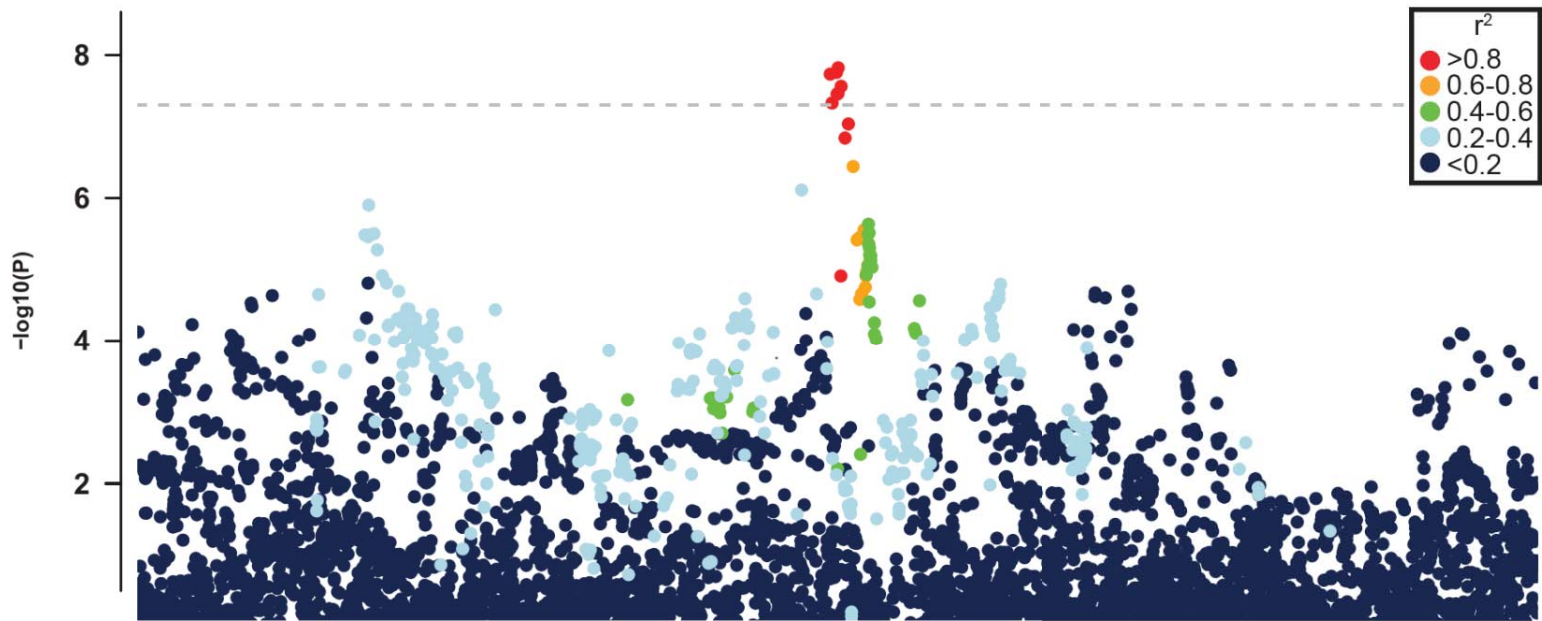
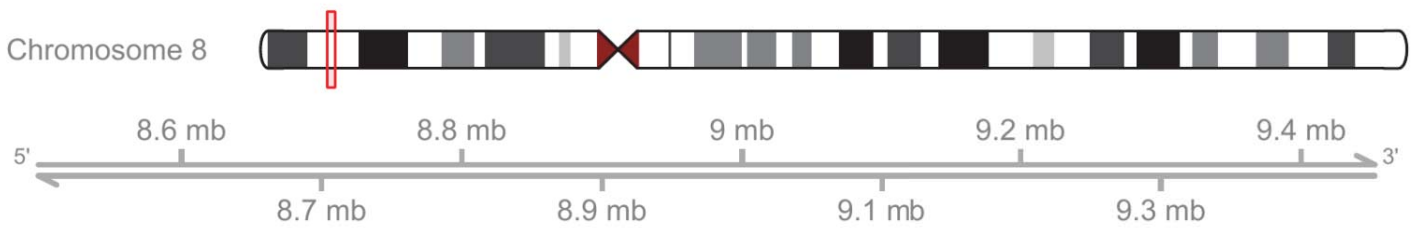
Uncorrected



Zscore of the effect allele: A



Lateral Orbitofrontal SA: rs1822951



CLDN23

ERI1

MFHAS1

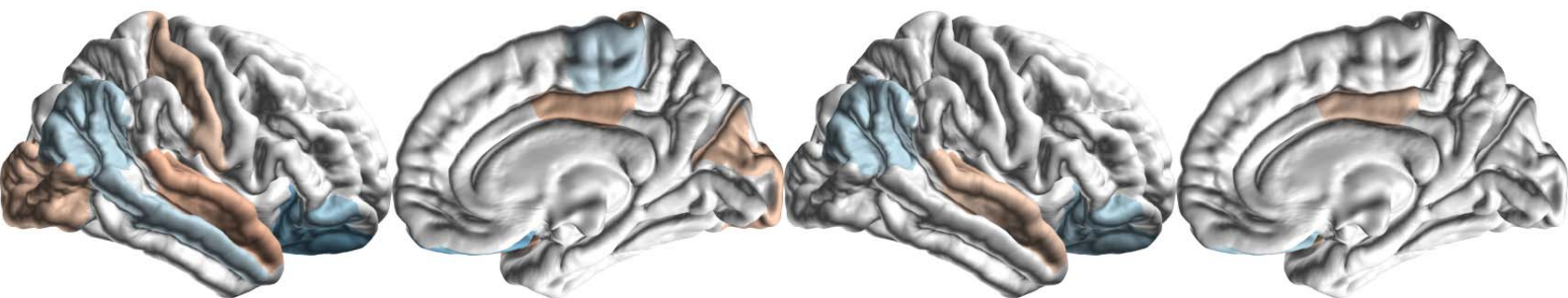
PPP1R3B

GZ>CP

CP>GZ

Corrected for total SA

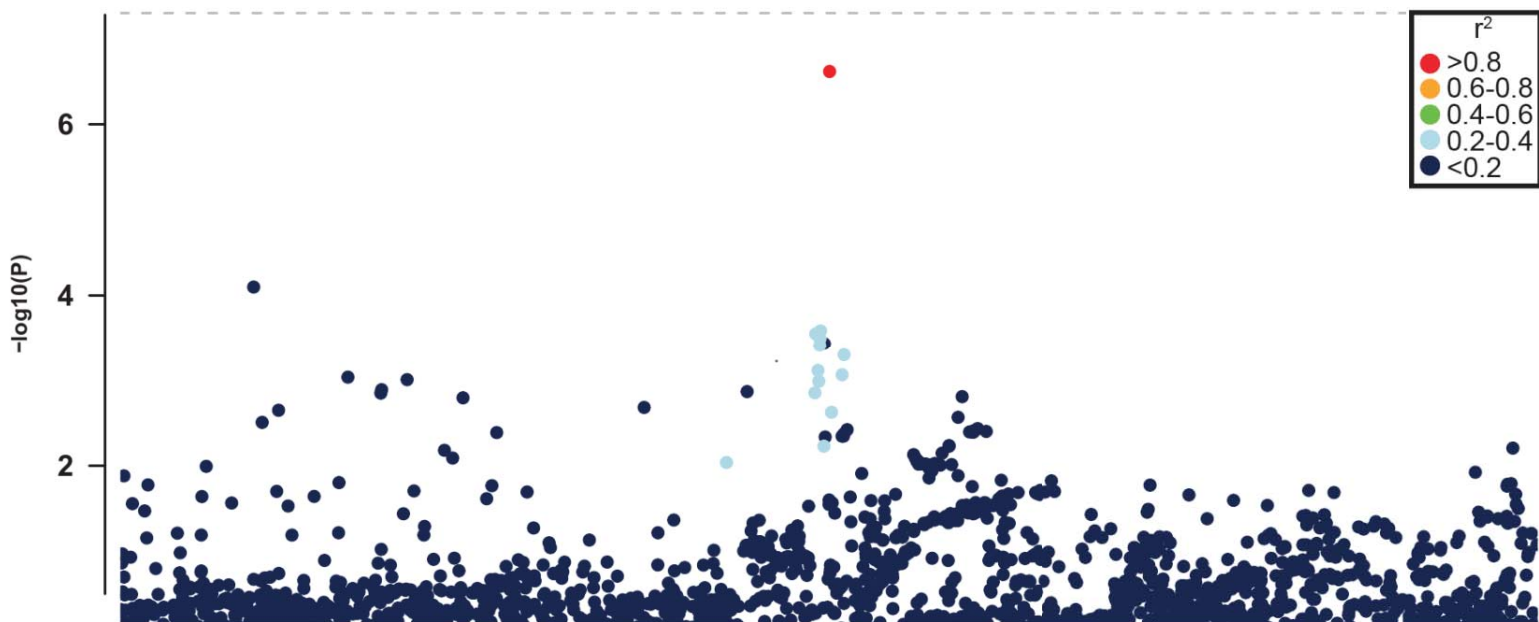
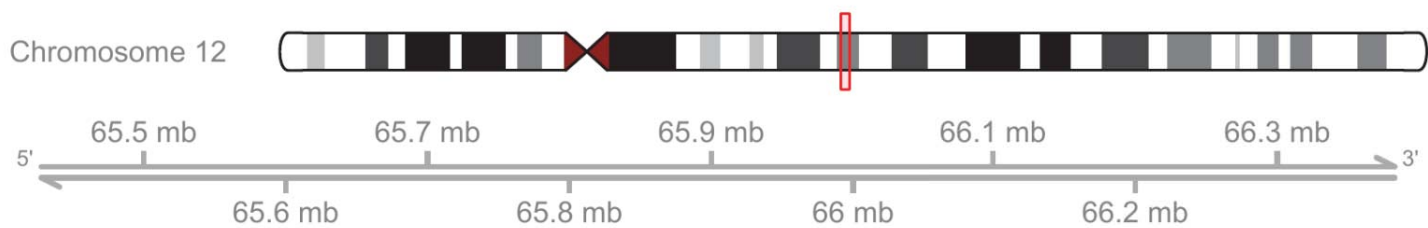
Uncorrected



Zscore of the effect allele: A

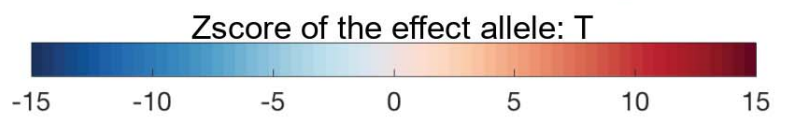
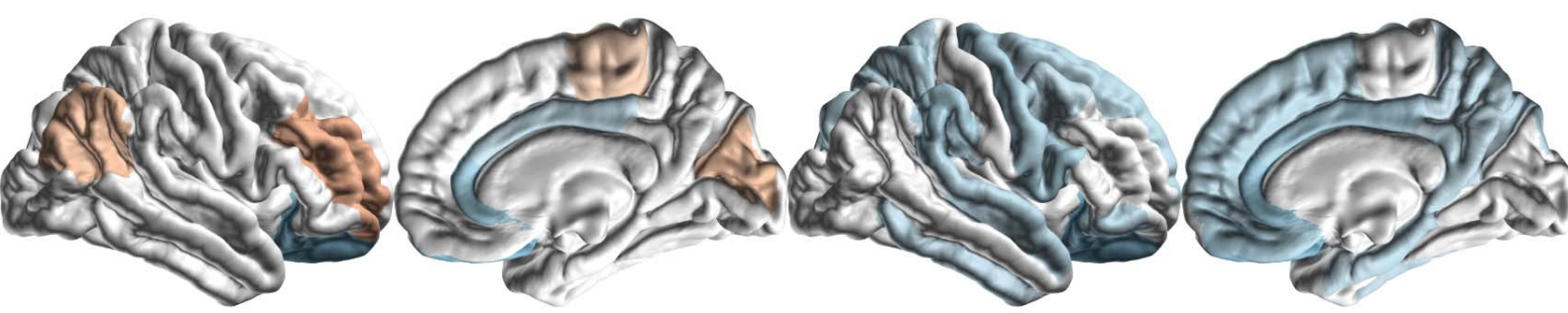
-15 -10 -5 0 5 10 15

Lateral Orbitofrontal SA: rs79487293

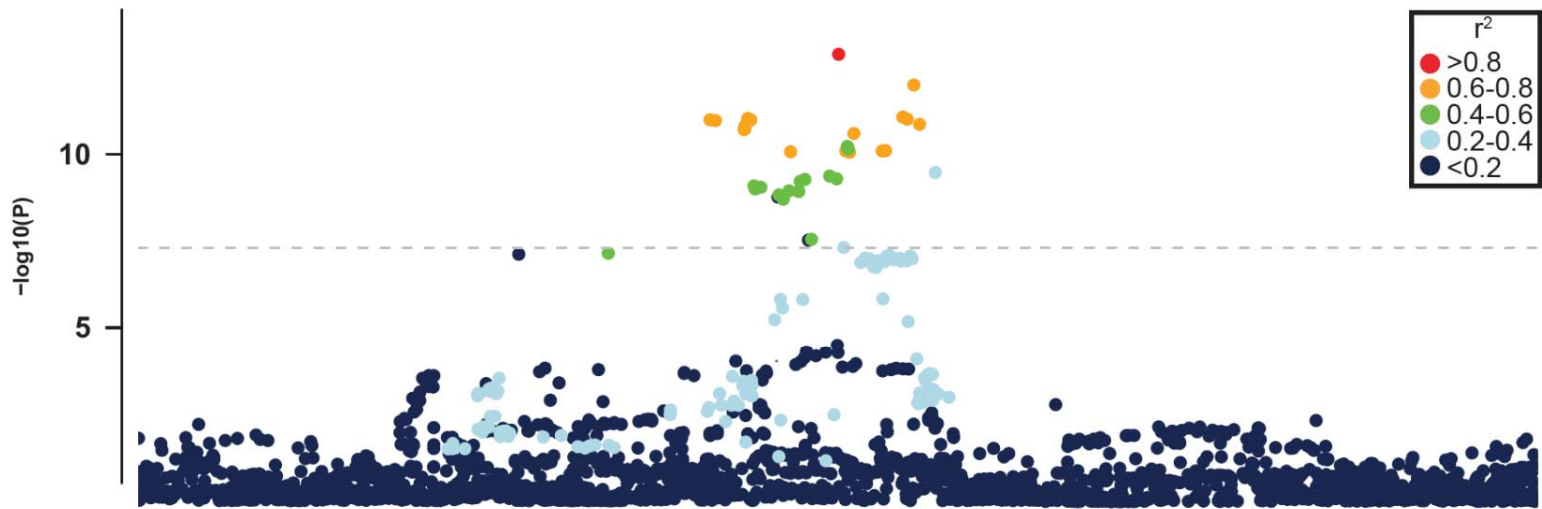
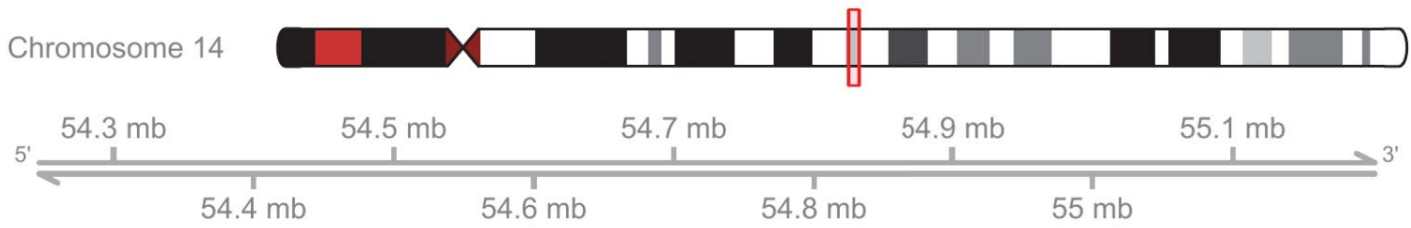


Corrected for total SA

Uncorrected



Lateral Orbitofrontal SA: rs2358483



BMP4

CDKN3

CNIH1

GMFB

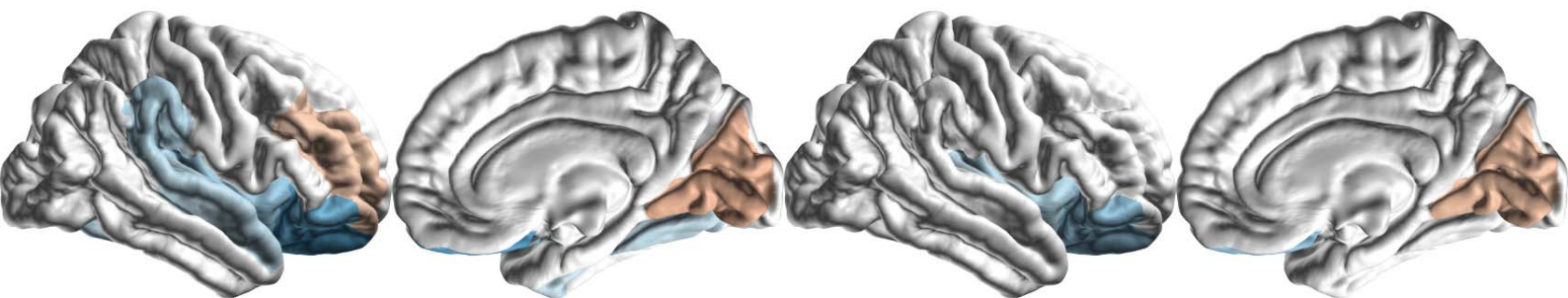
CGRRF1

CP>GZ

GZ>CP

Corrected for total SA

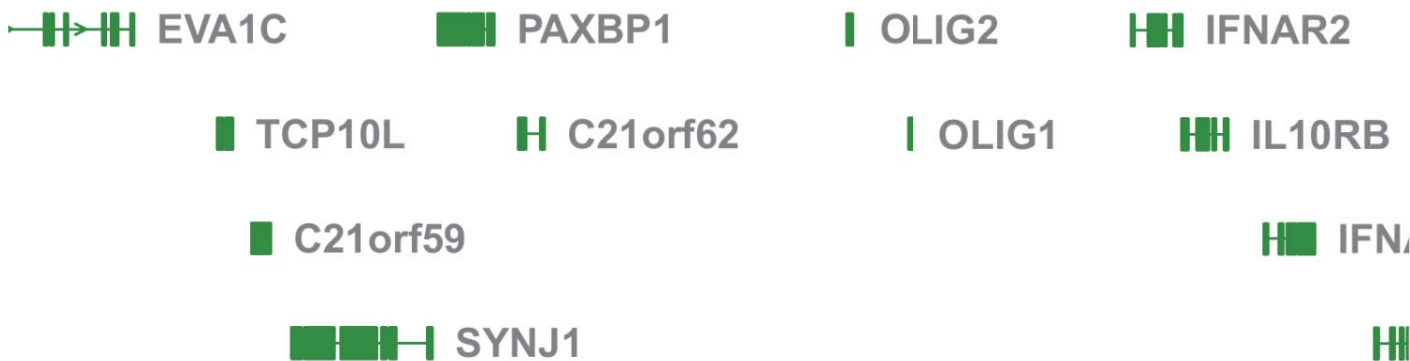
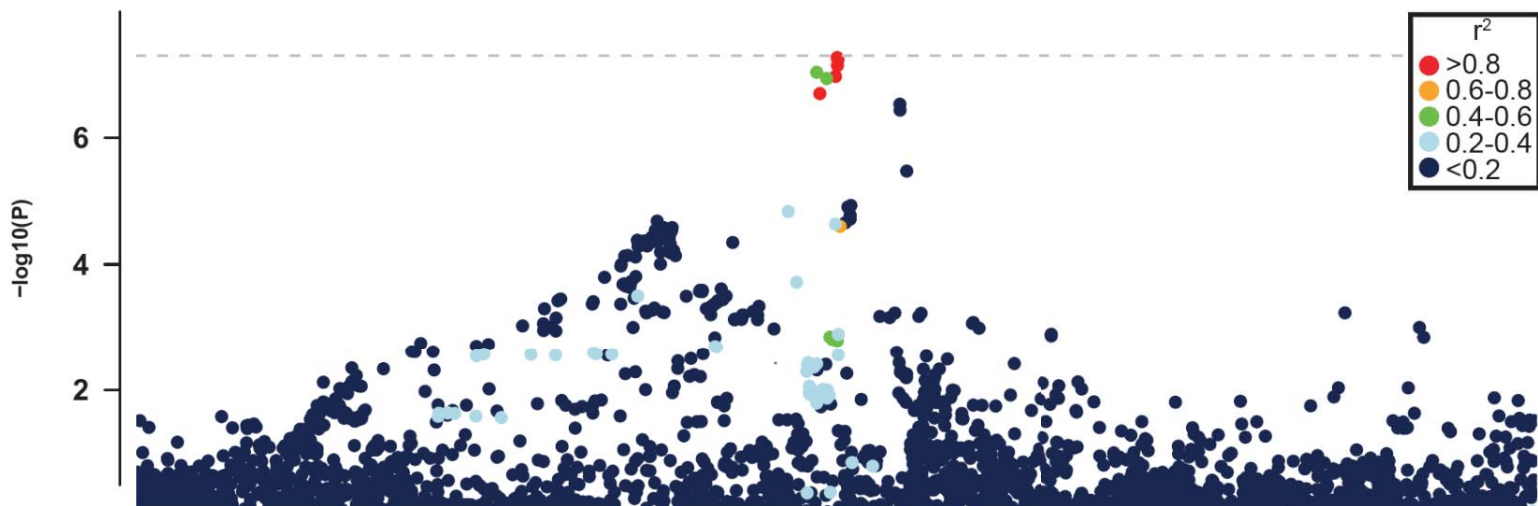
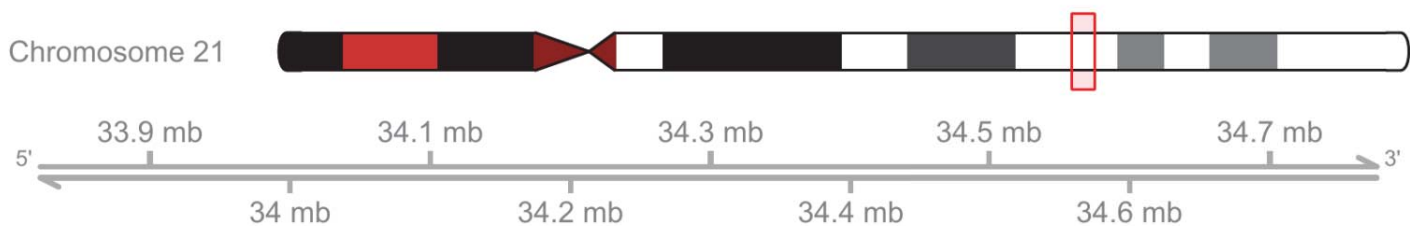
Uncorrected



Zscore of the effect allele: T

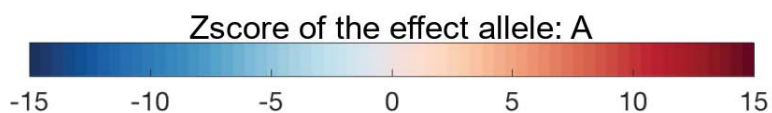
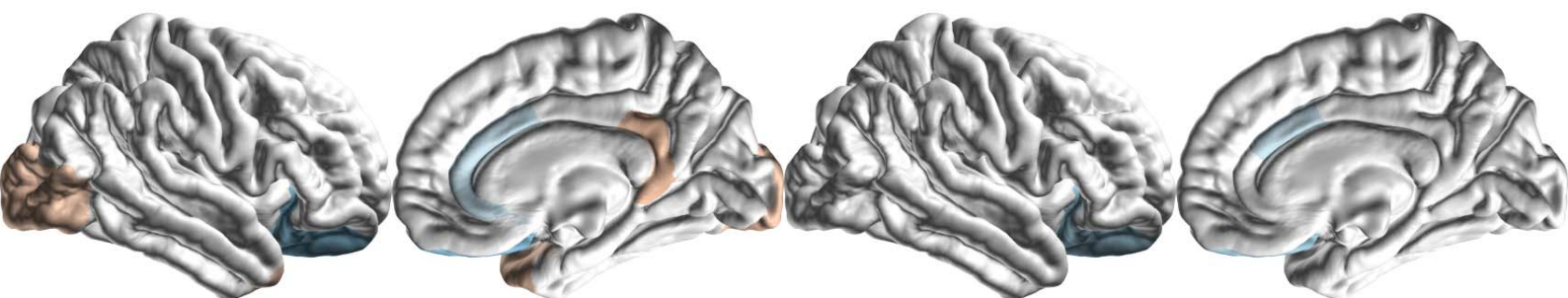


Lateral Orbitofrontal SA: rs12626790

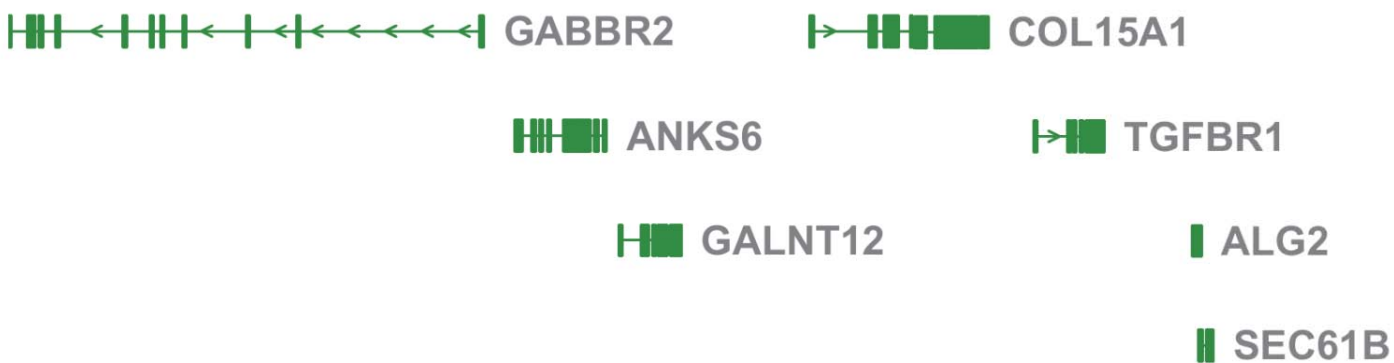
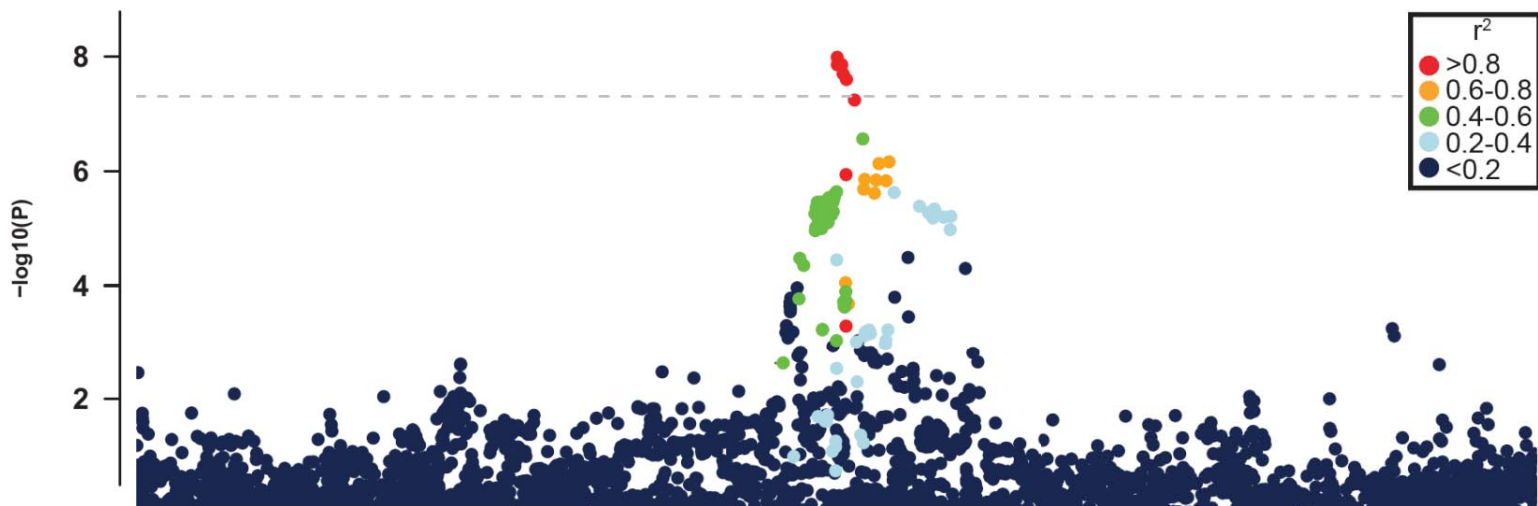
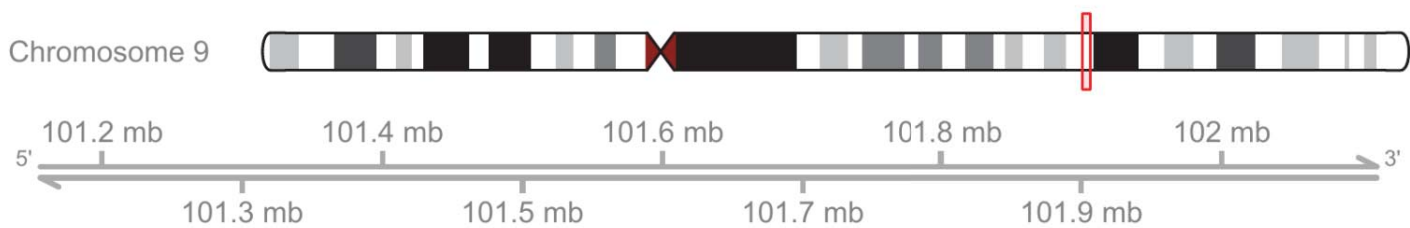


Corrected for total SA

Uncorrected

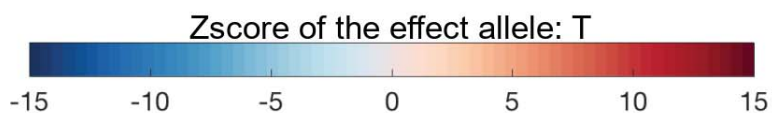
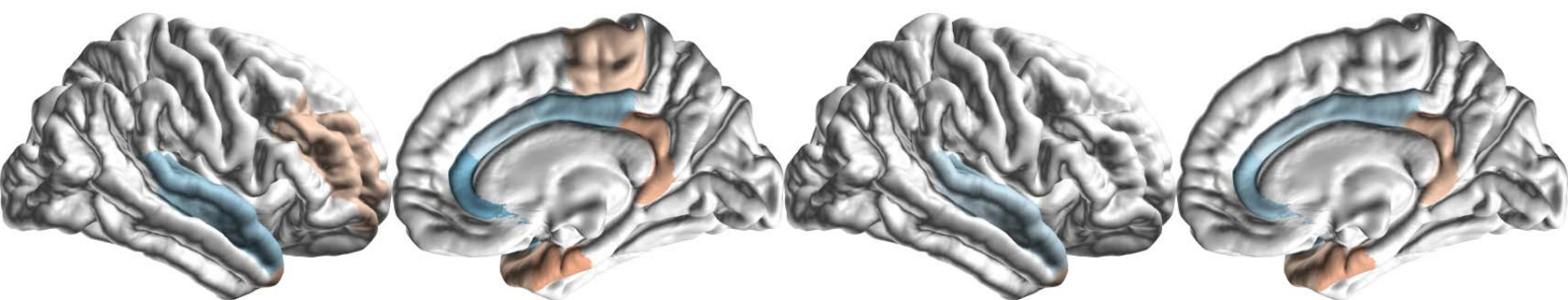


Rostral Anterior Cingulate SA: rs7874287



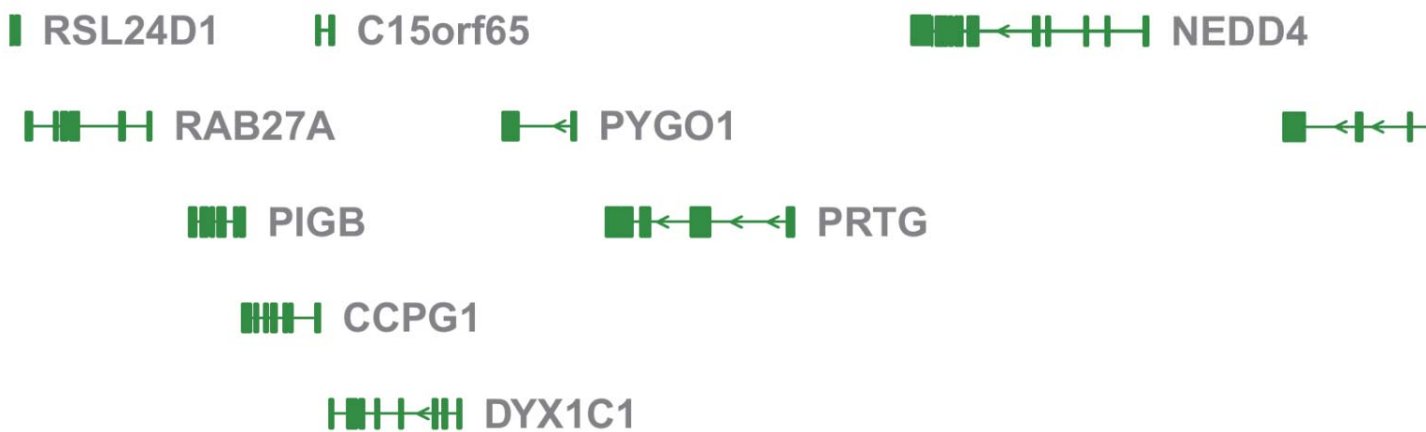
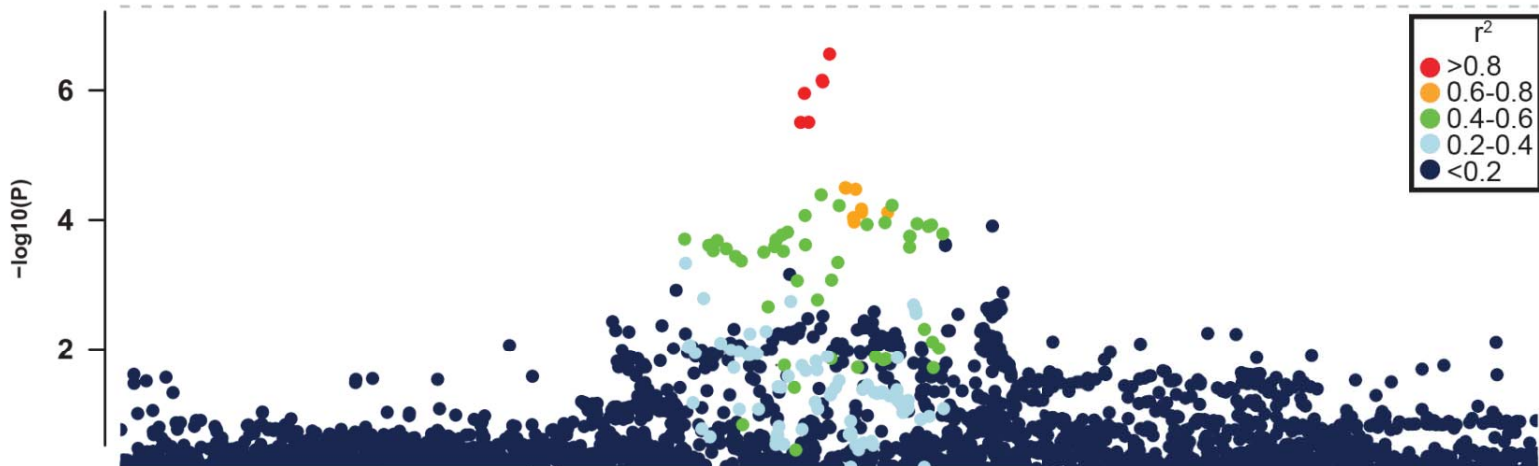
Corrected for total SA

Uncorrected



Rostral Anterior Cingulate SA: rs1986012

Chromosome 15

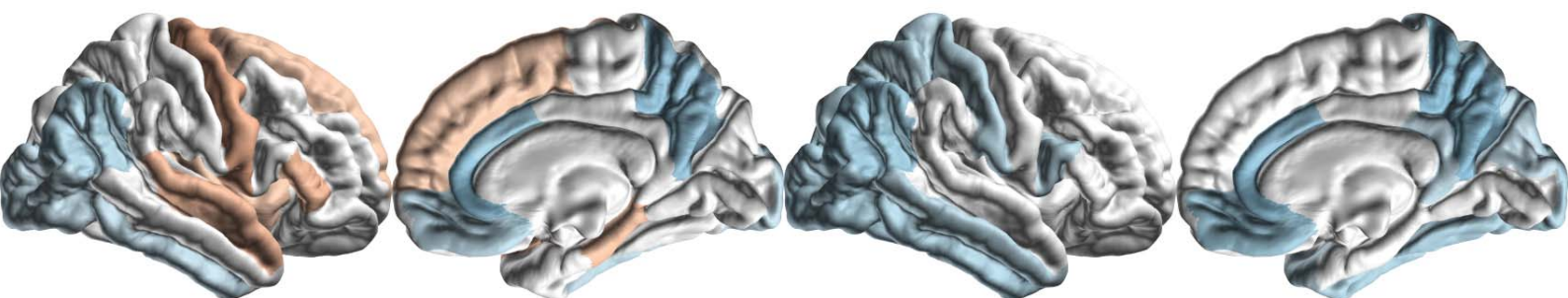


CP>GZ GZ>CP



Corrected for total SA

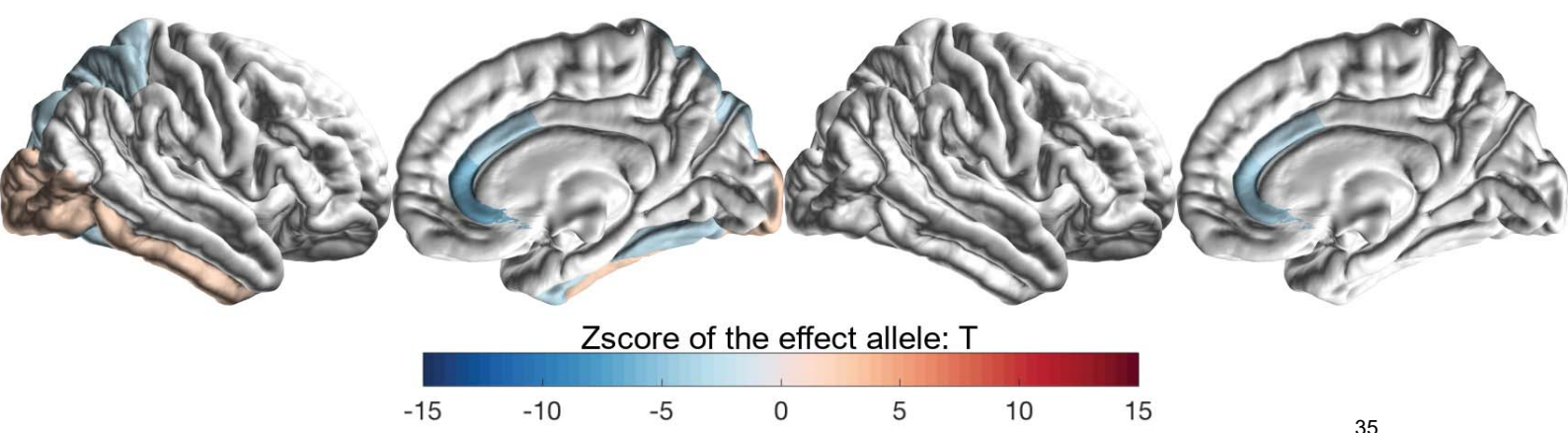
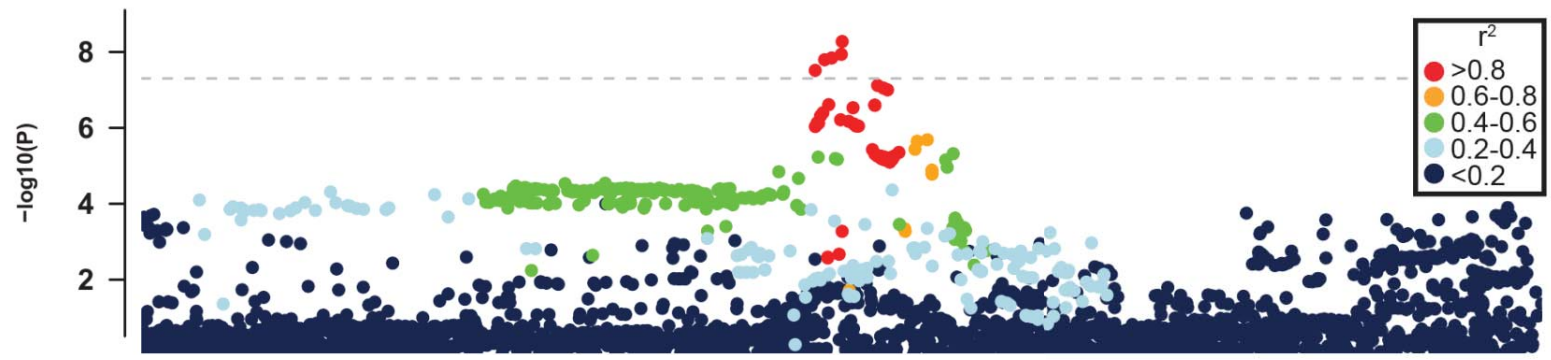
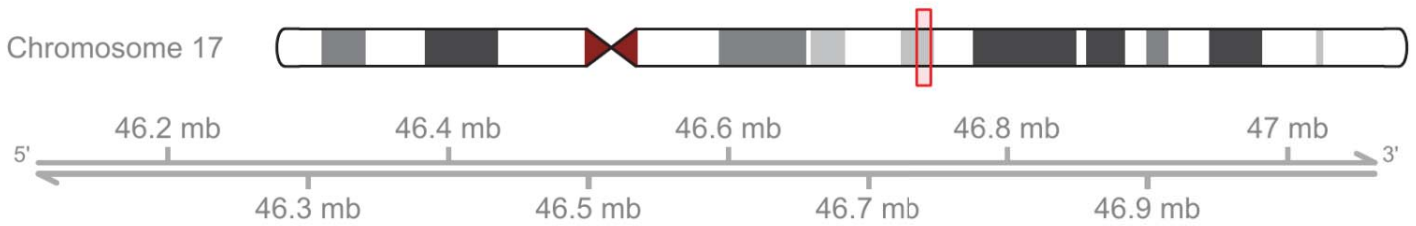
Uncorrected



Zscore of the effect allele: T

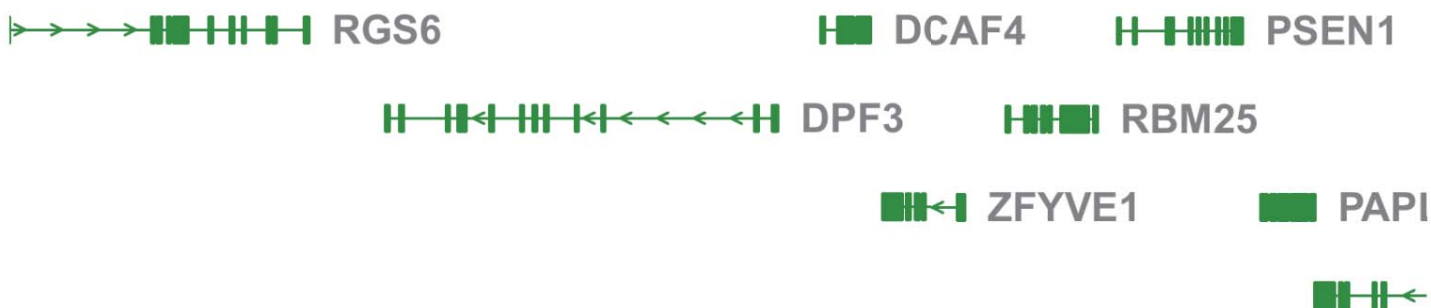
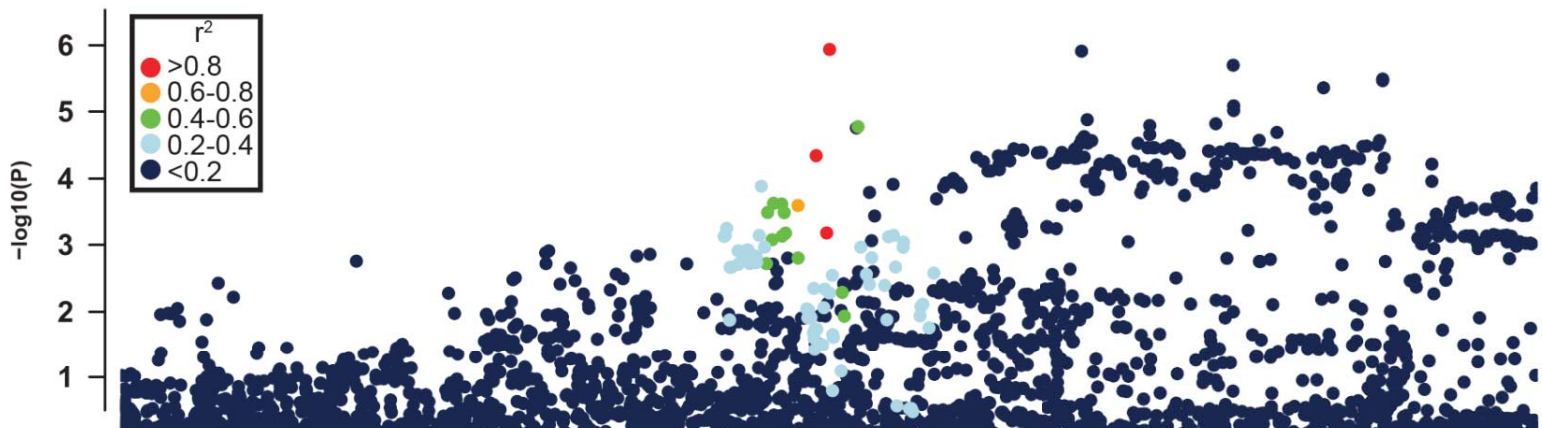
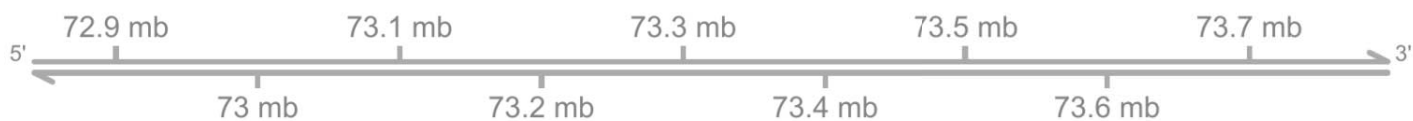


Rostral Anterior Cingulate SA: rs2202895



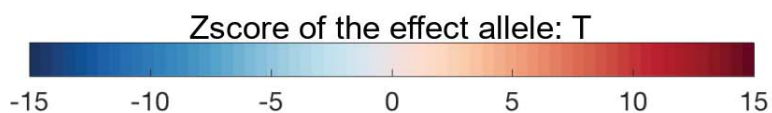
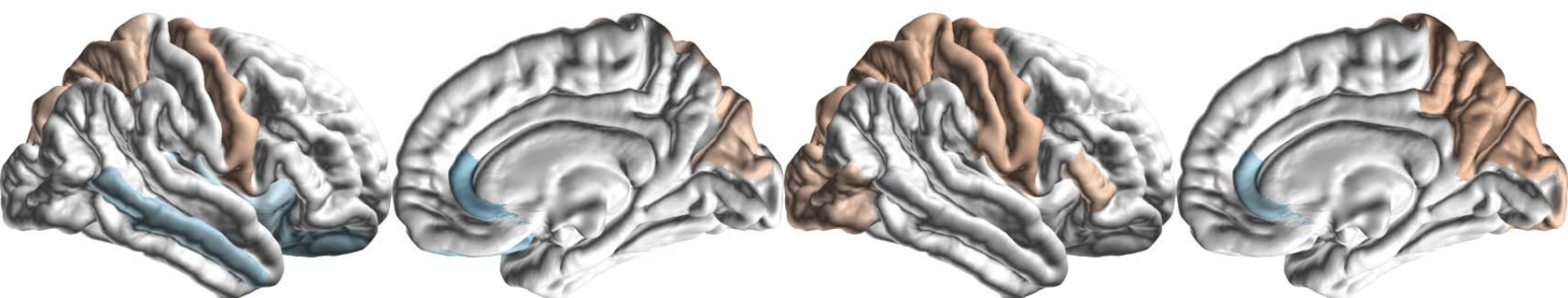
Rostral Anterior Cingulate TH: rs4899448

Chromosome 14

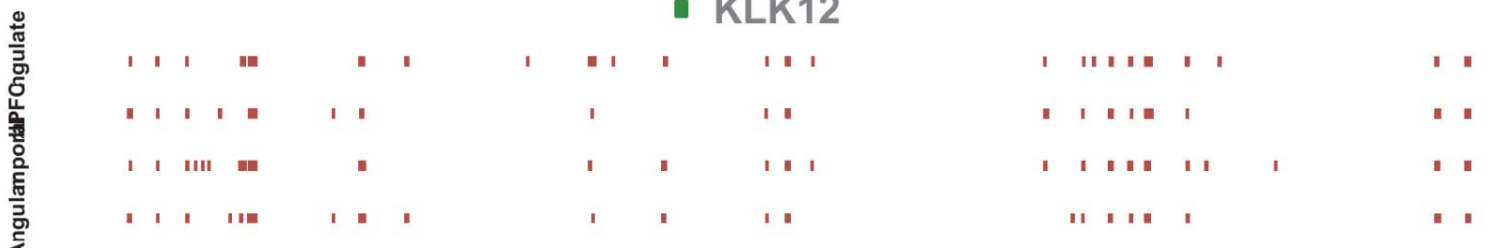
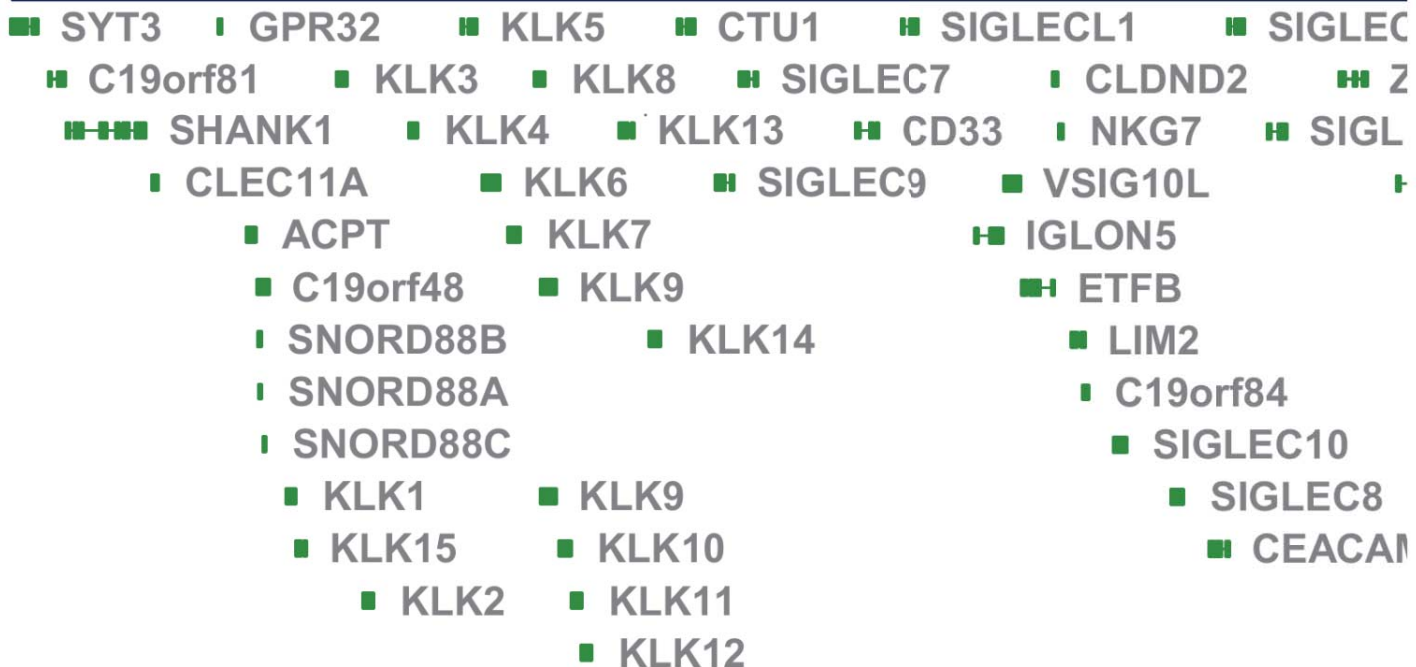
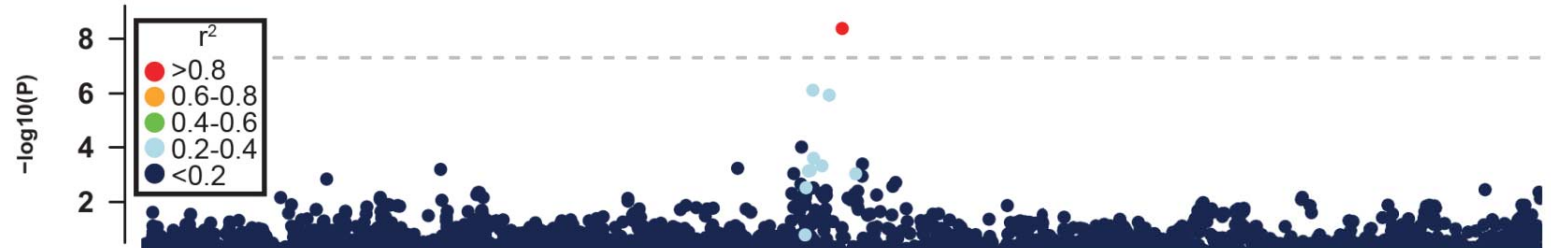
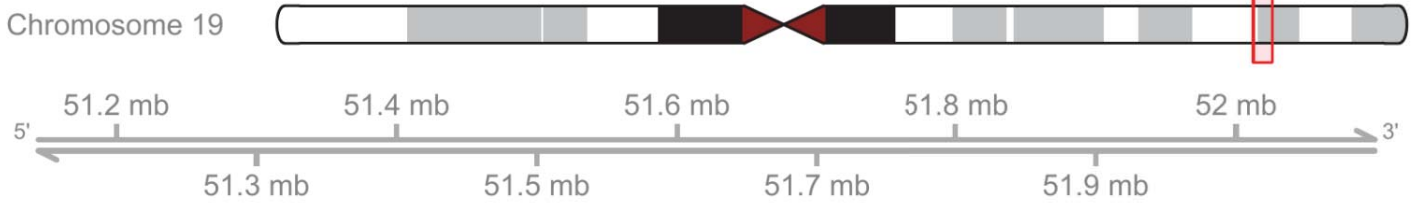


Corrected for average TH

Uncorrected

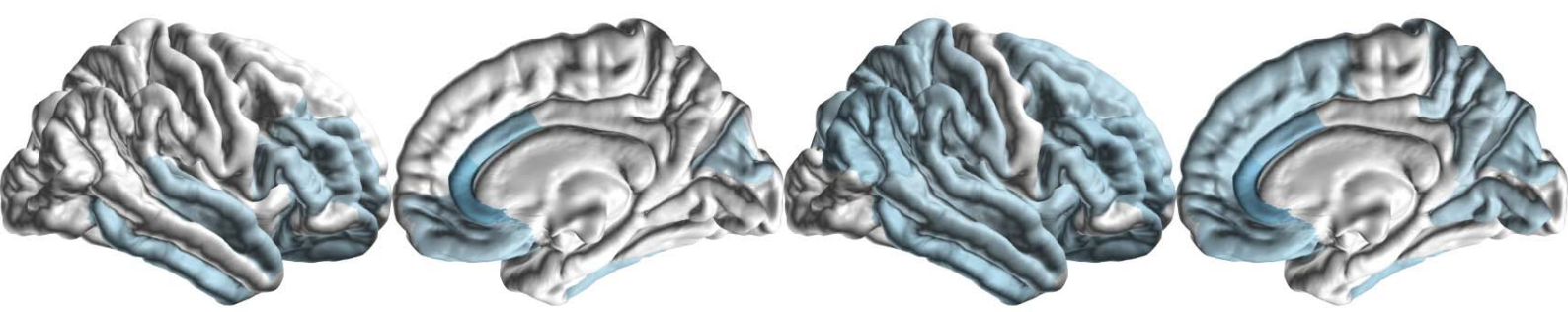


Rostral Anterior Cingulate TH: rs62115964



Corrected for average TH

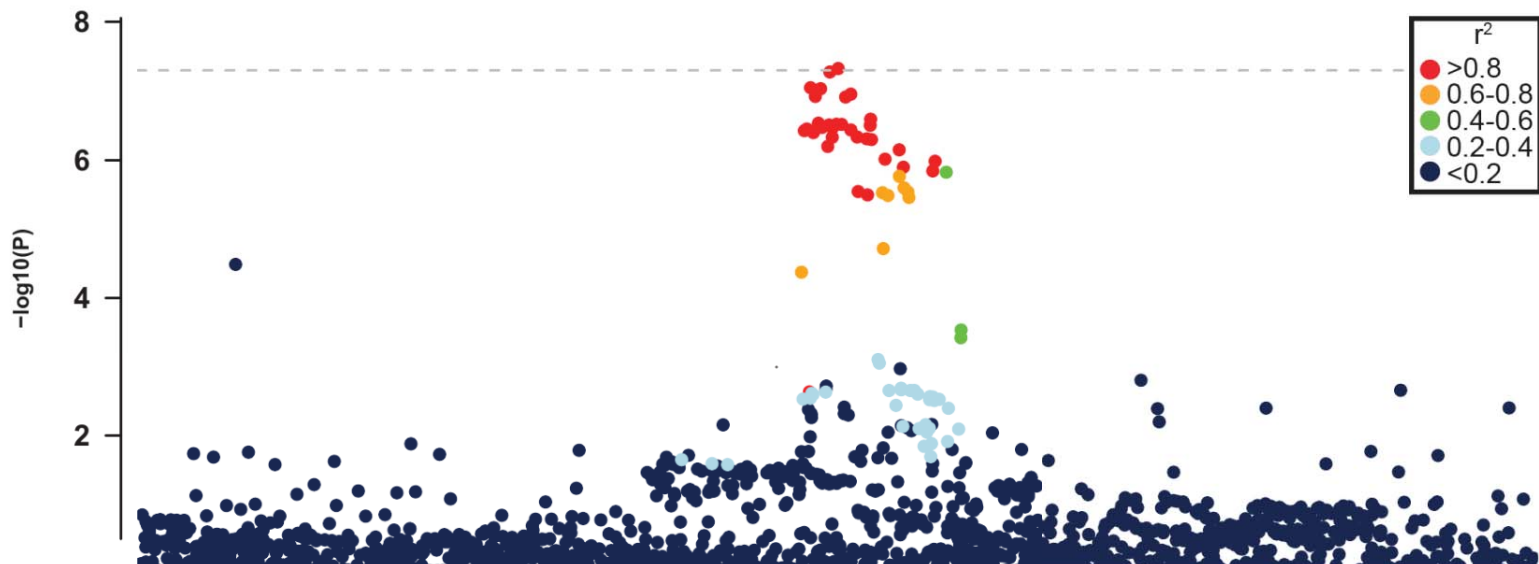
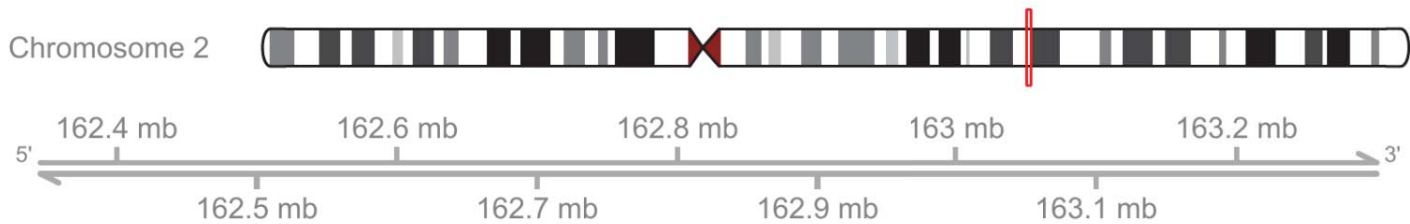
Uncorrected



Zscore of the effect allele: A

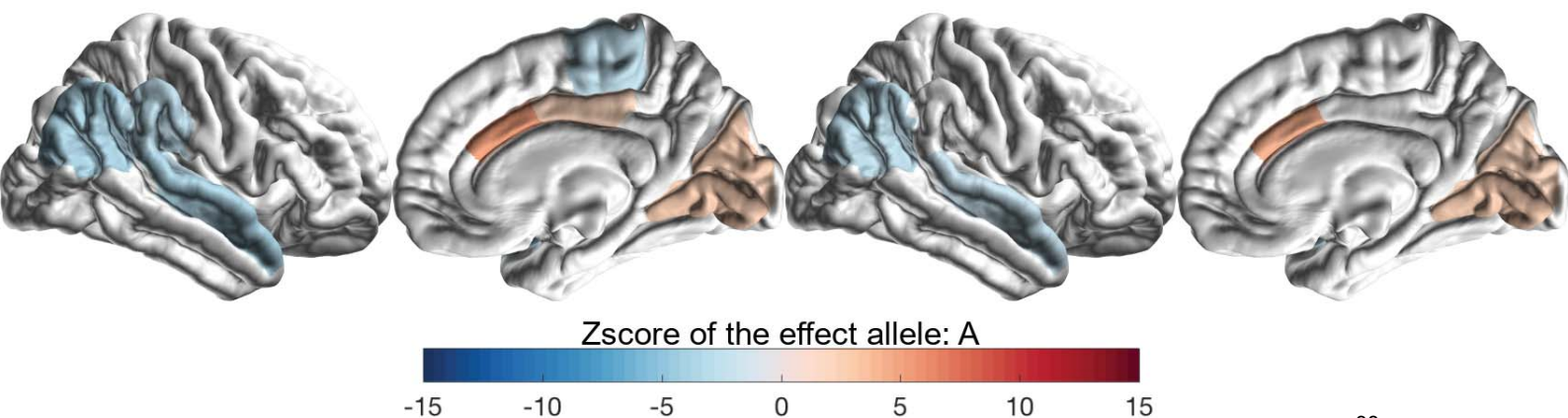


Caudal Anterior Cingulate SA: rs13021985



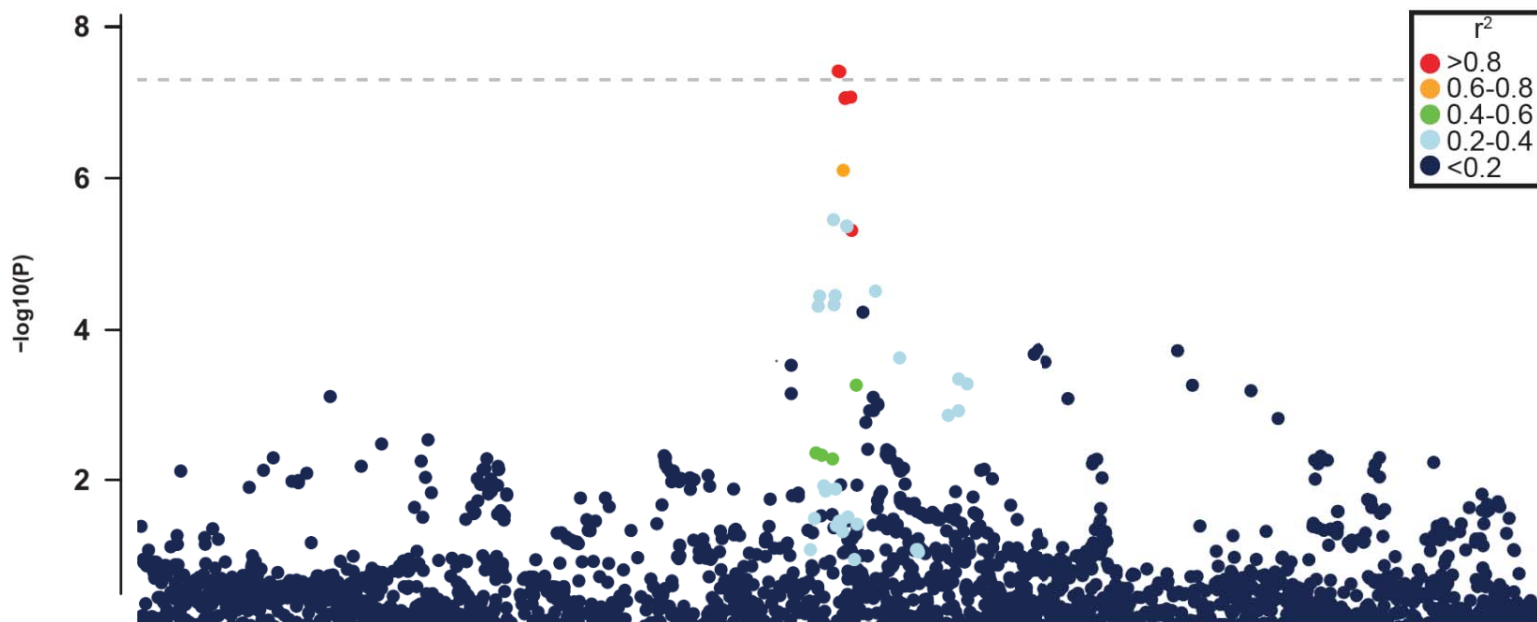
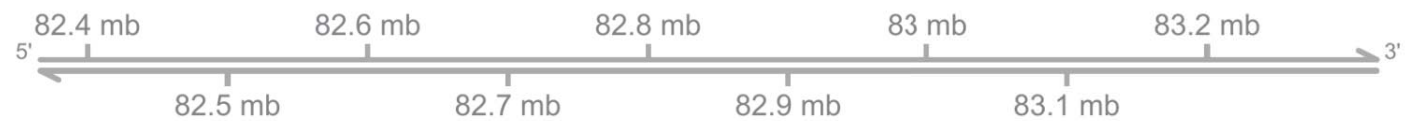
Corrected for total SA

Uncorrected



Caudal Anterior Cingulate SA: rs7728751

Chromosome 5



TMEM167A

VCAN

HAPLN1

XRCC4

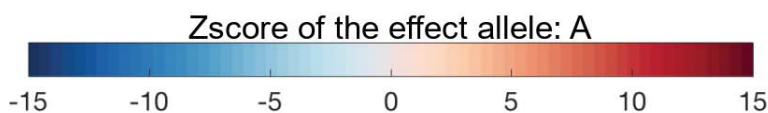
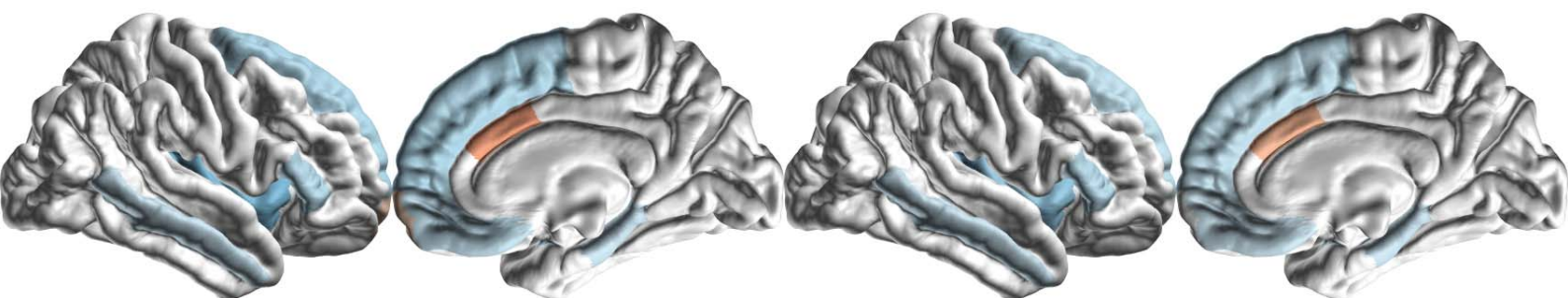
HAPLN1

GZ>CP

CP>GZ

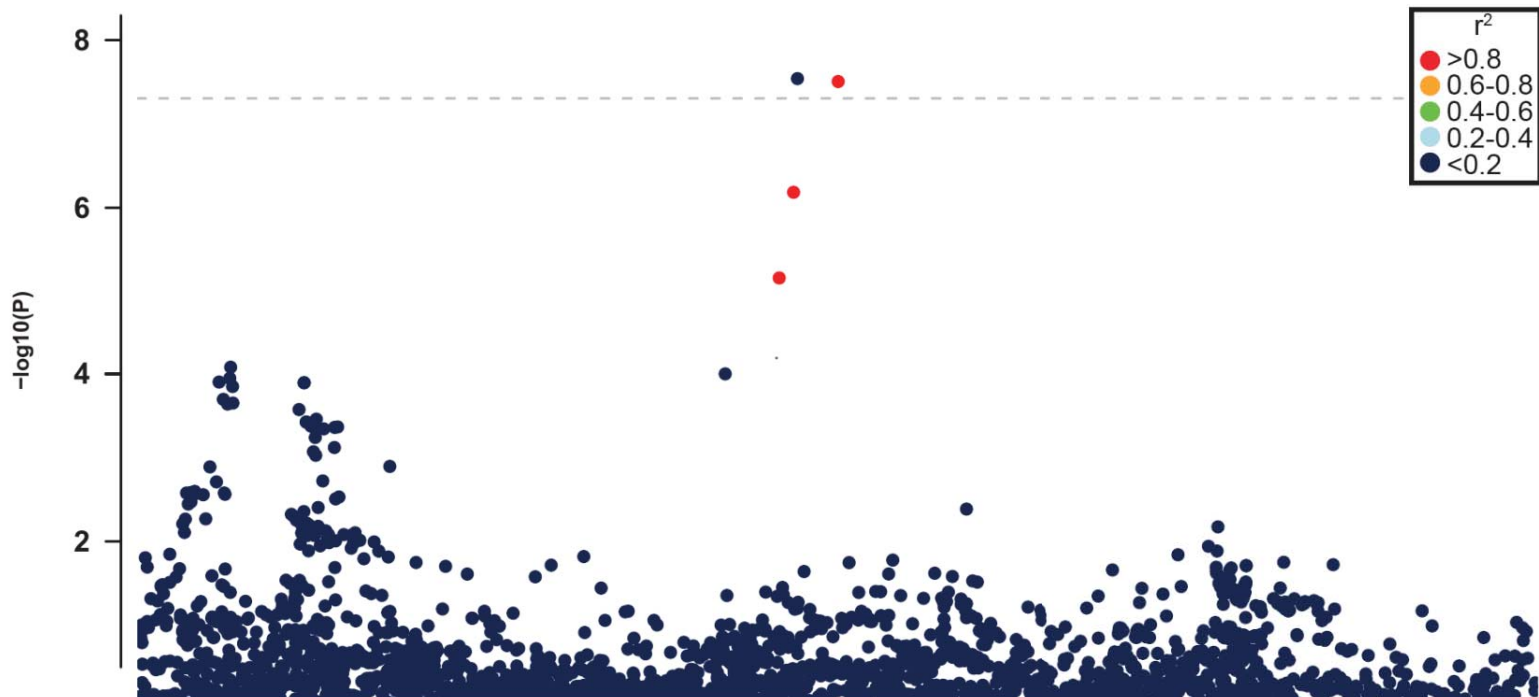
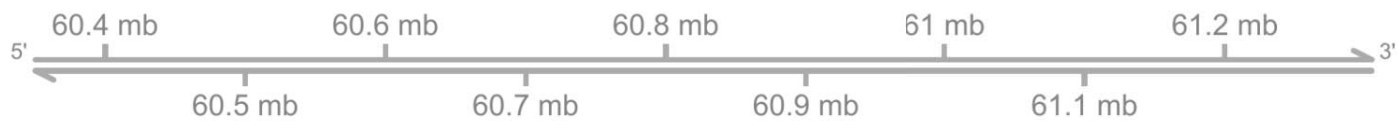
Corrected for total SA

Uncorrected



Caudal Anterior Cingulate SA: rs80241863

Chromosome 16

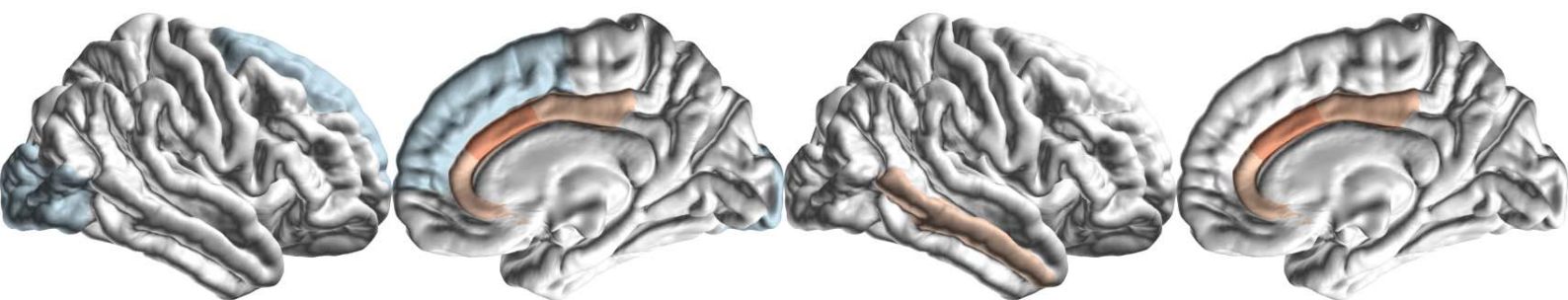


GZ>CP

CP>GZ

Corrected for total SA

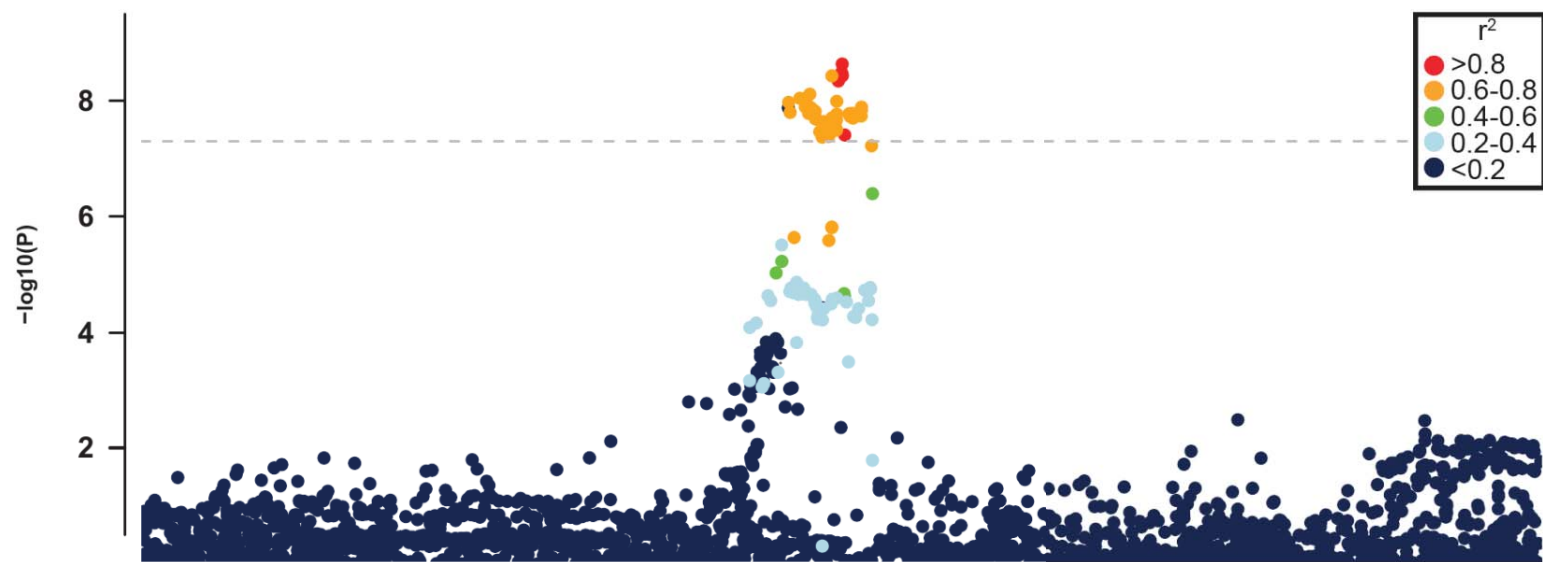
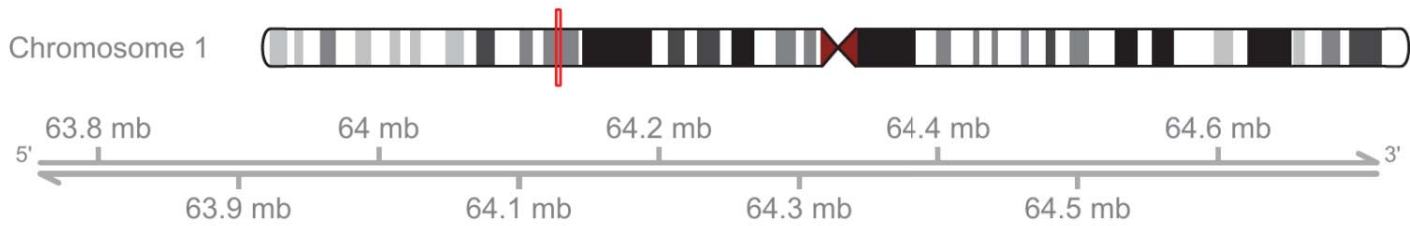
Uncorrected



Zscore of the effect allele: A

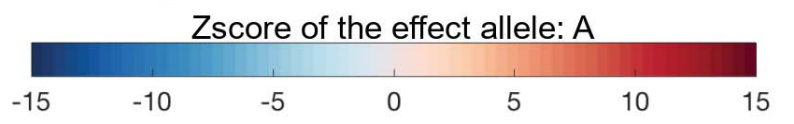
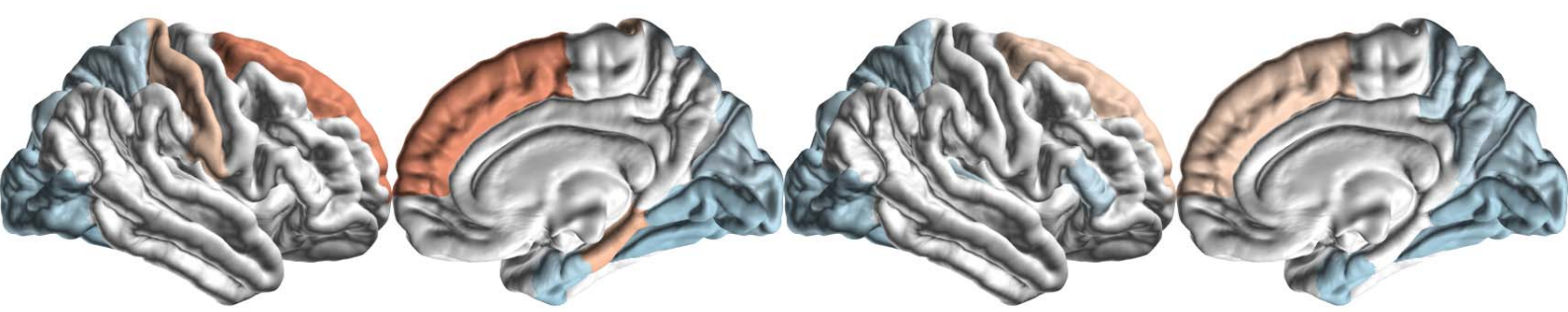


Superior Frontal SA: rs4915928



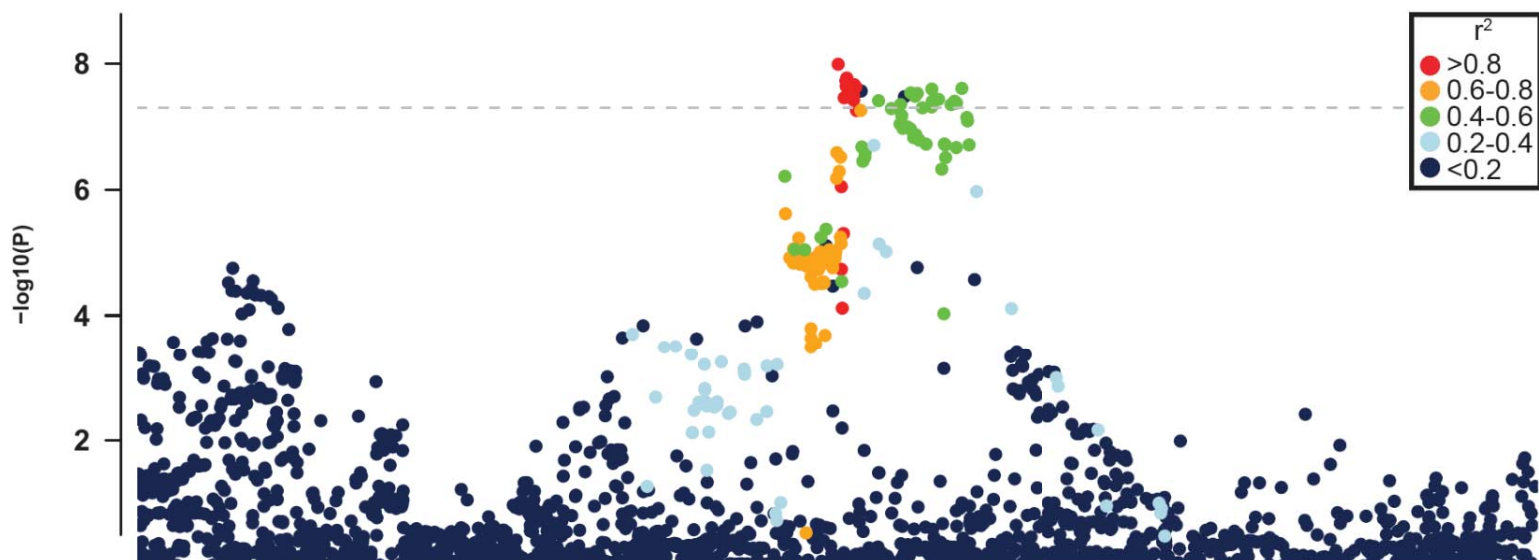
Corrected for total SA

Uncorrected



Superior Frontal SA: rs4492207

Chromosome 6



TPD52L1

HEY2

HINT3

HDDC2

NCOA7

TRMT11

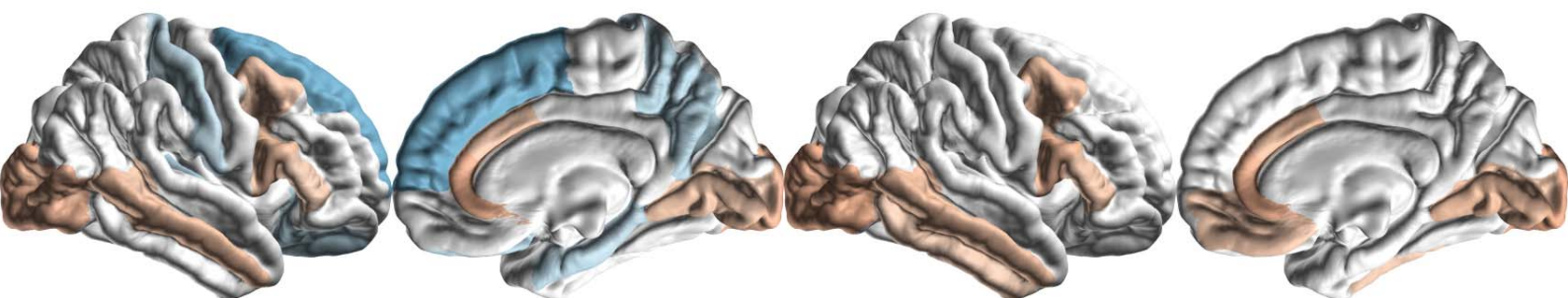
GZ>CP



CP>GZ

Corrected for total SA

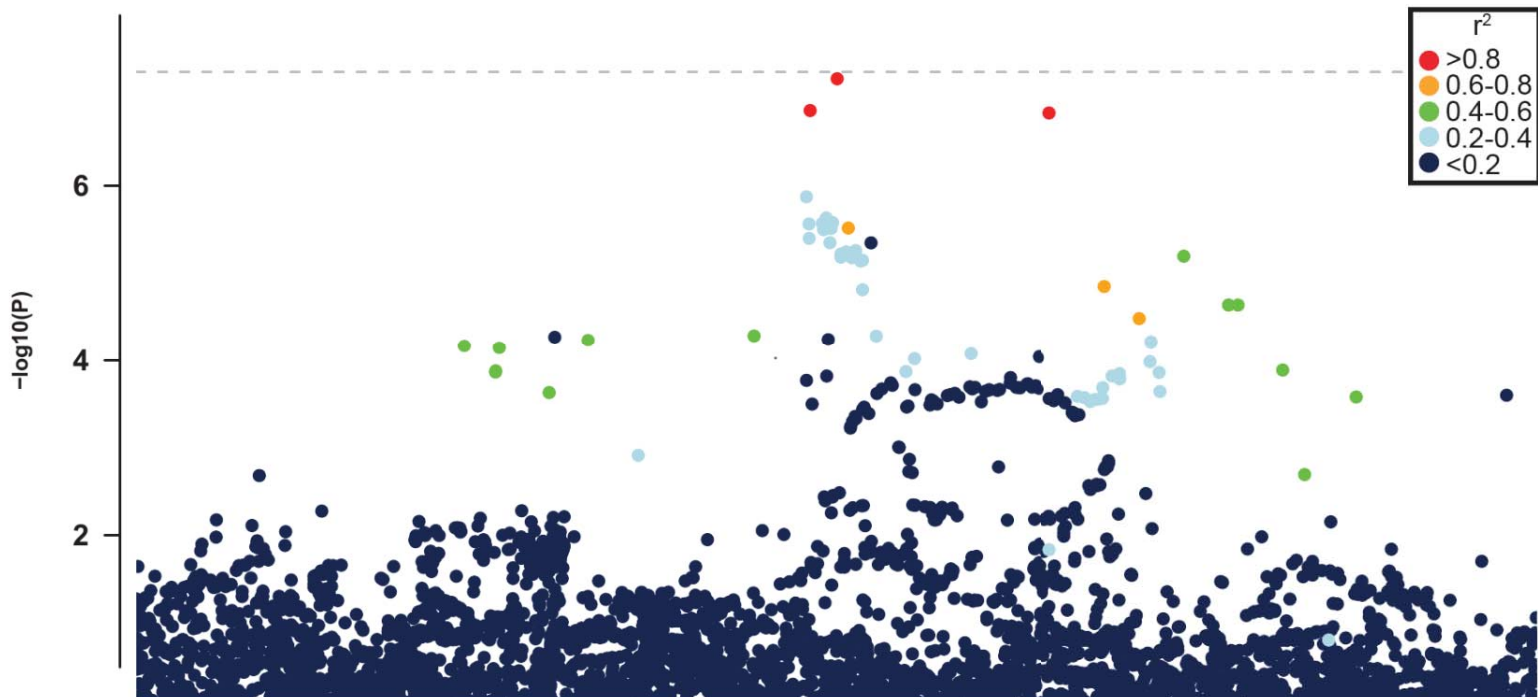
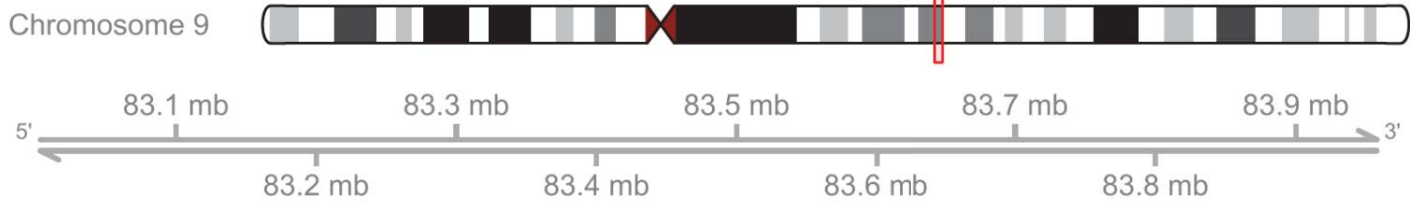
Uncorrected



Zscore of the effect allele: A



Superior Frontal SA: rs76696867



GZ>CP

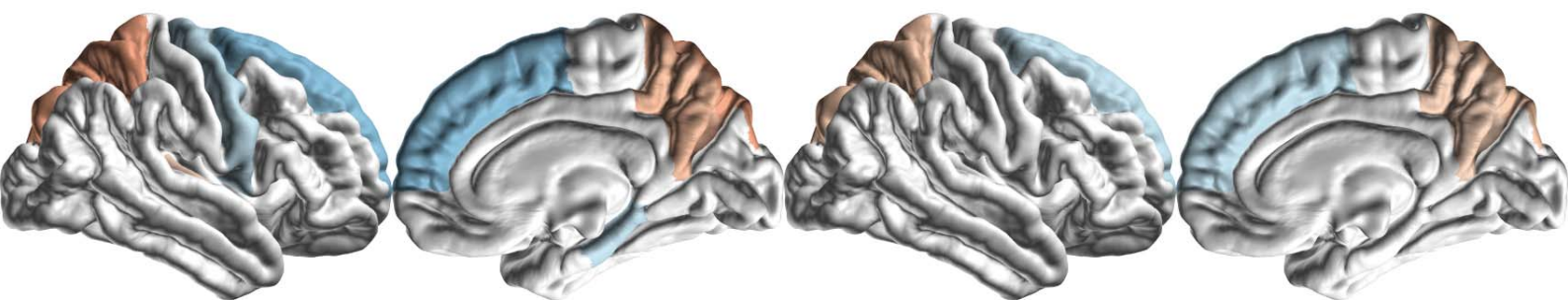


CP>GZ

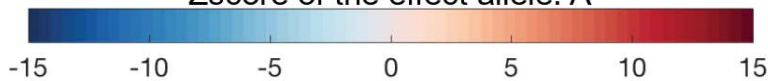


Corrected for total SA

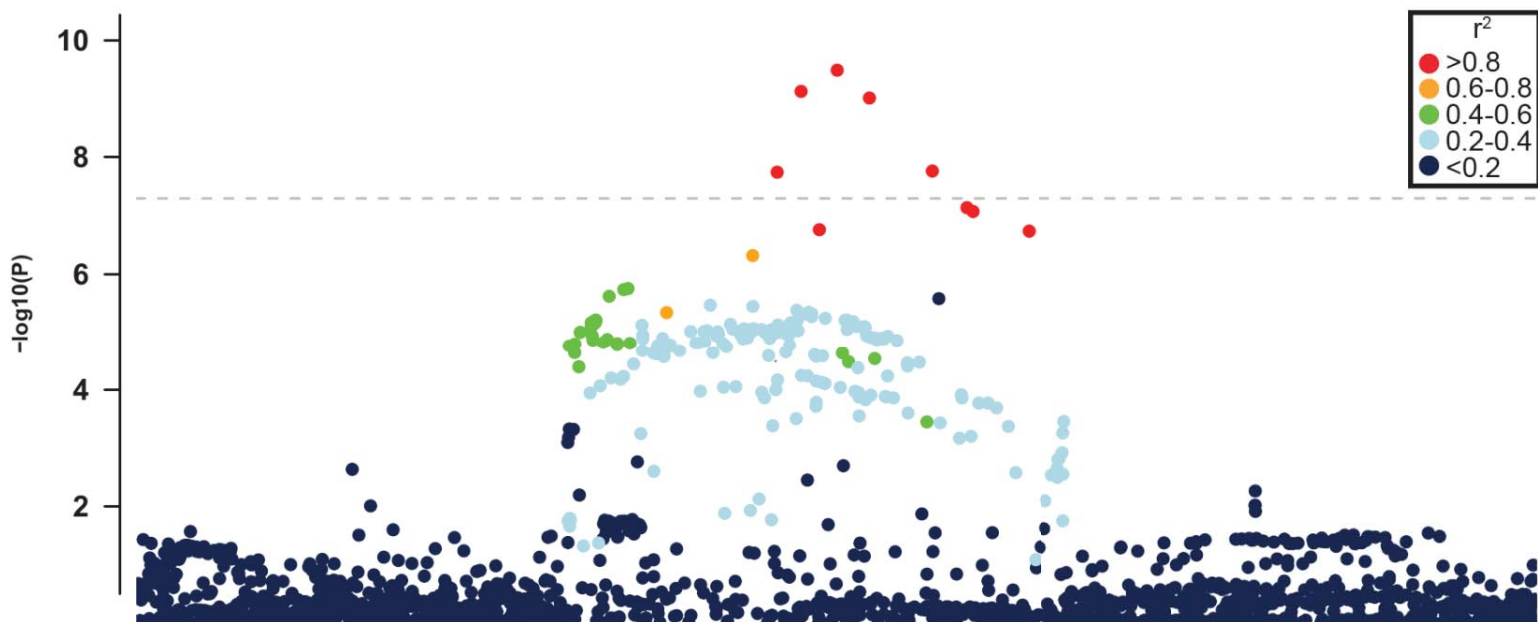
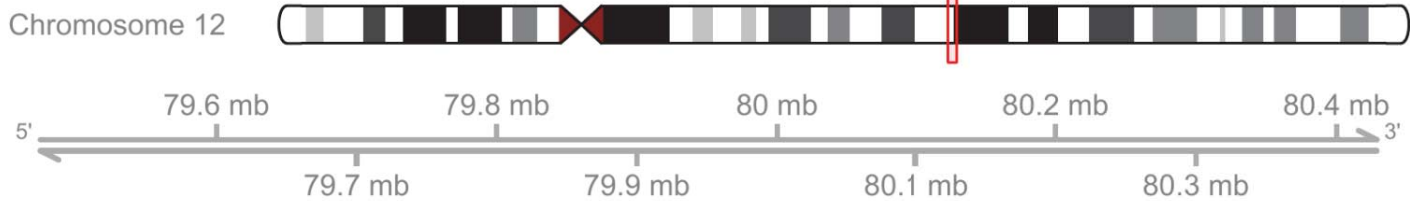
Uncorrected



Zscore of the effect allele: A

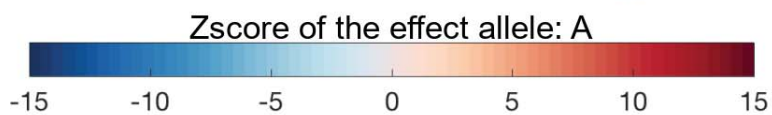
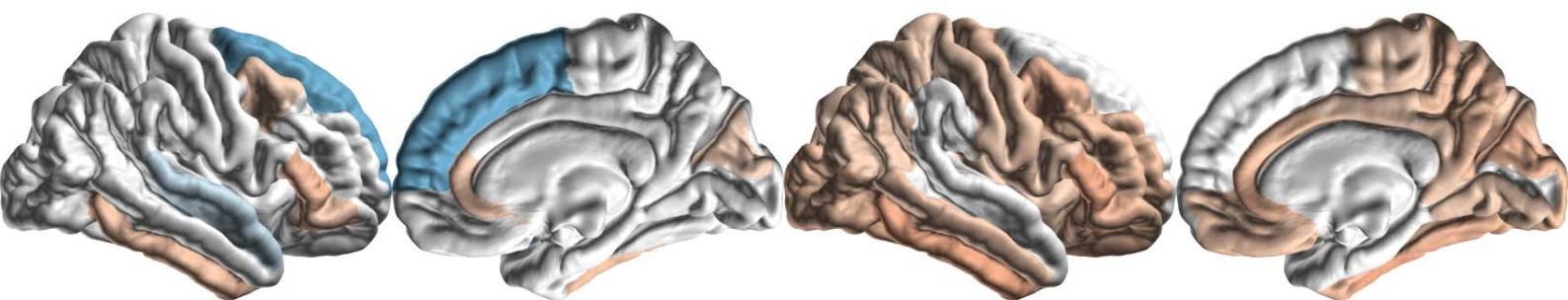


Superior Frontal SA: rs4842266

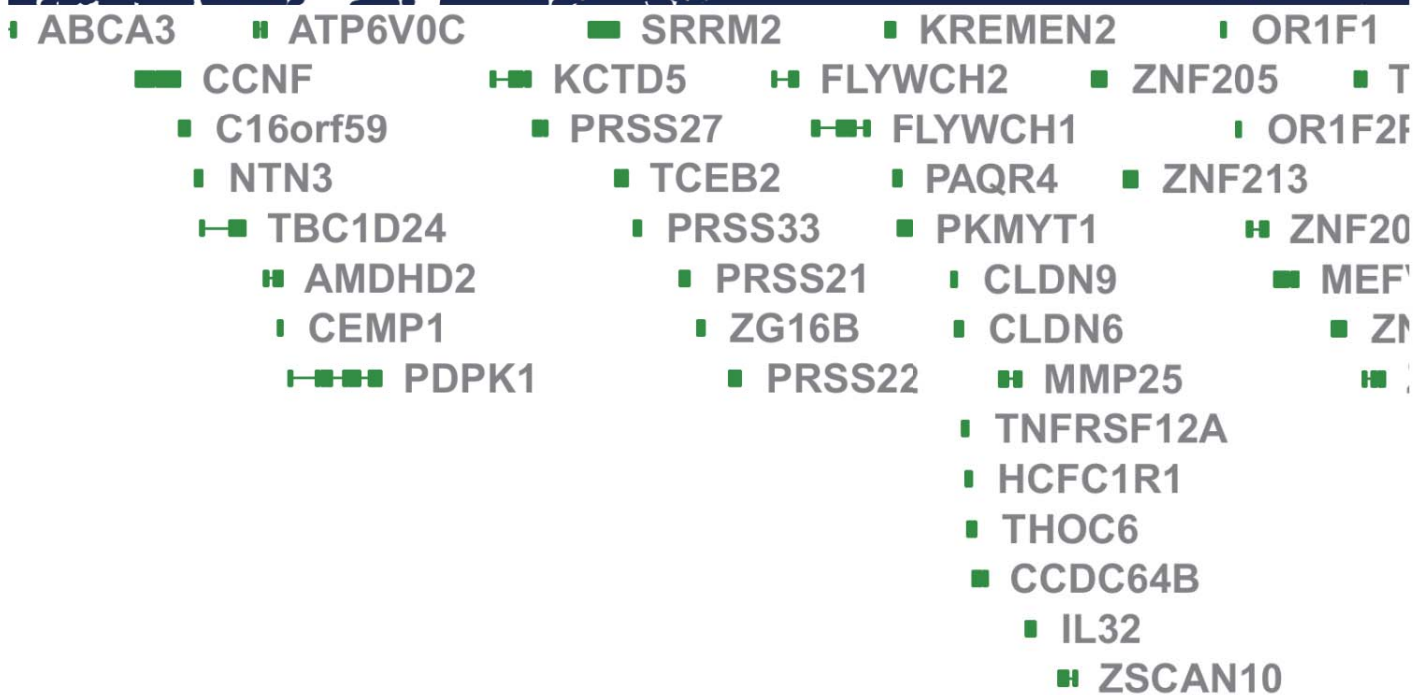
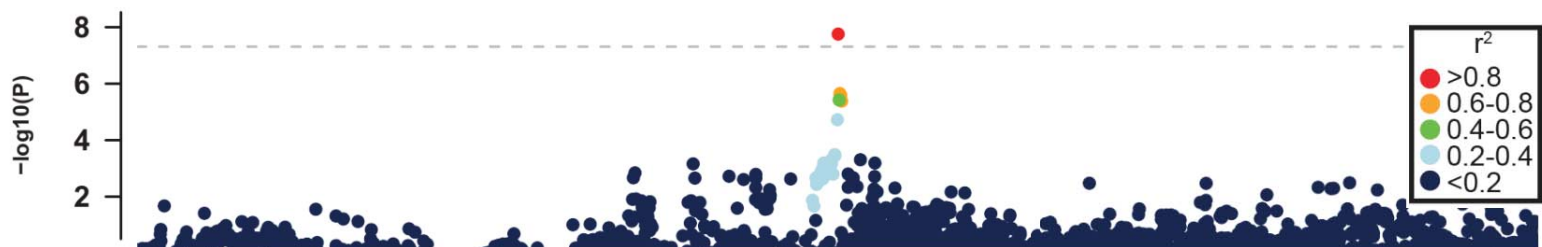
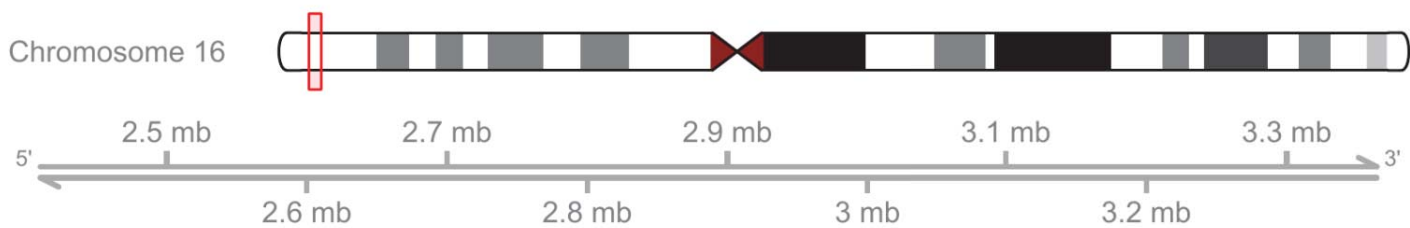


Corrected for total SA

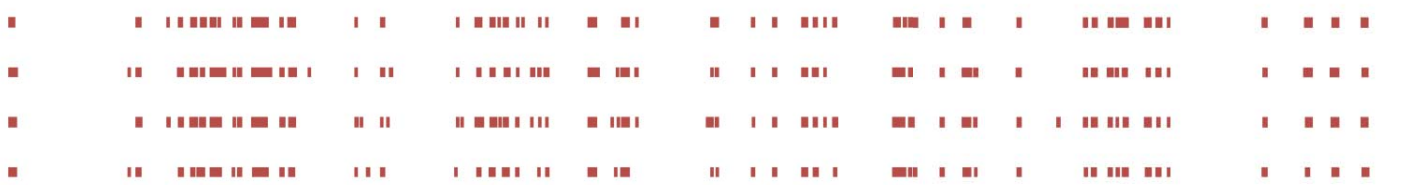
Uncorrected



Superior Frontal TH: rs9929219

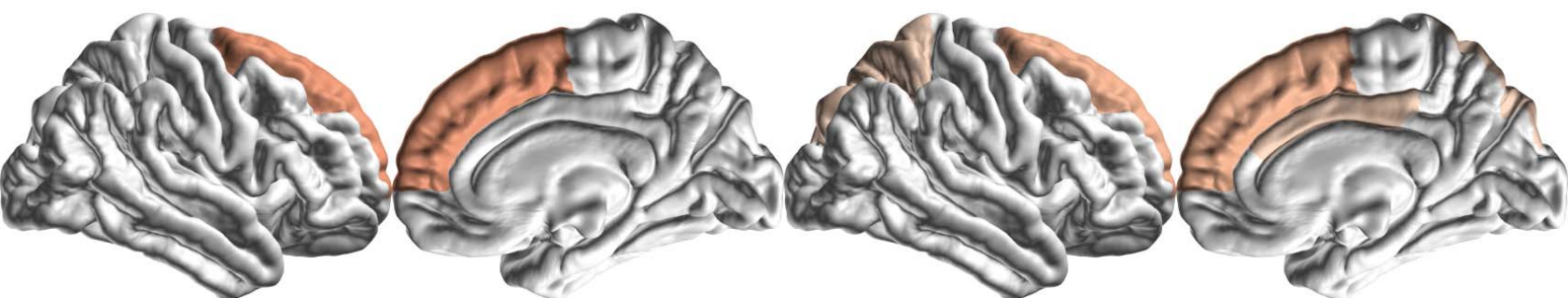


AngiampolifFGulate



Corrected for average TH

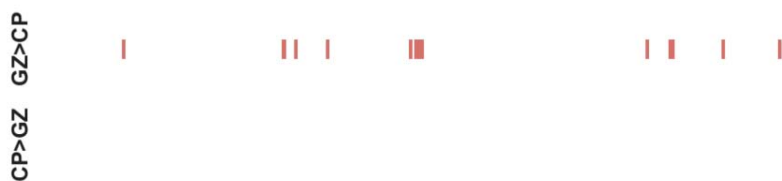
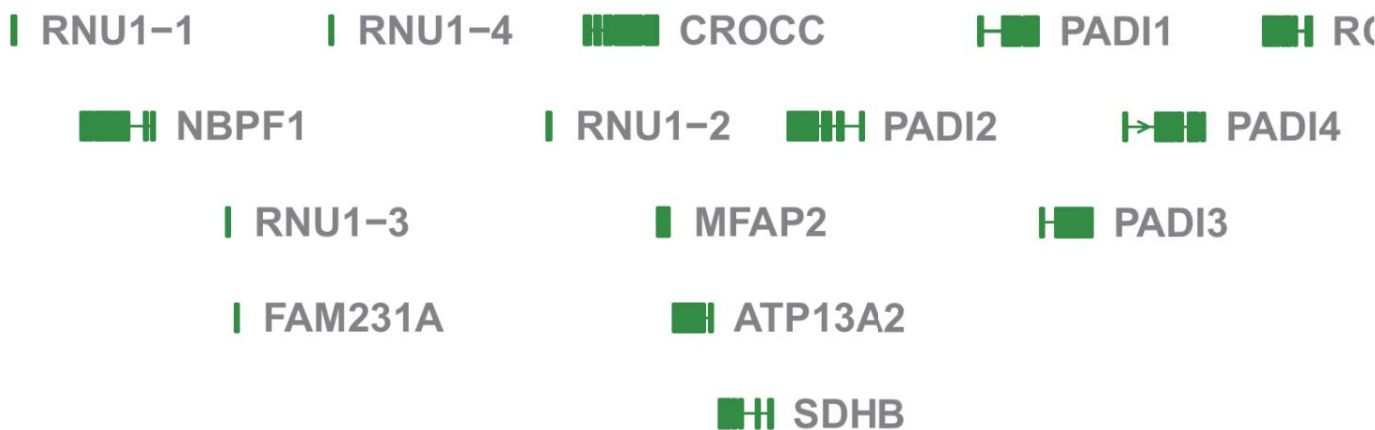
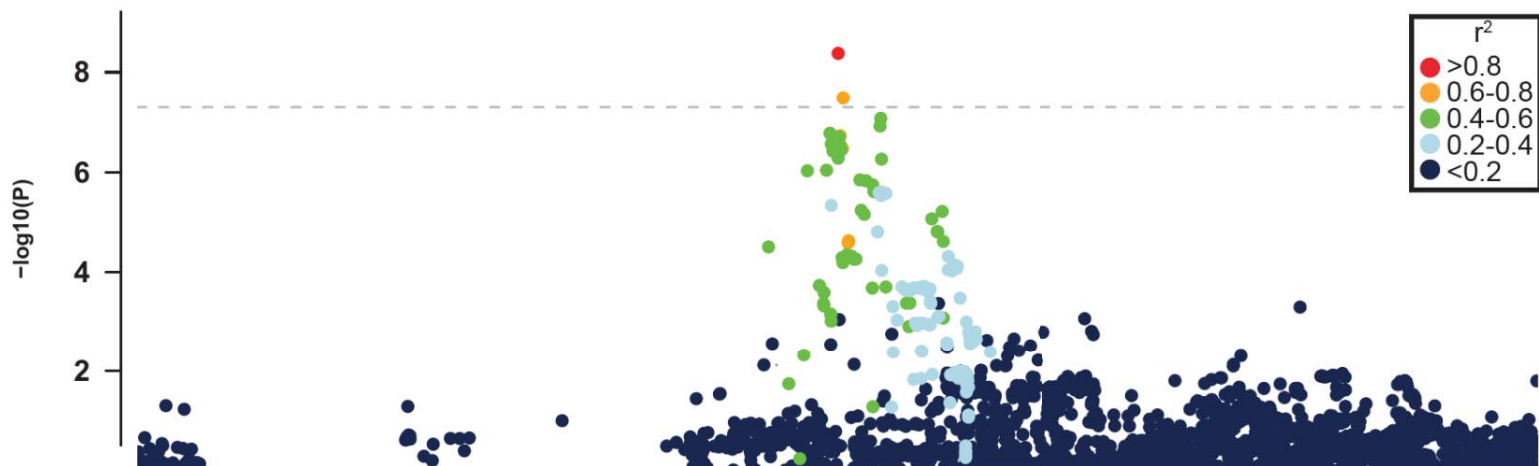
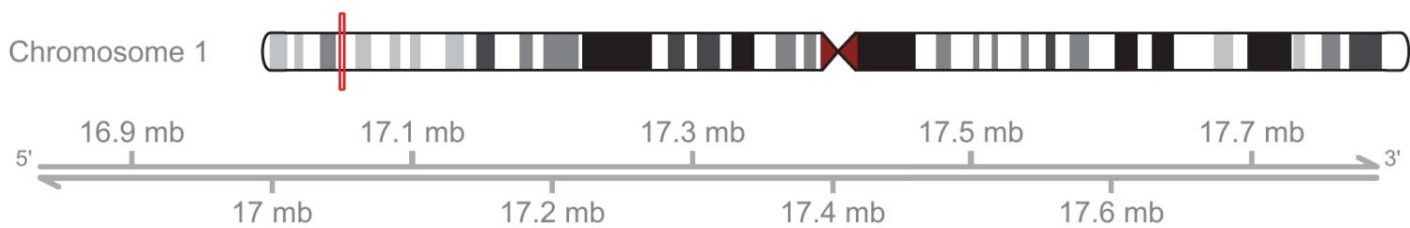
Uncorrected



Zscore of the effect allele: A

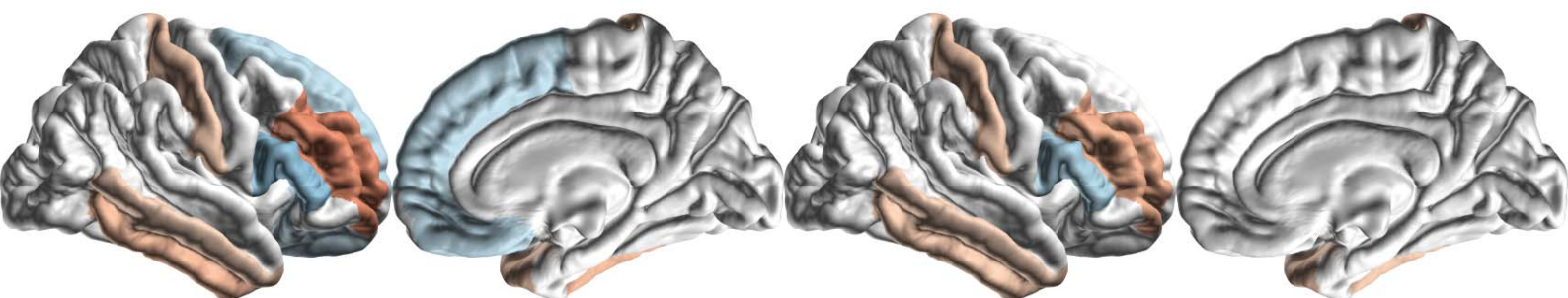


Rostral Middle Frontal SA: rs6682671

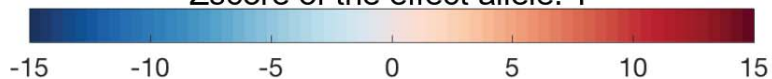


Corrected for total SA

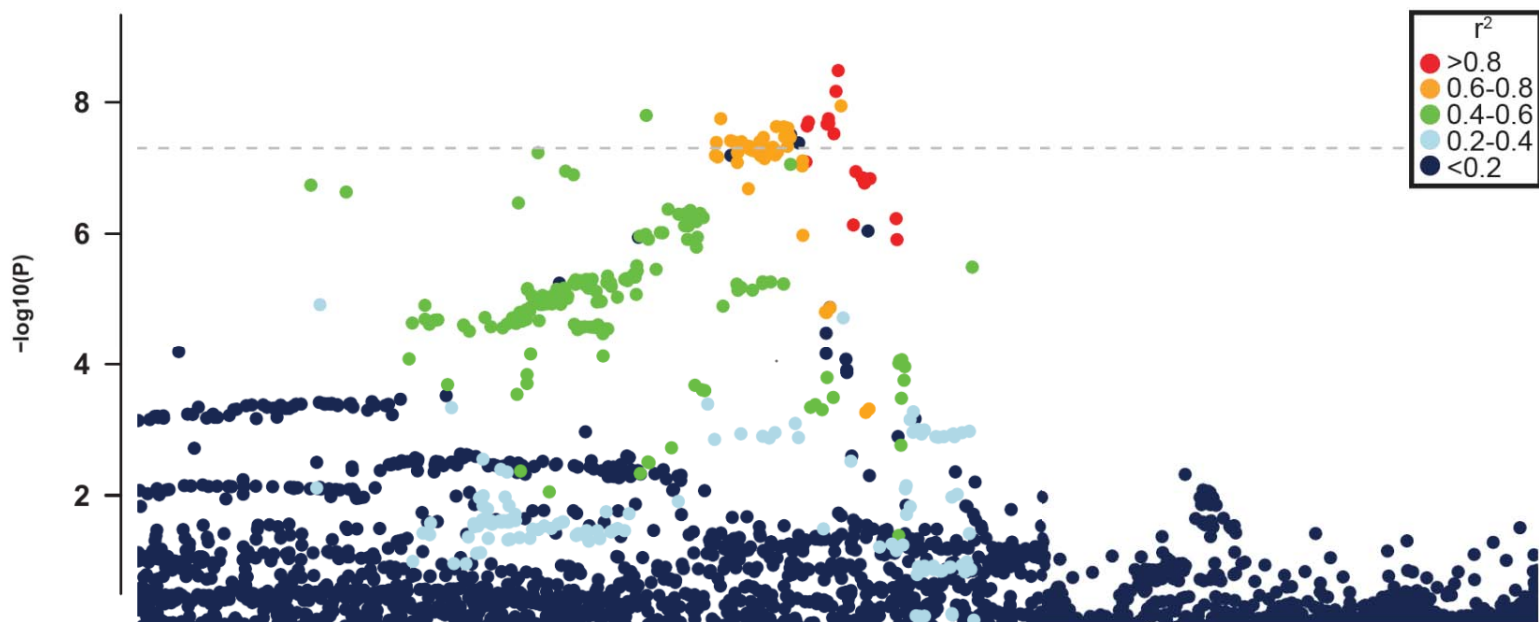
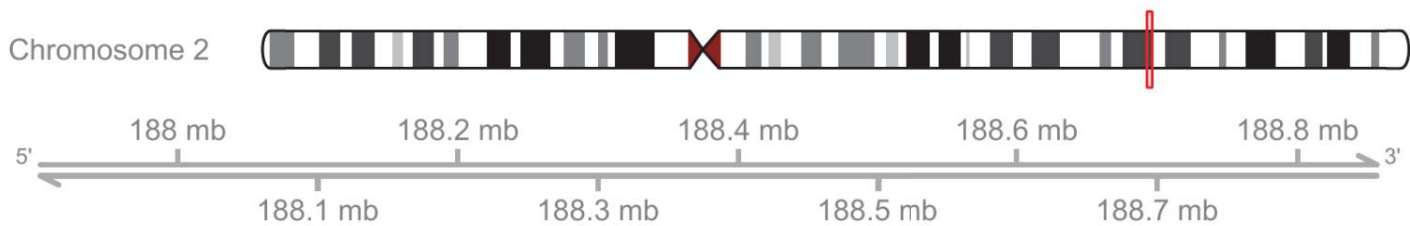
Uncorrected



Zscore of the effect allele: T



Rostral Middle Frontal SA: rs35612915

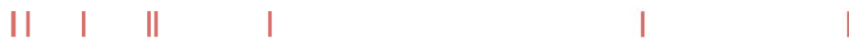


CALCRL

TFPI

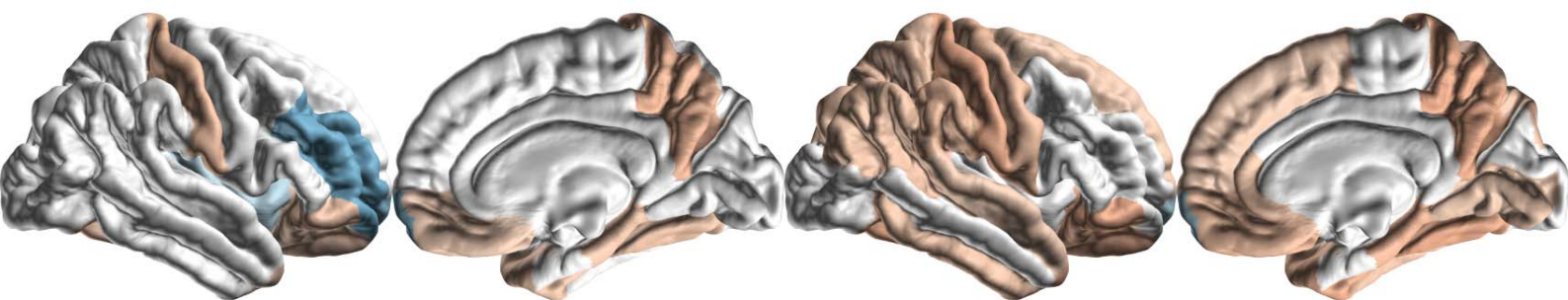
GZ>CP

CP>GZ



Corrected for total SA

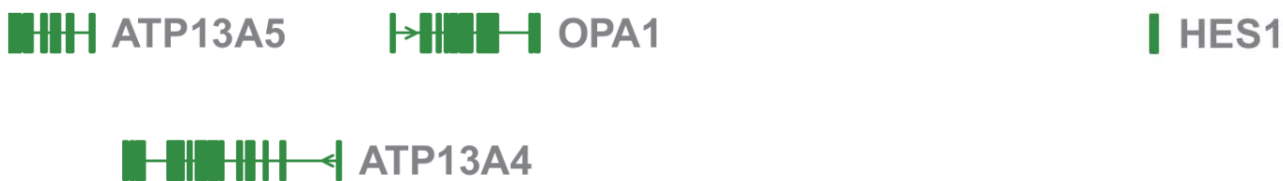
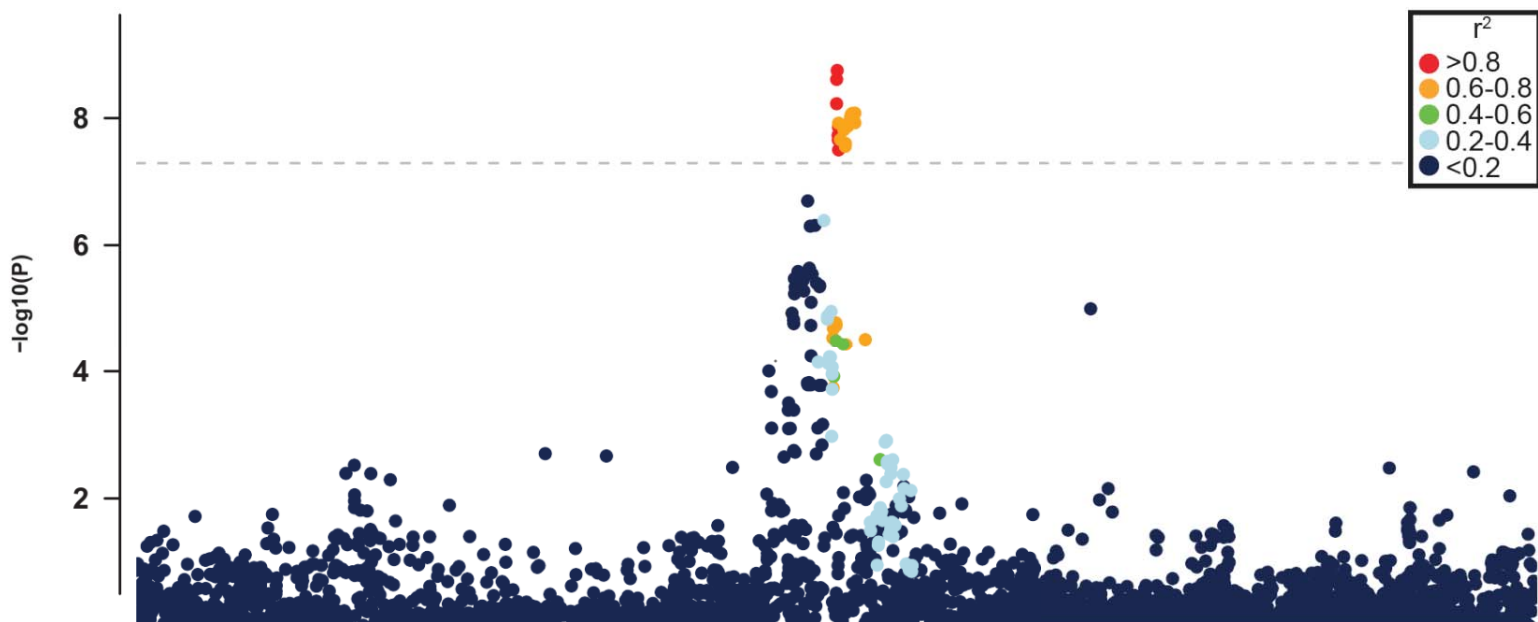
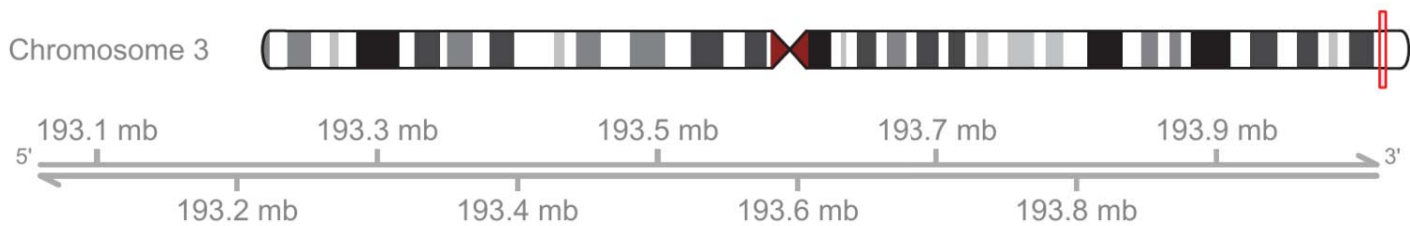
Uncorrected



Zscore of the effect allele: A

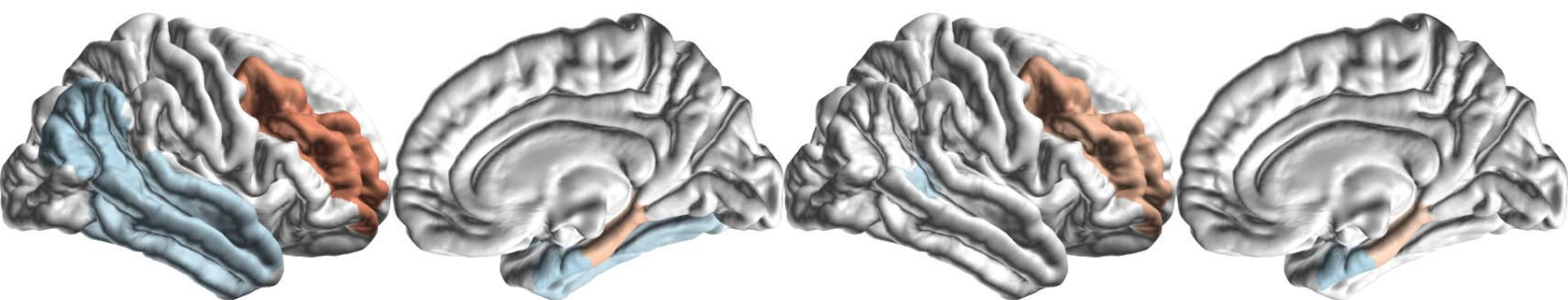


Rostral Middle Frontal SA: rs1165645

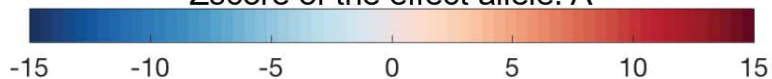


Corrected for total SA

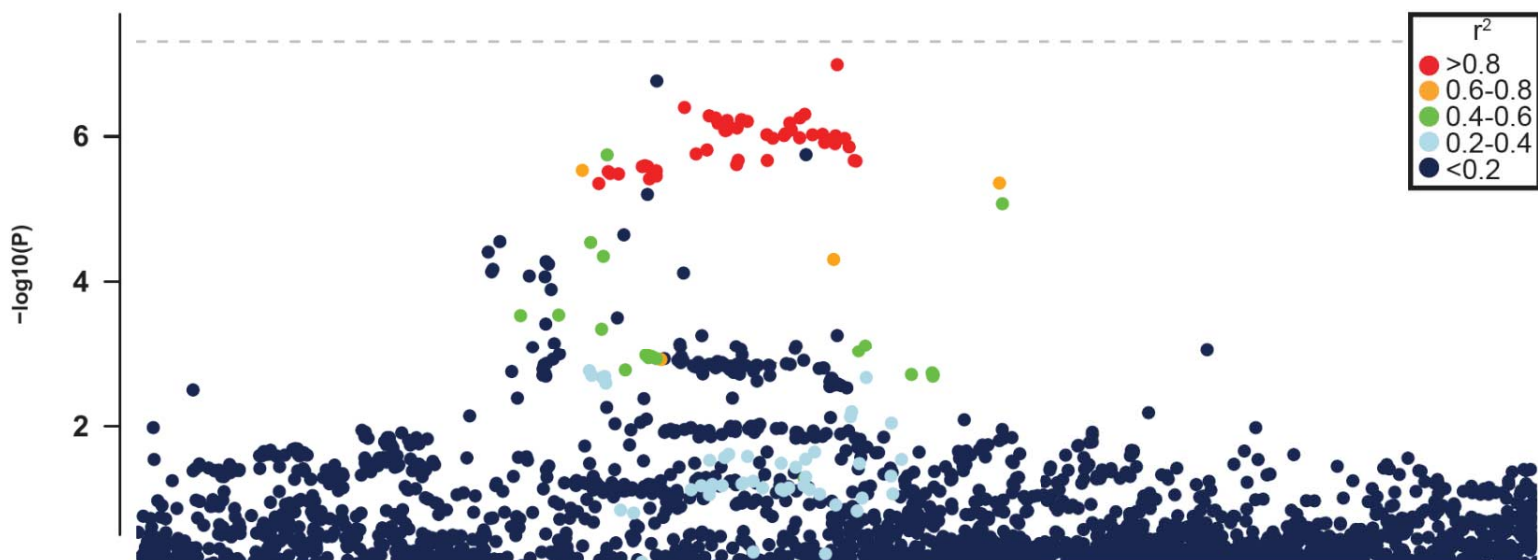
Uncorrected



Zscore of the effect allele: A



Rostral Middle Frontal SA: rs7731360

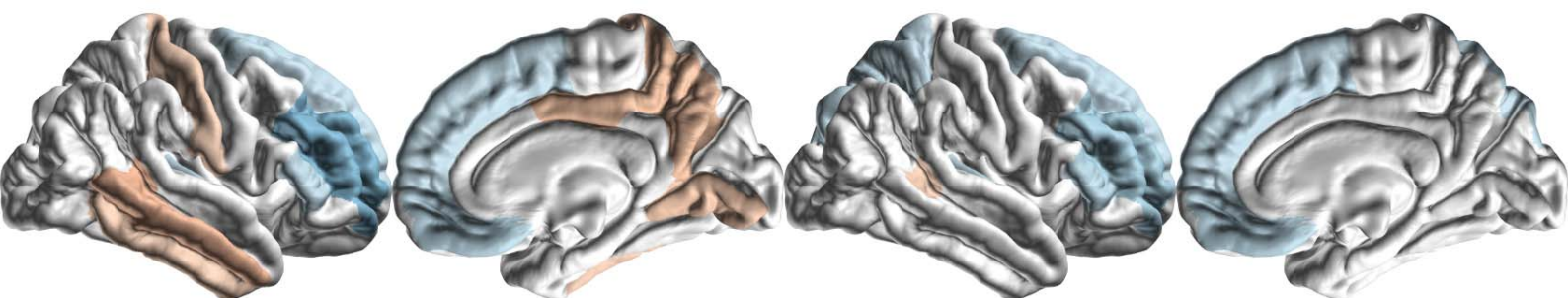


GZ>CP
CP>GZ



Corrected for total SA

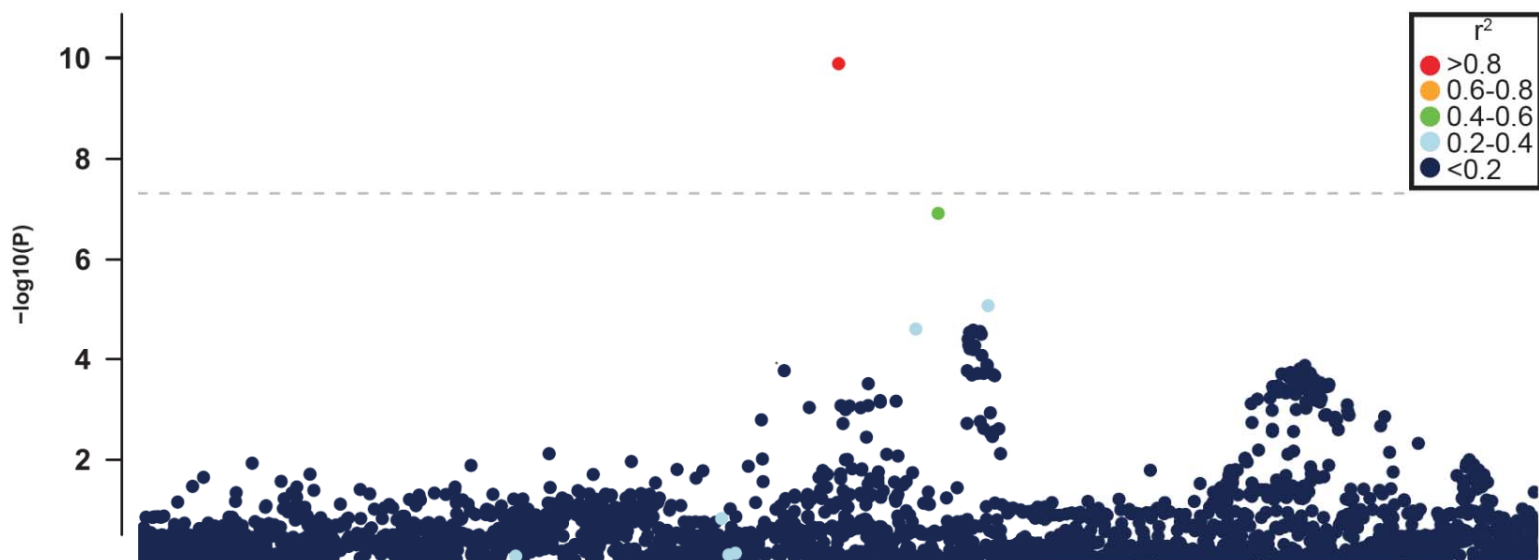
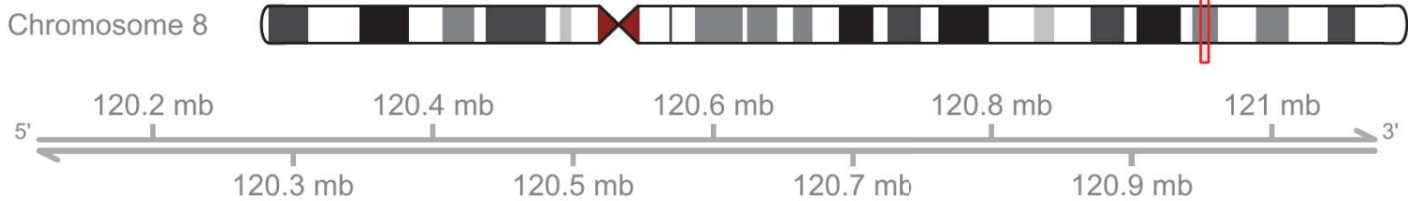
Uncorrected



Zscore of the effect allele: A

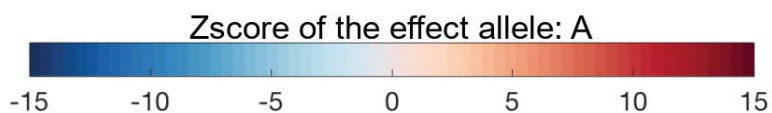
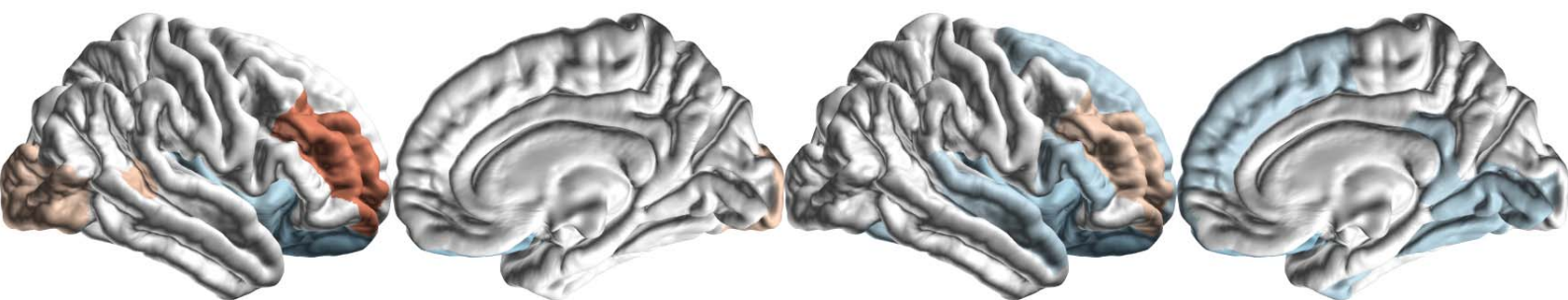


Rostral Middle Frontal SA: rs10283100

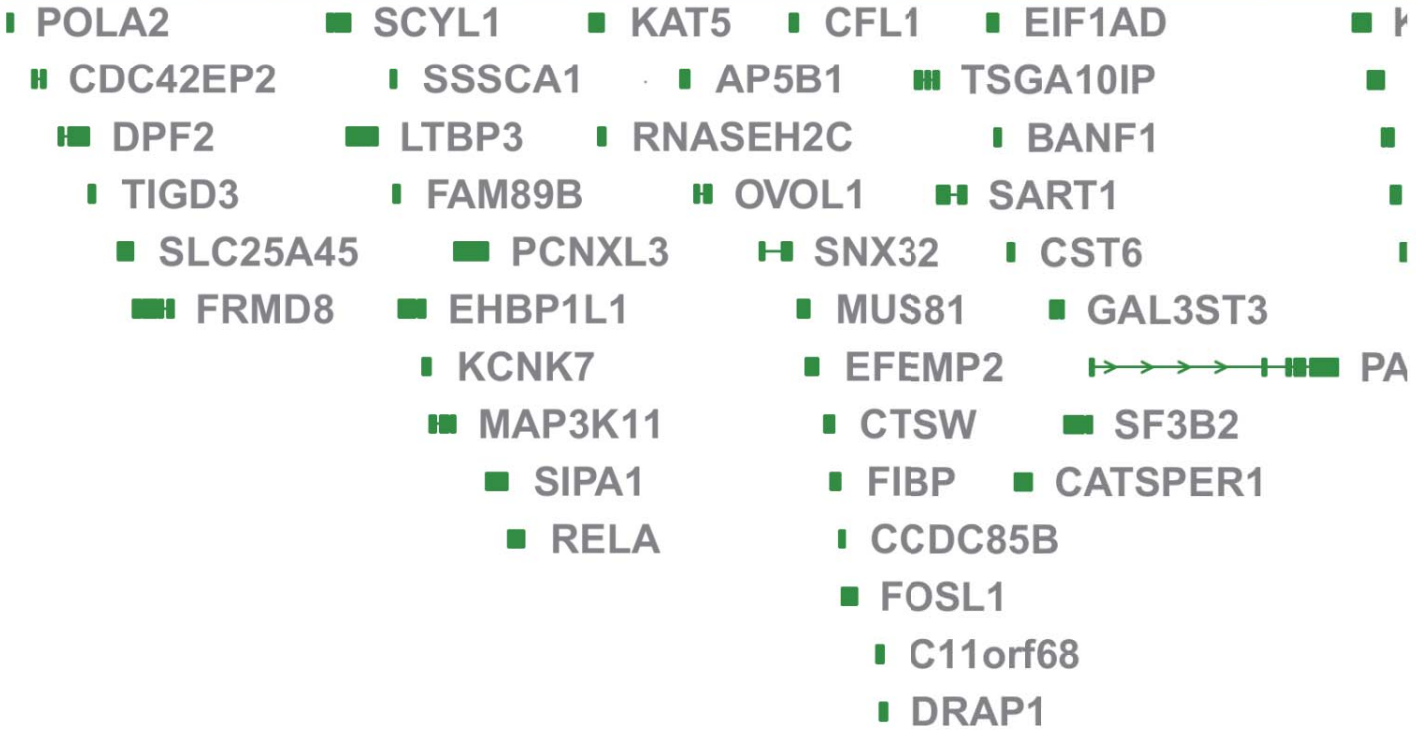
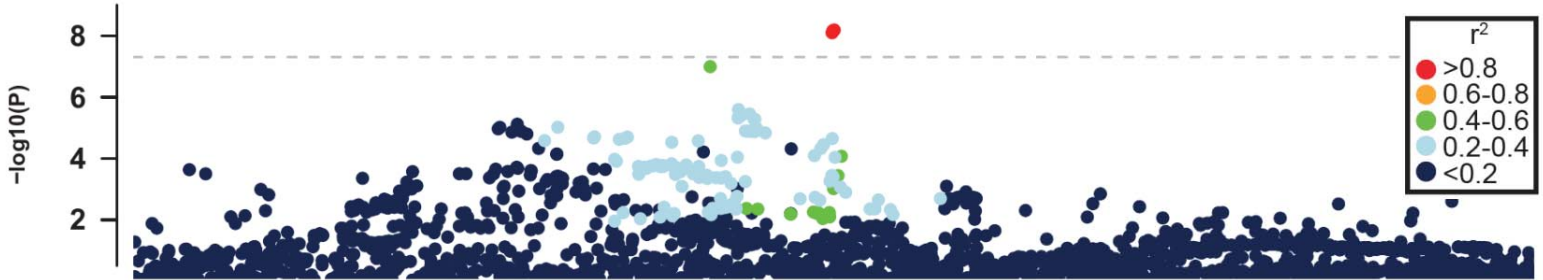
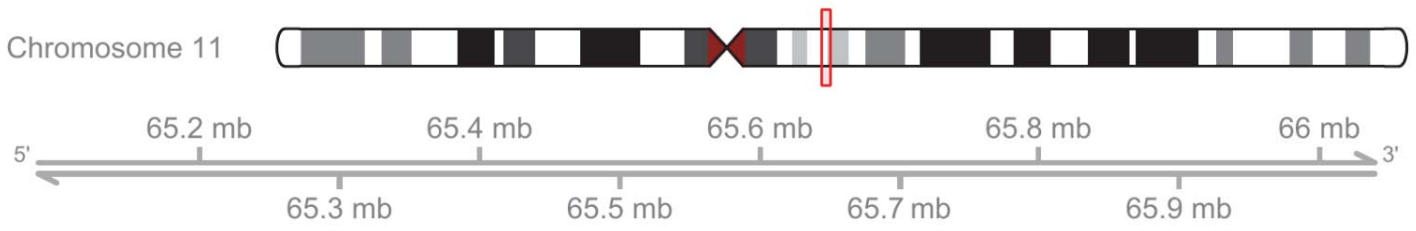


Corrected for total SA

Uncorrected



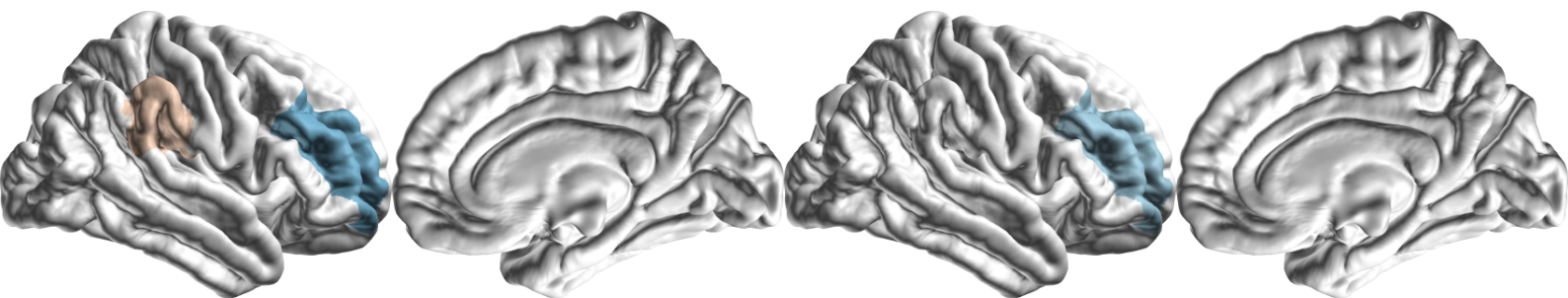
Rostral Middle Frontal SA: rs2276133



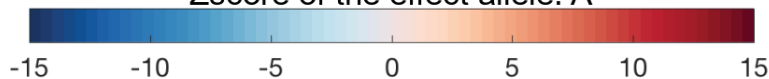
CP>GZZ>CP

Corrected for total SA

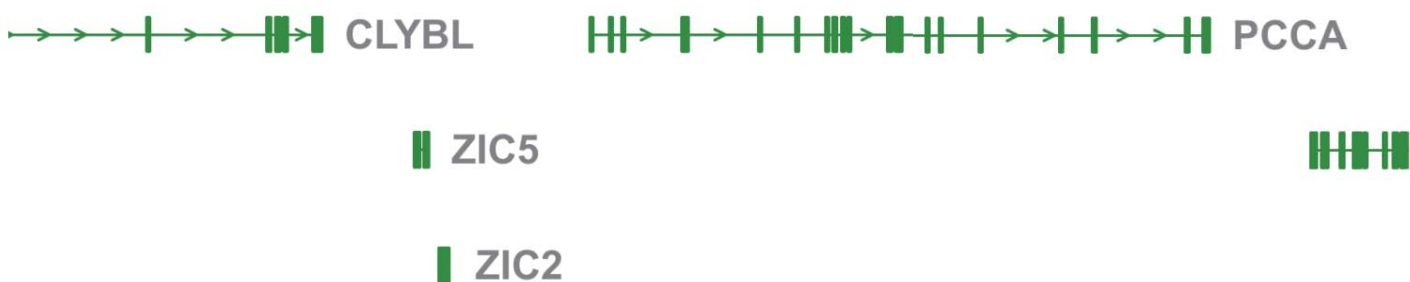
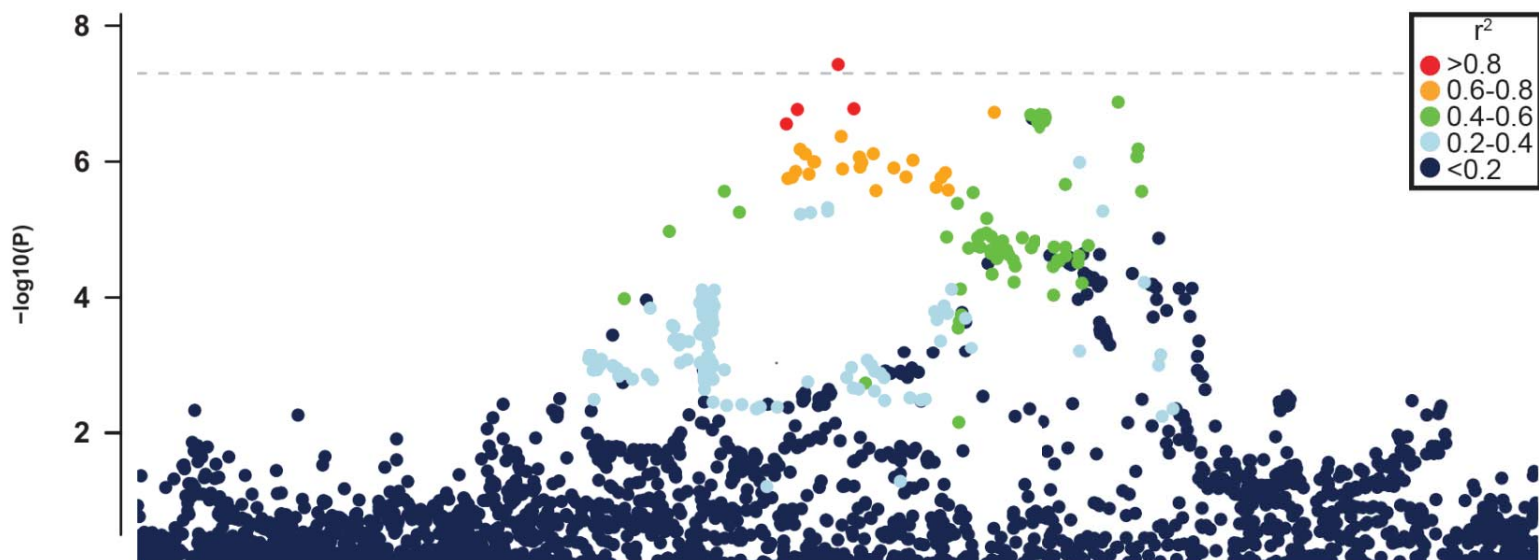
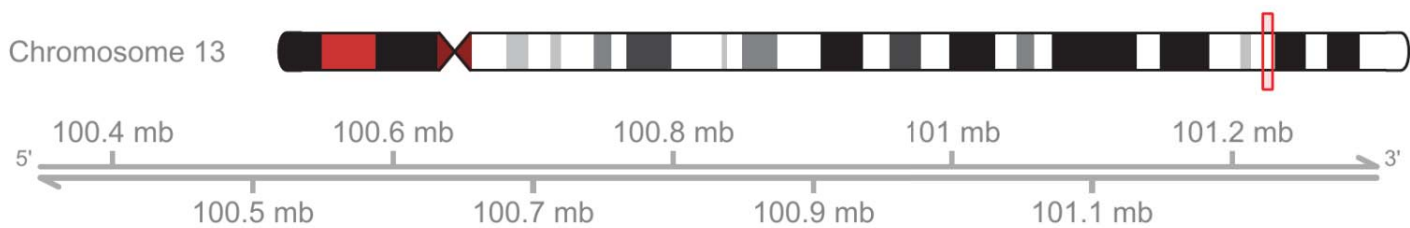
Uncorrected



Zscore of the effect allele: A

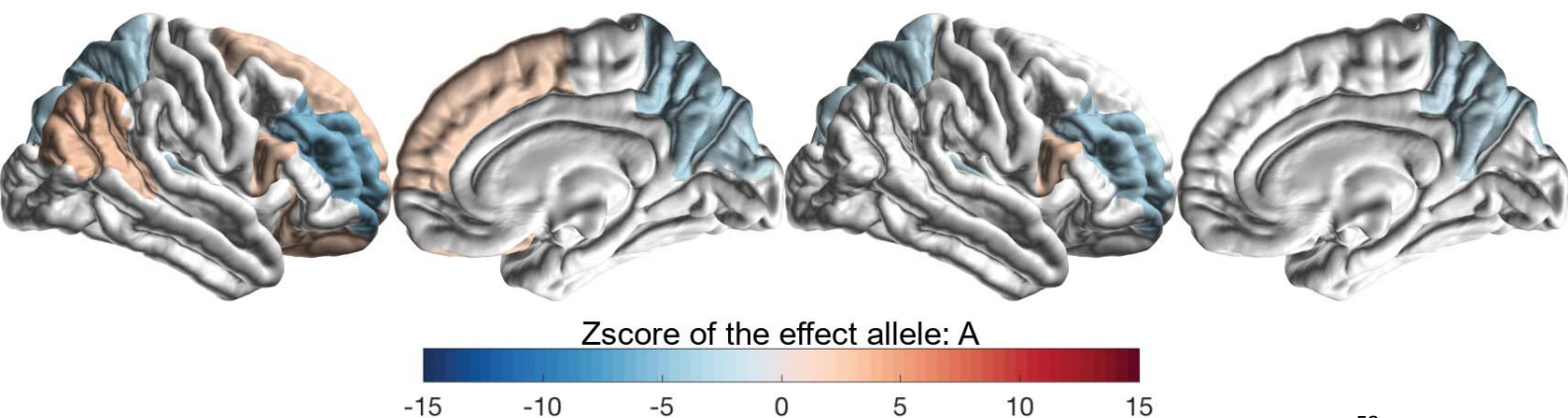


Rostral Middle Frontal SA: rs4772268

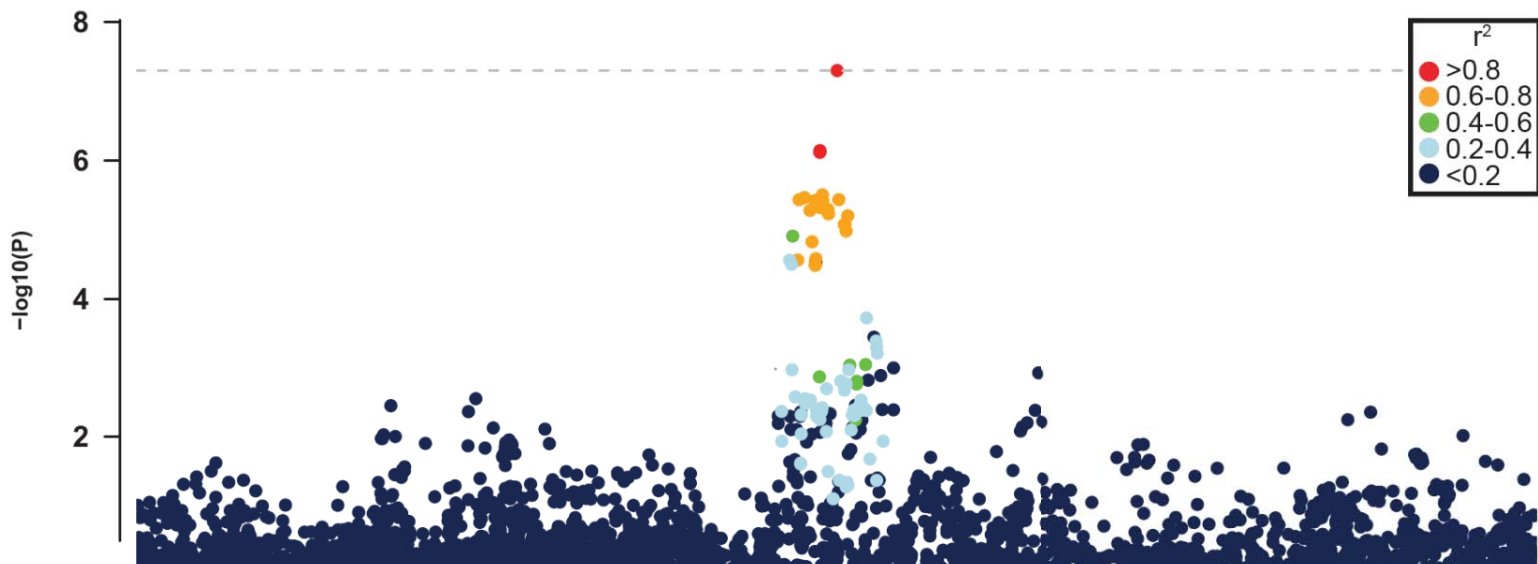
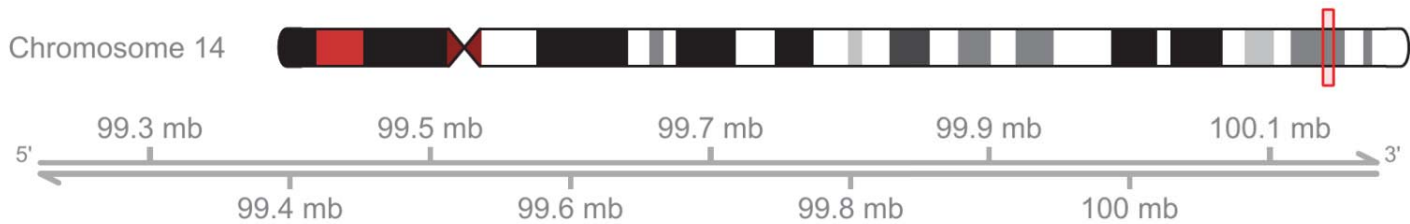


Corrected for total SA

Uncorrected



Rostral Middle Frontal SA: rs1257415

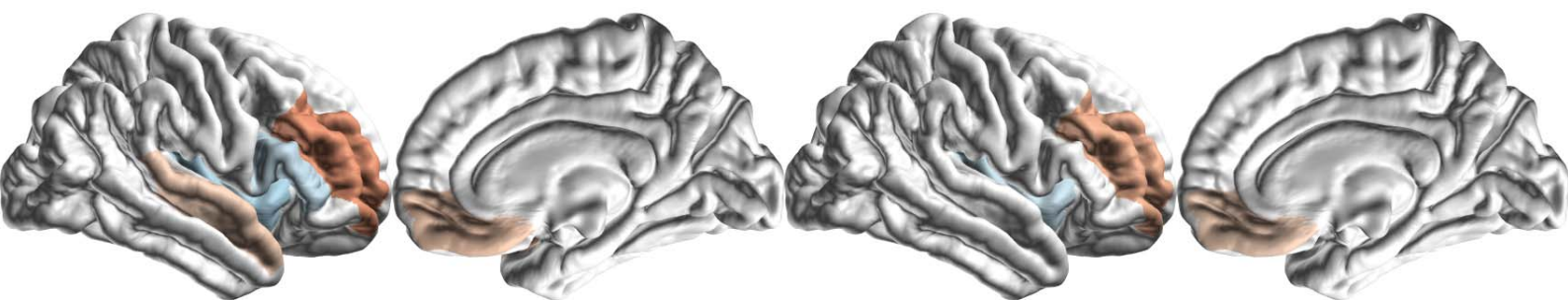


CP>GZ
GZ>CP

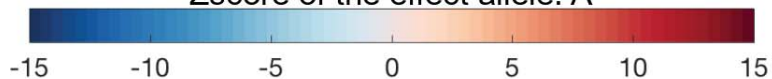


Corrected for total SA

Uncorrected

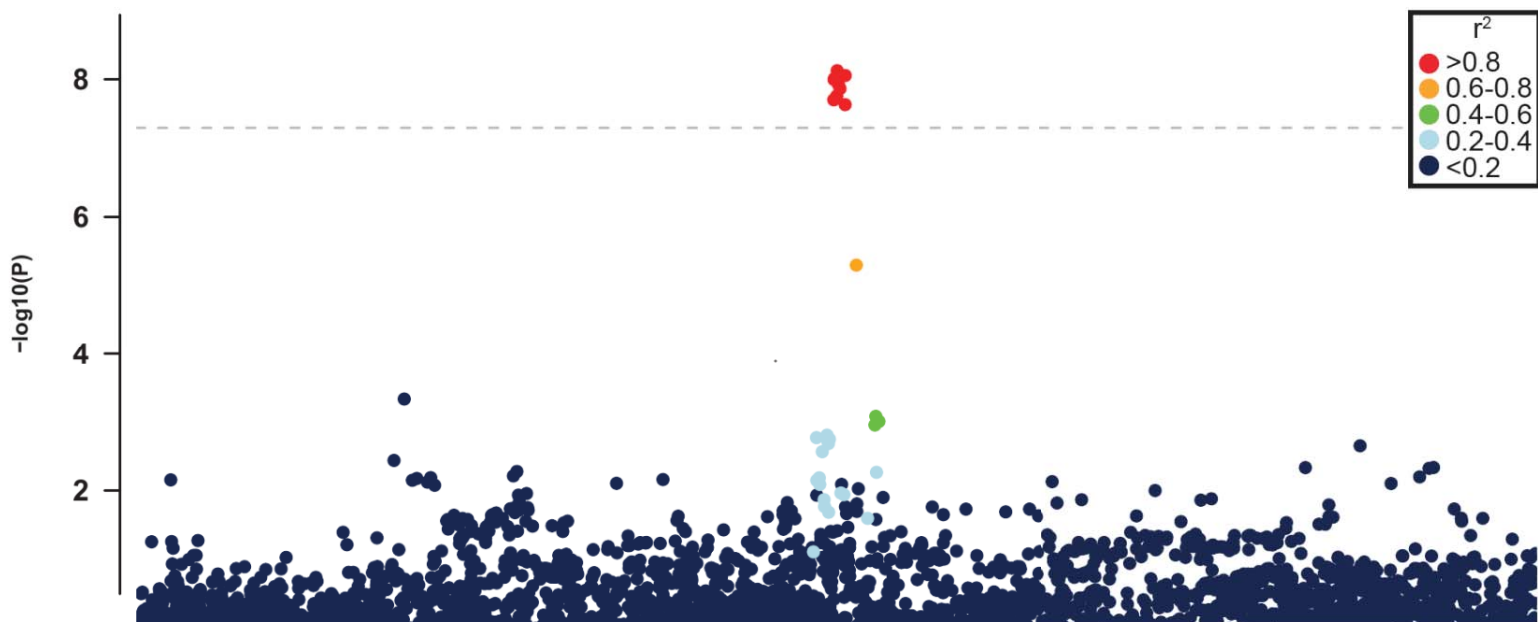
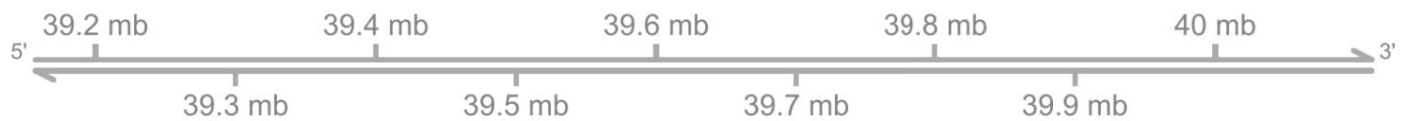


Zscore of the effect allele: A



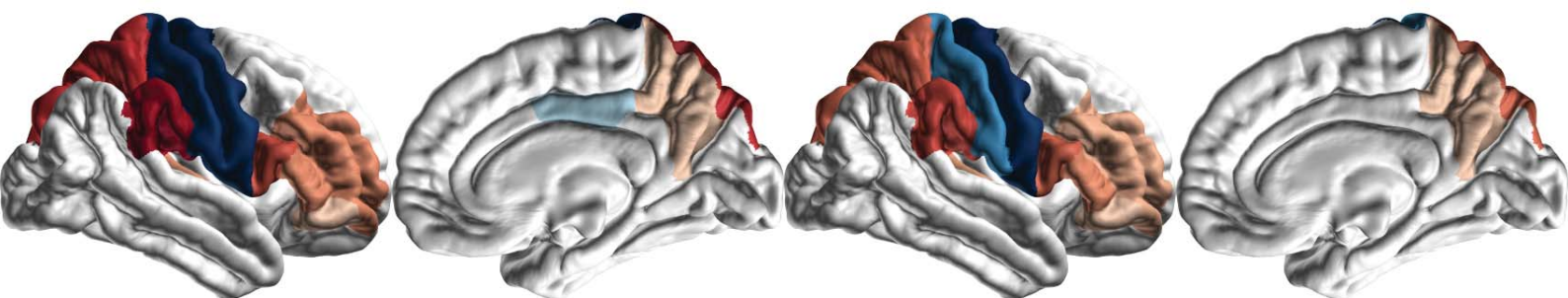
Rostral Middle Frontal SA: rs1080066

Chromosome 15

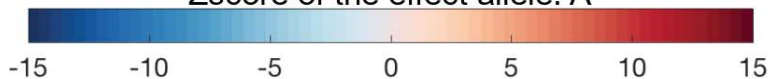


Corrected for total SA

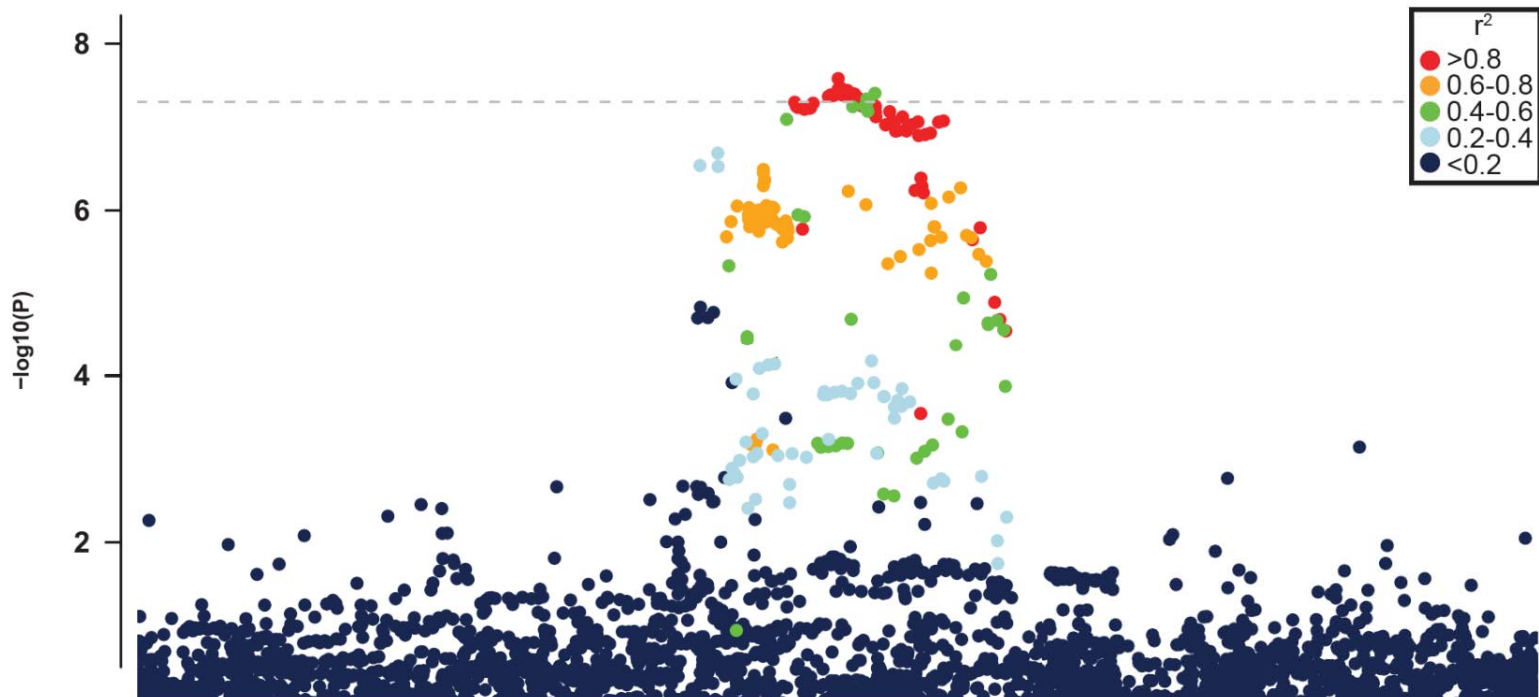
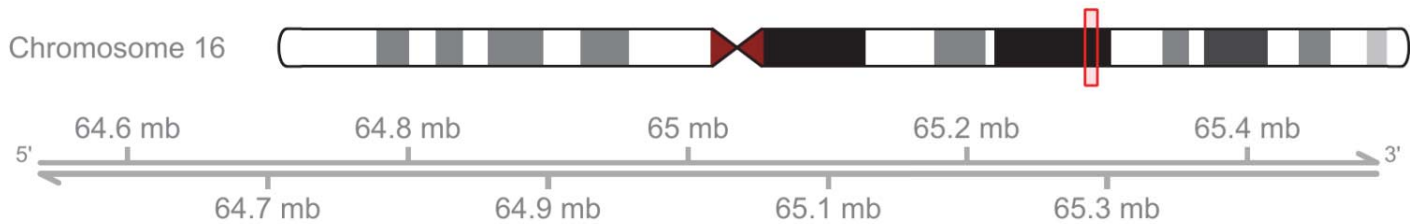
Uncorrected



Zscore of the effect allele: A



Rostral Middle Frontal SA: rs40115

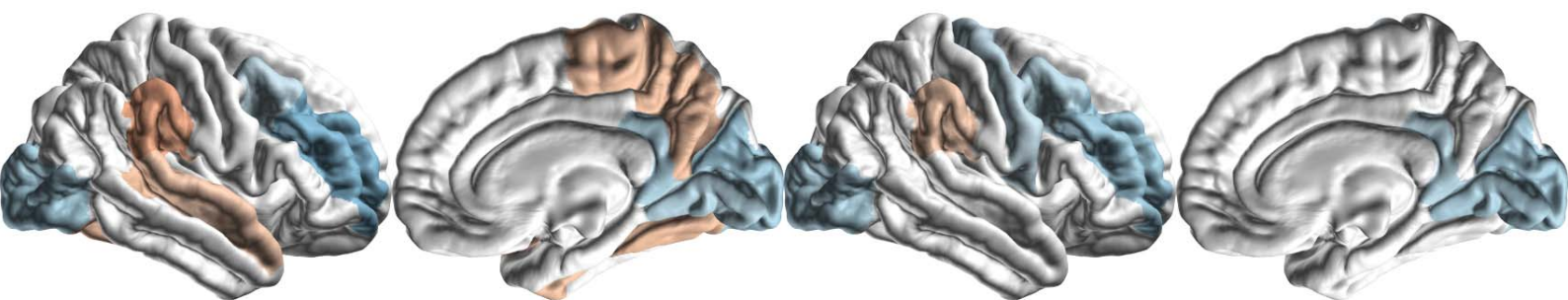


GZ>CP

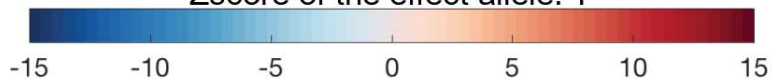
CP>GZ

Corrected for total SA

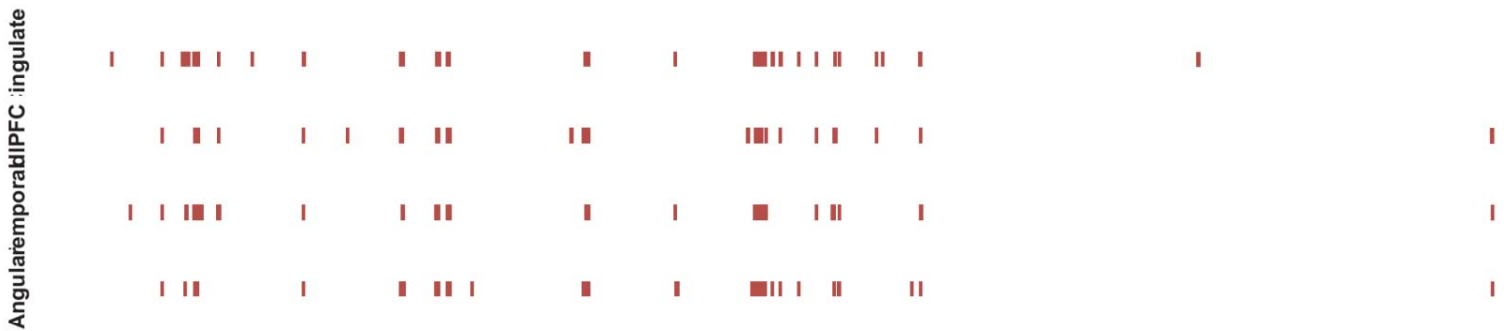
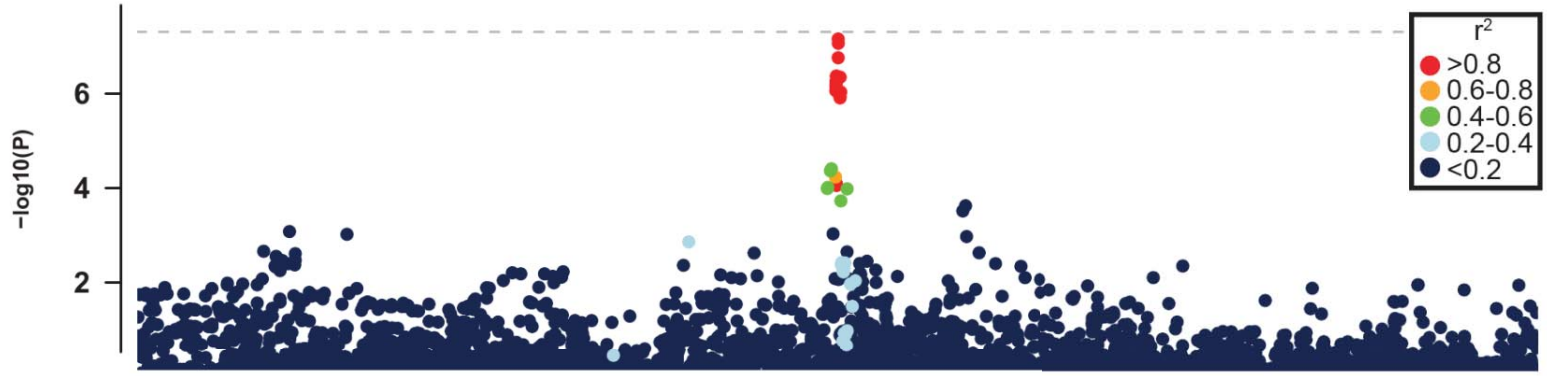
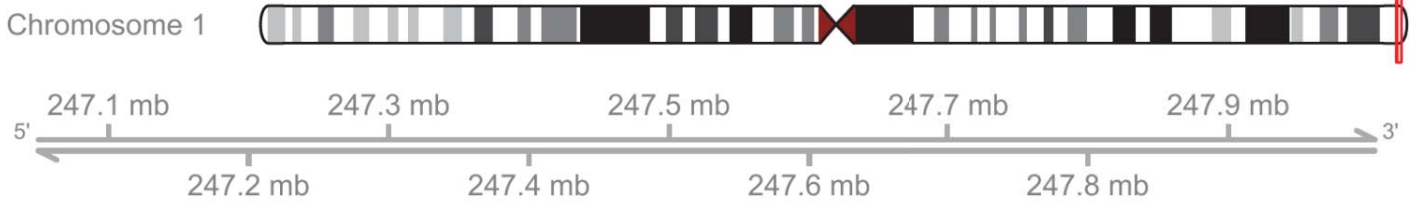
Uncorrected



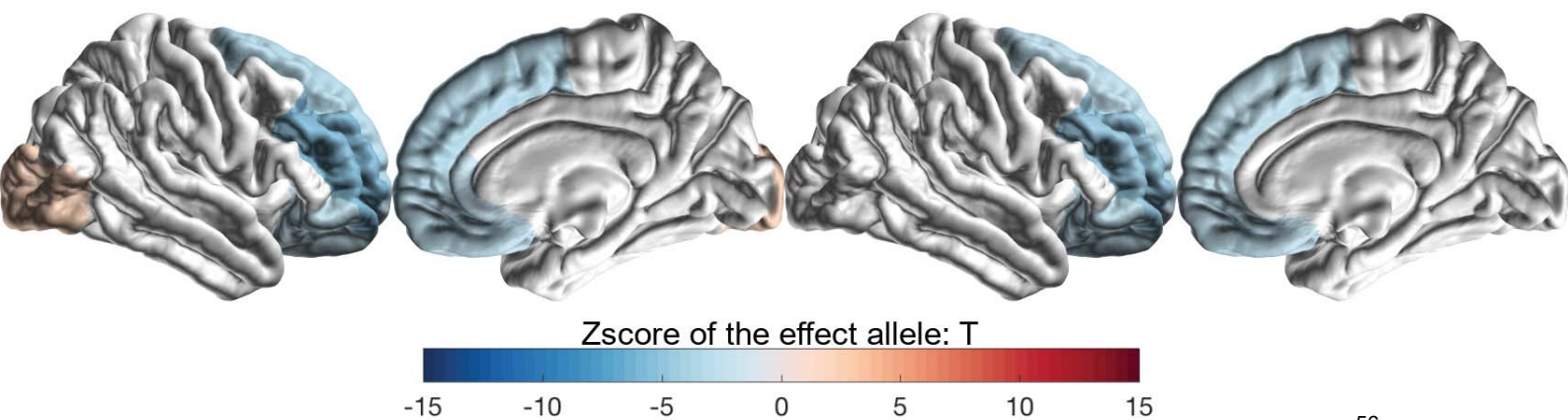
Zscore of the effect allele: T



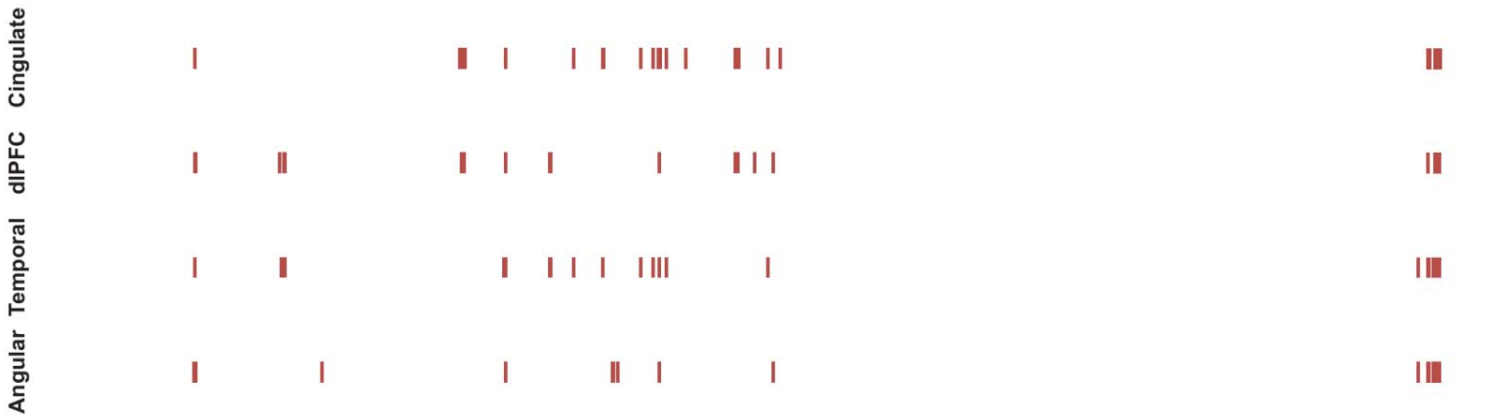
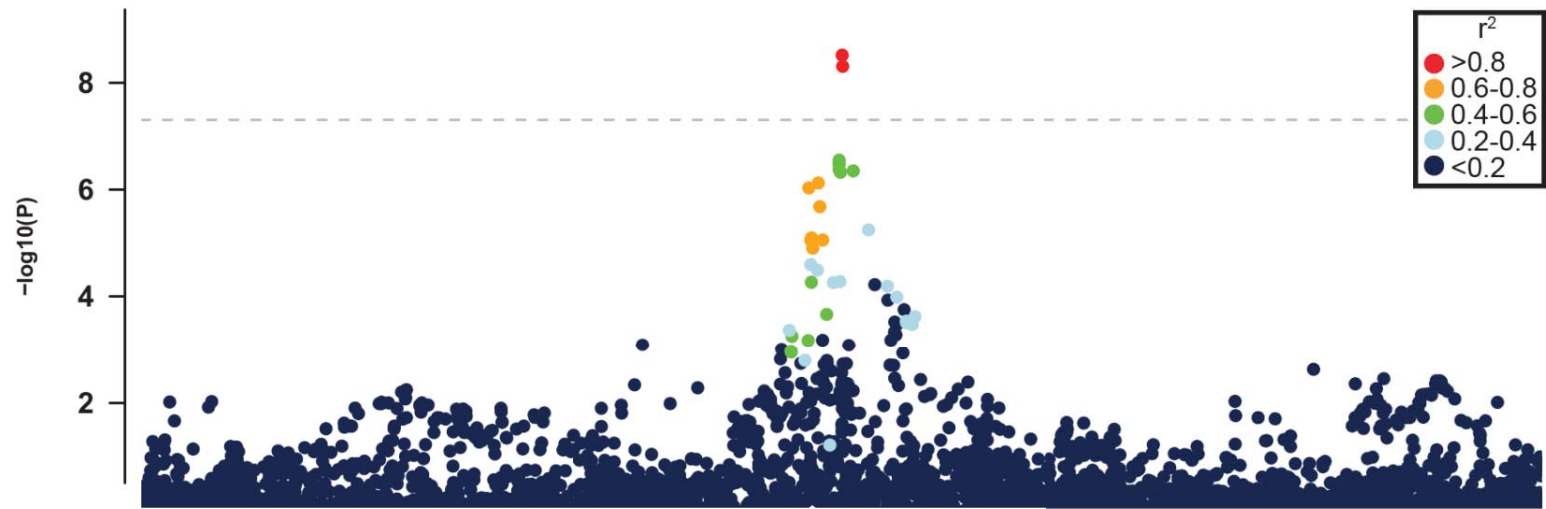
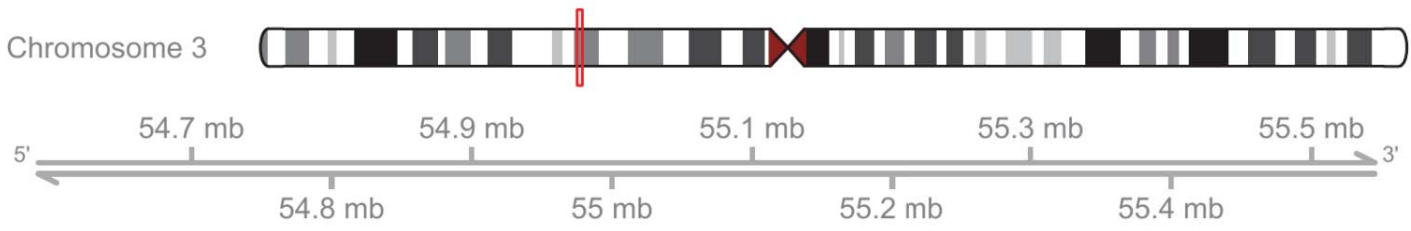
Rostral Middle Frontal TH: rs6671321



Uncorrected

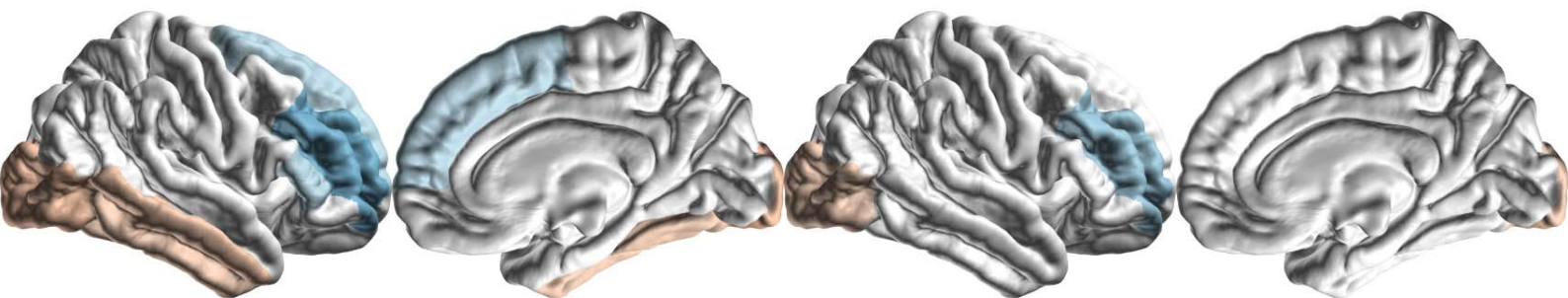


Rostral Middle Frontal TH: rs5008703

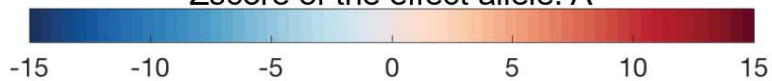


Corrected for average TH

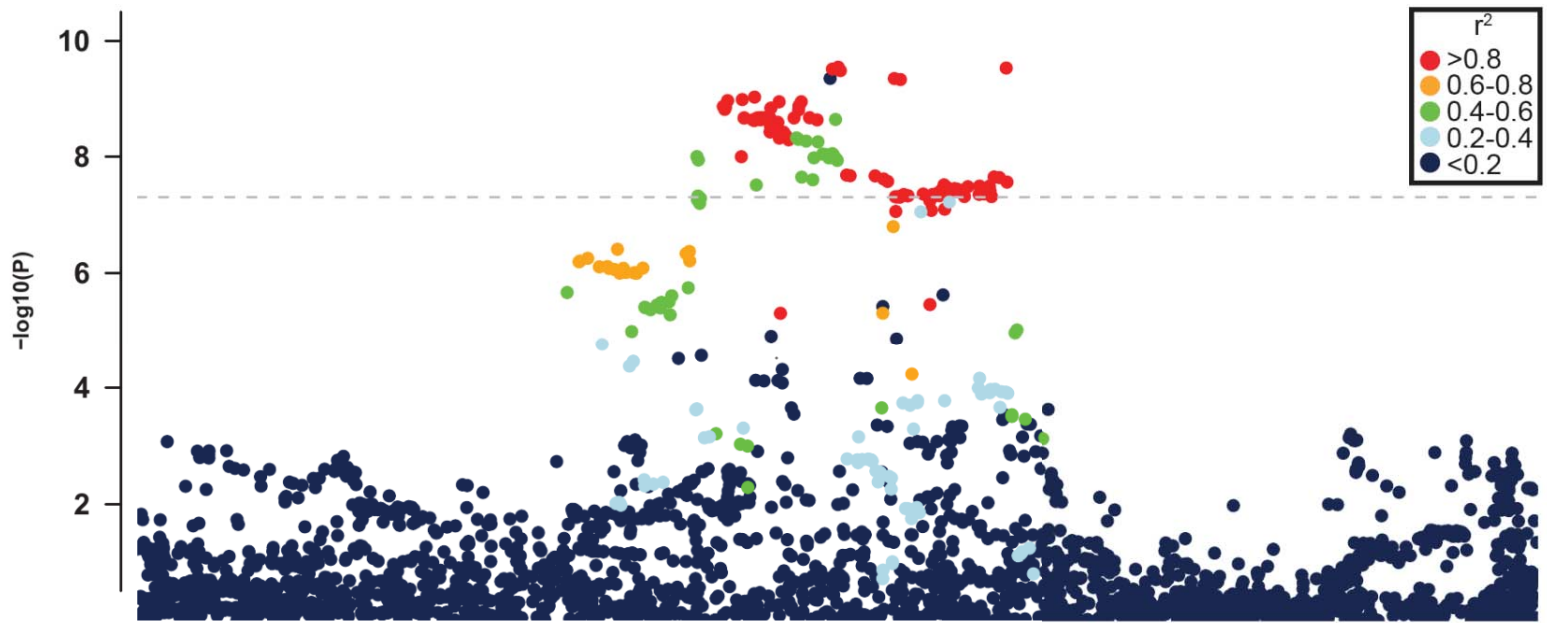
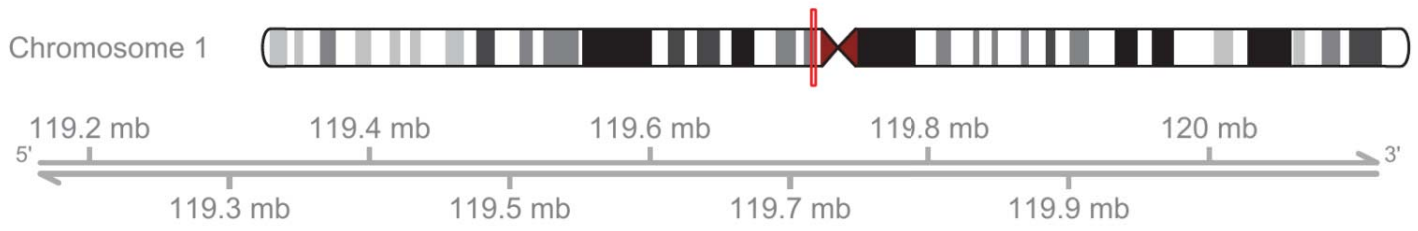
Uncorrected



Zscore of the effect allele: A



Pars Orbitalis SA: rs12064109



TBX15

HAO2 HSD3

WARS2

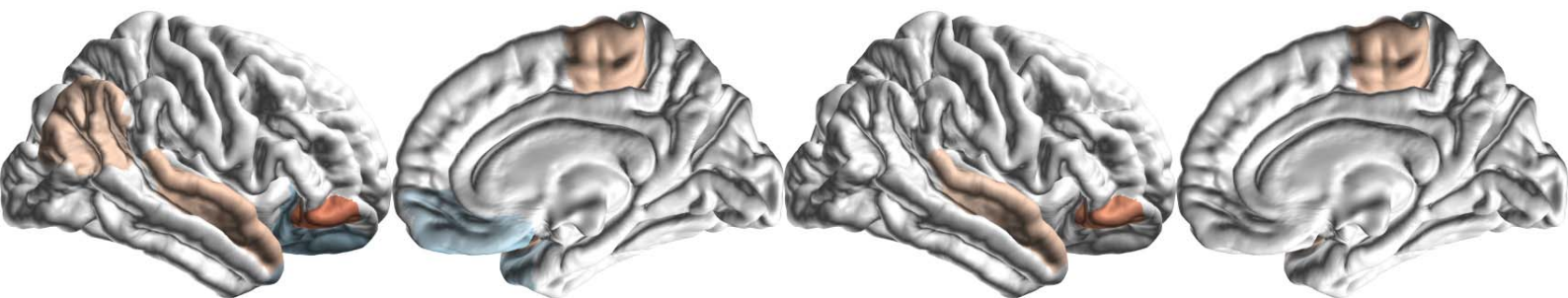
HSD3B2

GZ>CP

CP>GZ

Corrected for total SA

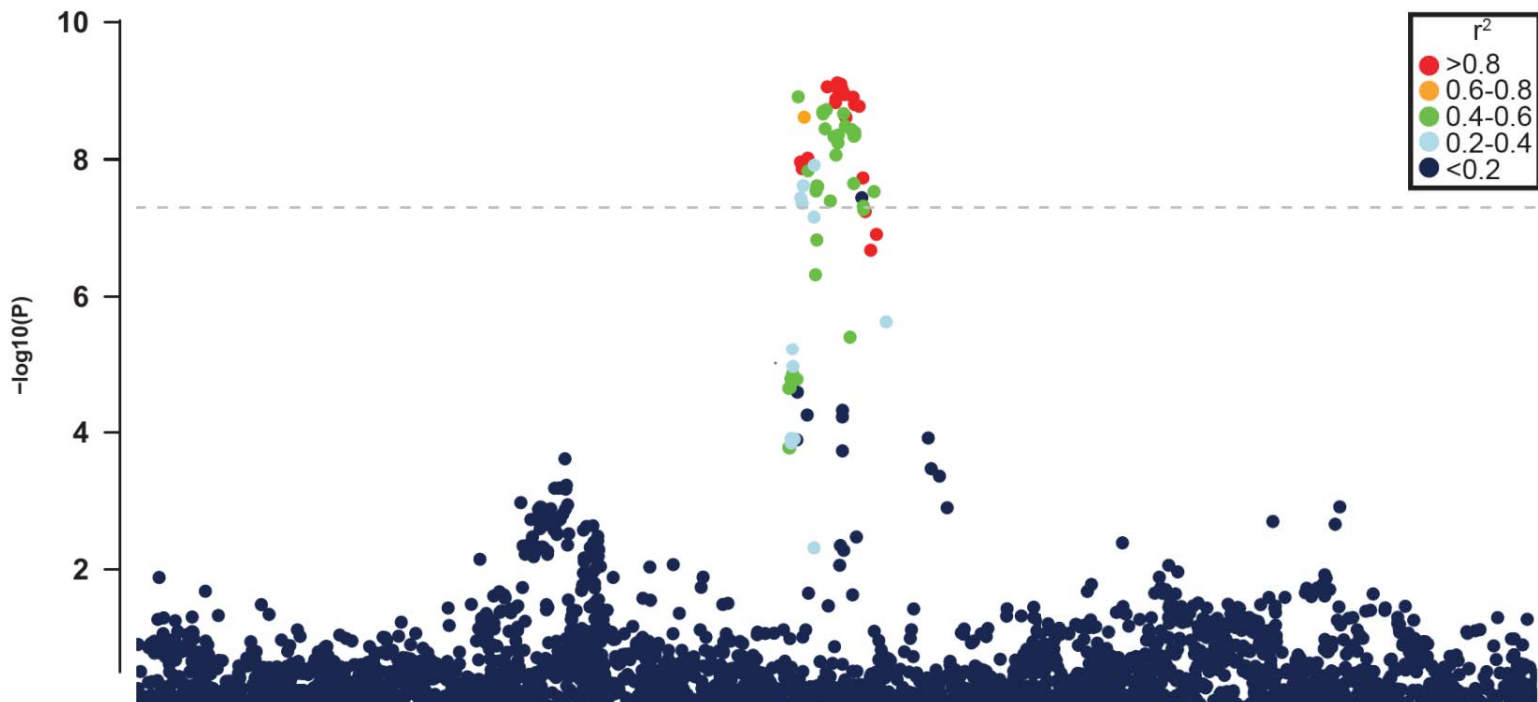
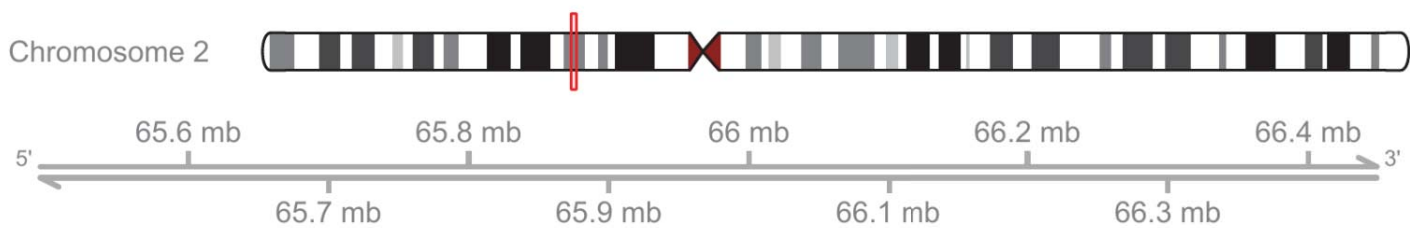
Uncorrected



Zscore of the effect allele: T



Pars Orbitalis SA: rs2287283



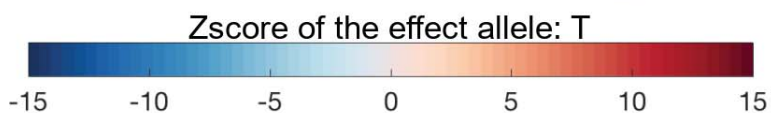
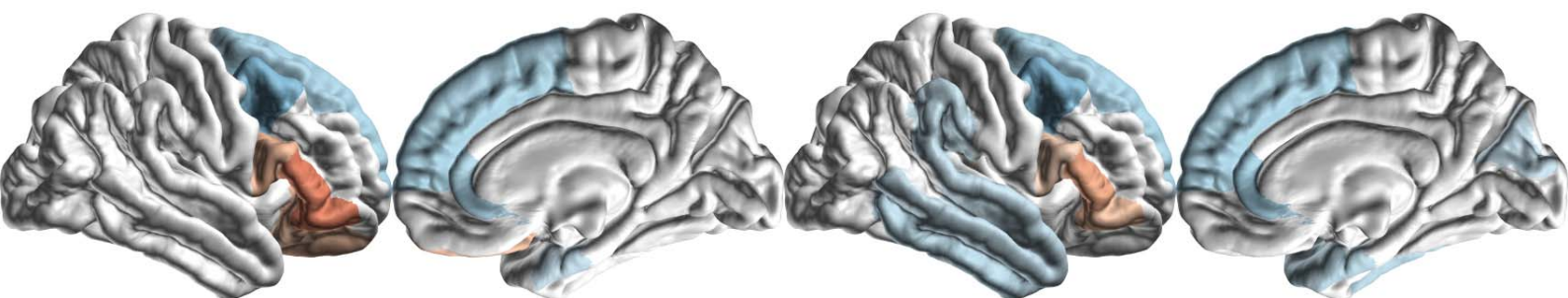
AC1R2 ← SPRED2

GZ>CP

CP>GZ

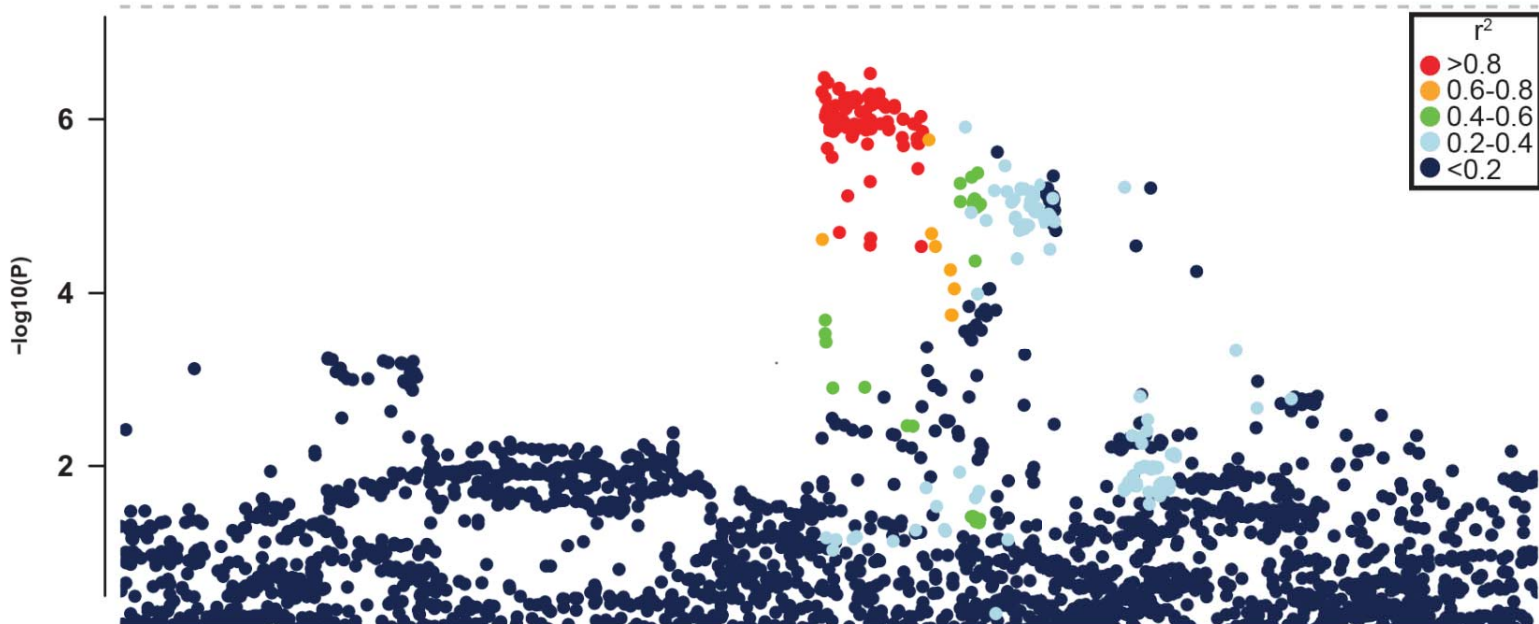
Corrected for total SA

Uncorrected



Pars Orbitalis SA: rs6551313

Chromosome 3



HTR1F GG G ZNF654

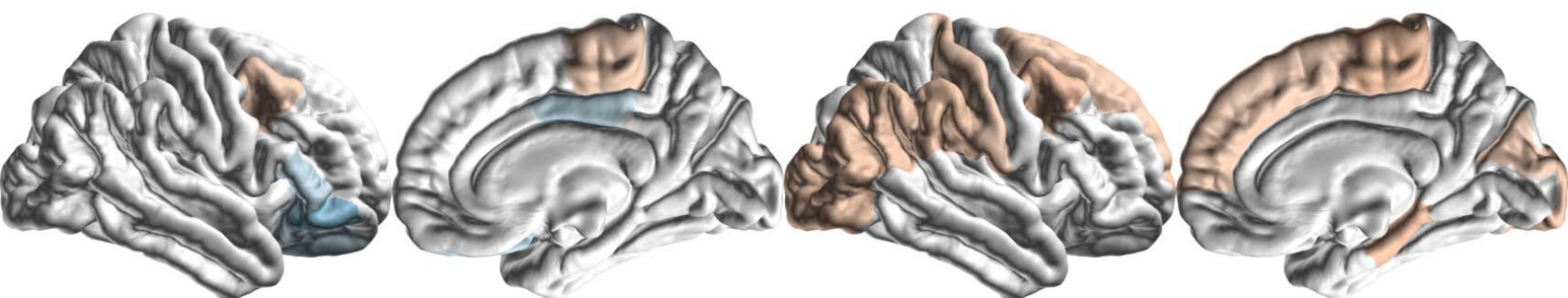
C3orf38

GZ>CP

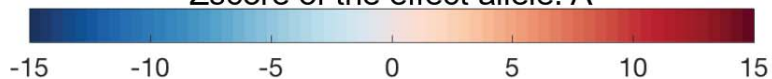
CP>GZ

Corrected for total SA

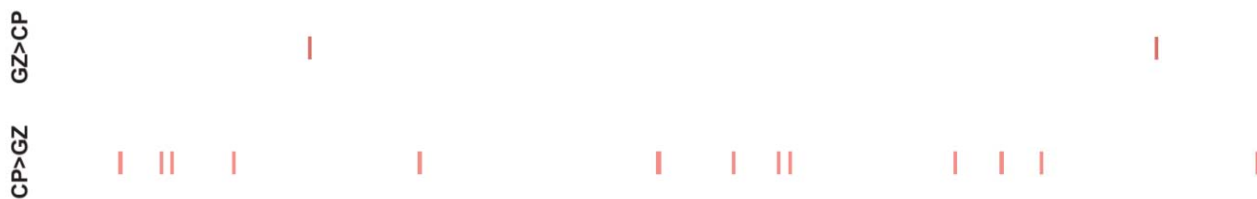
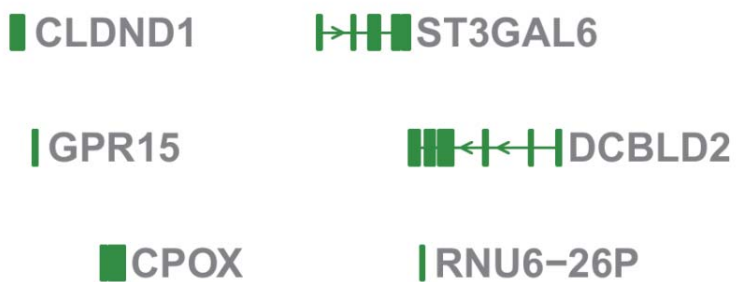
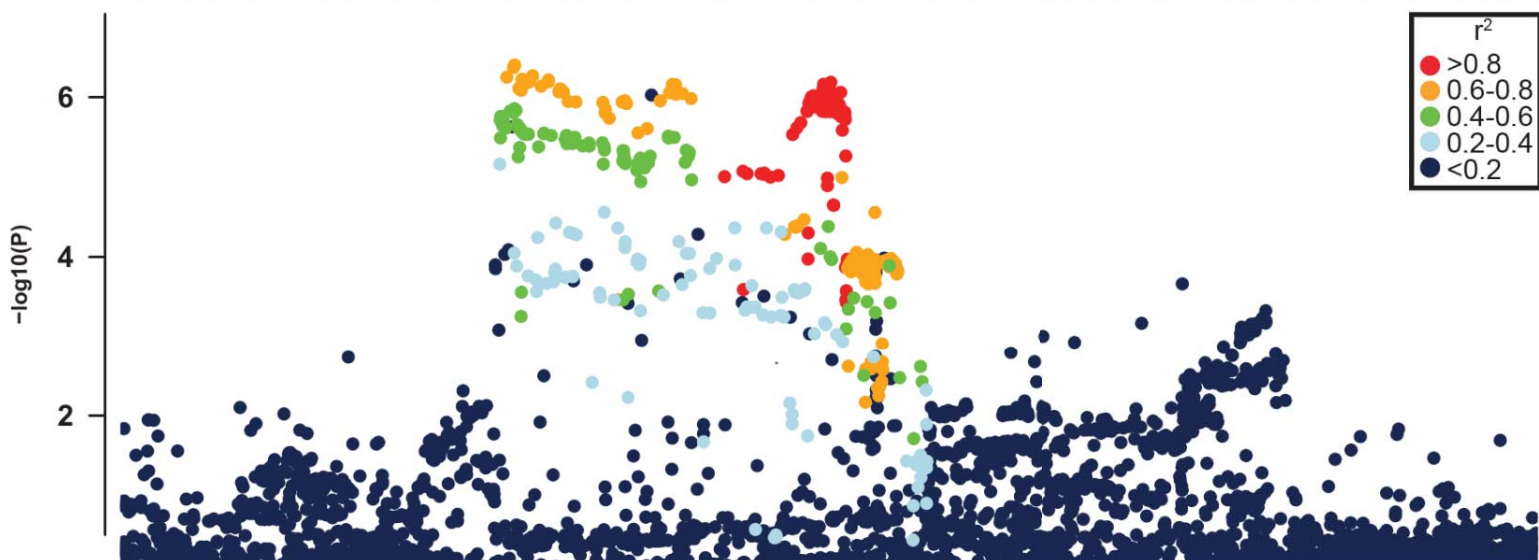
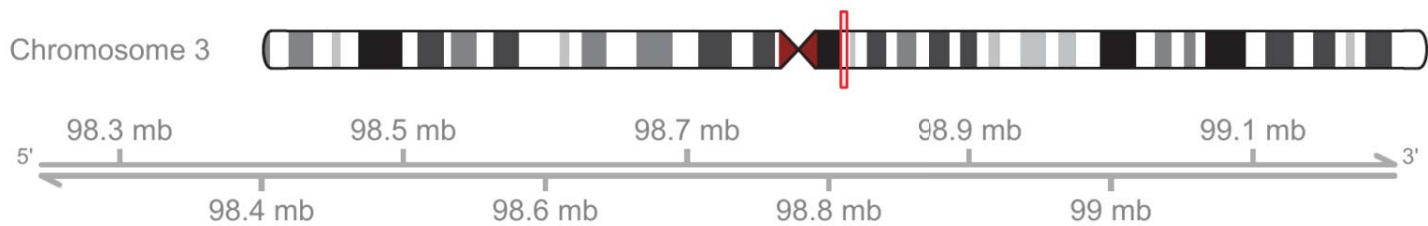
Uncorrected



Zscore of the effect allele: A

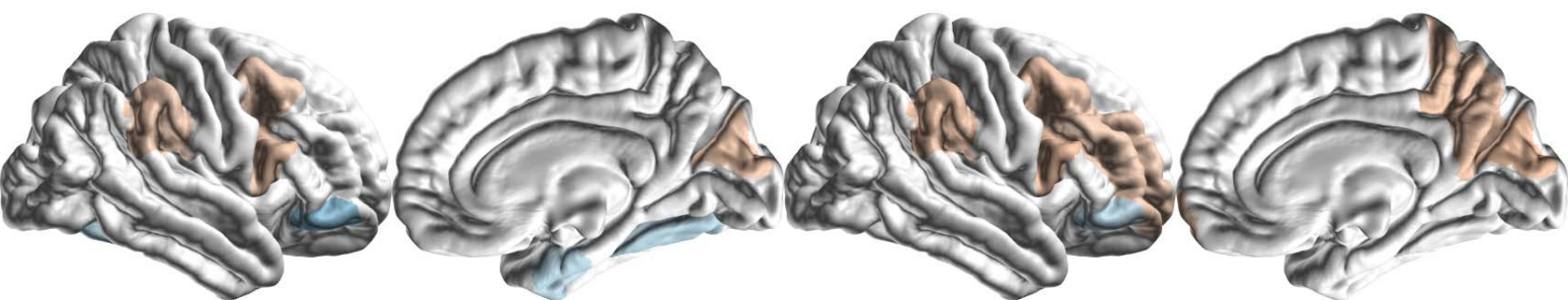


Pars Orbitalis SA: rs2262601



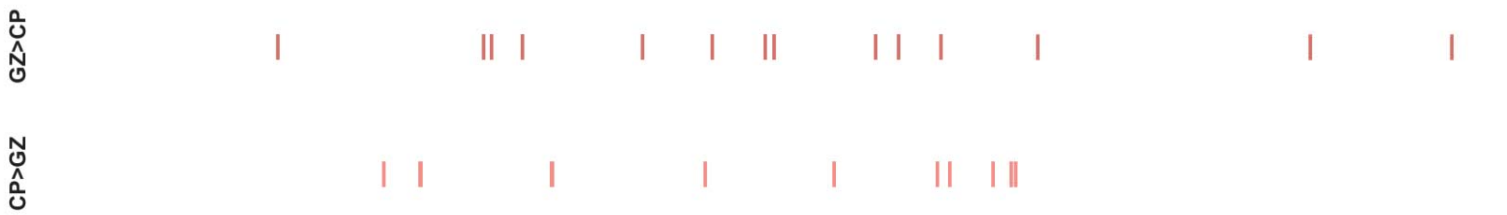
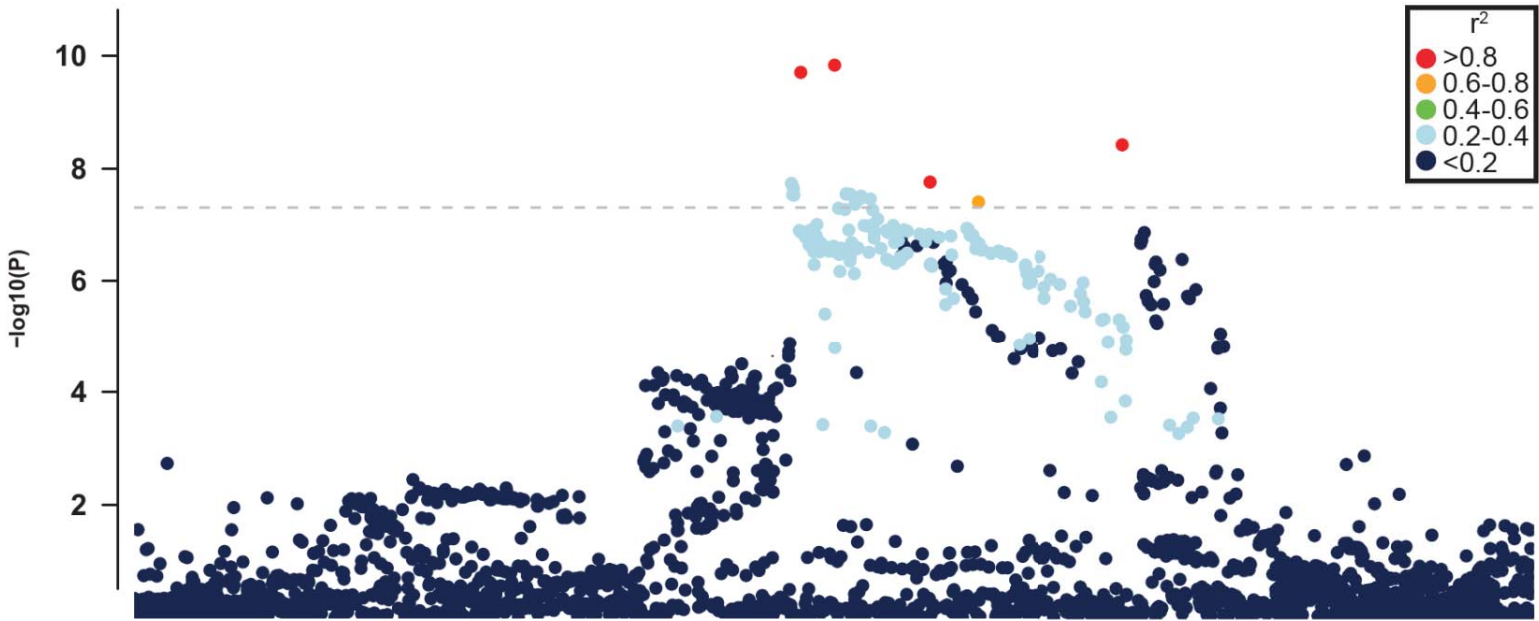
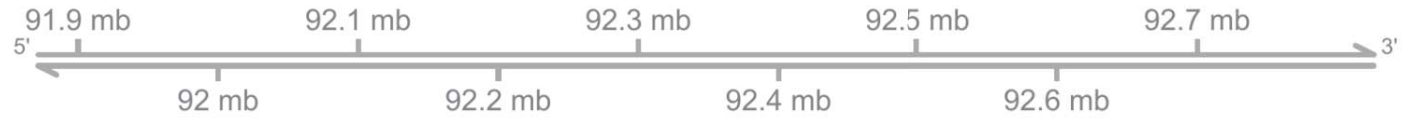
Corrected for total SA

Uncorrected



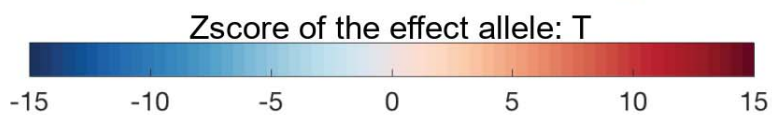
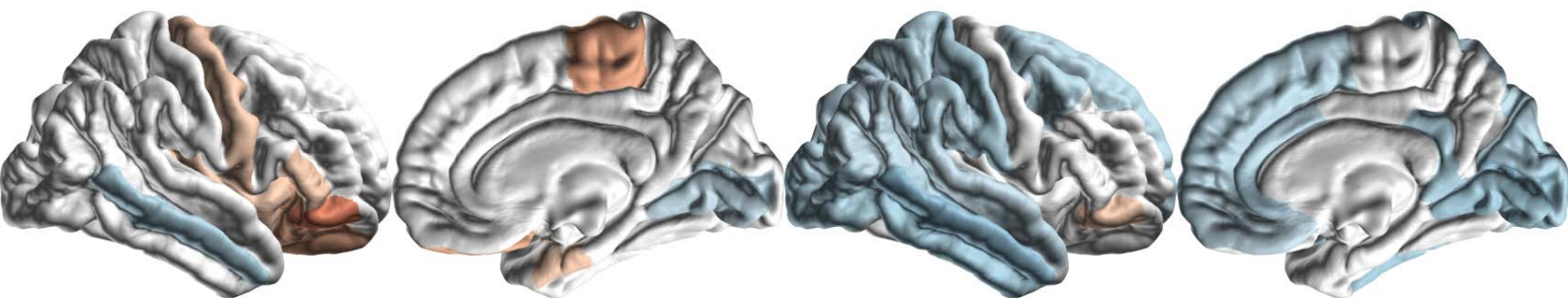
Pars Orbitalis SA: rs61901866

Chromosome 11

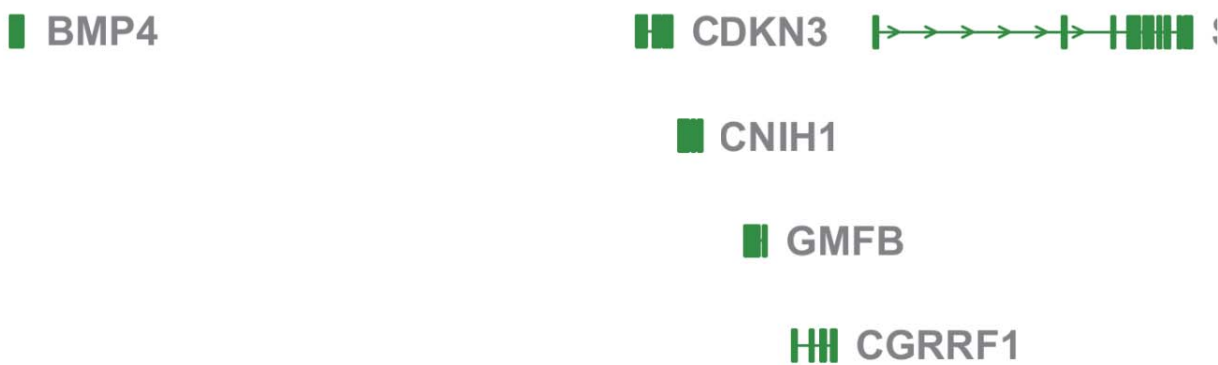
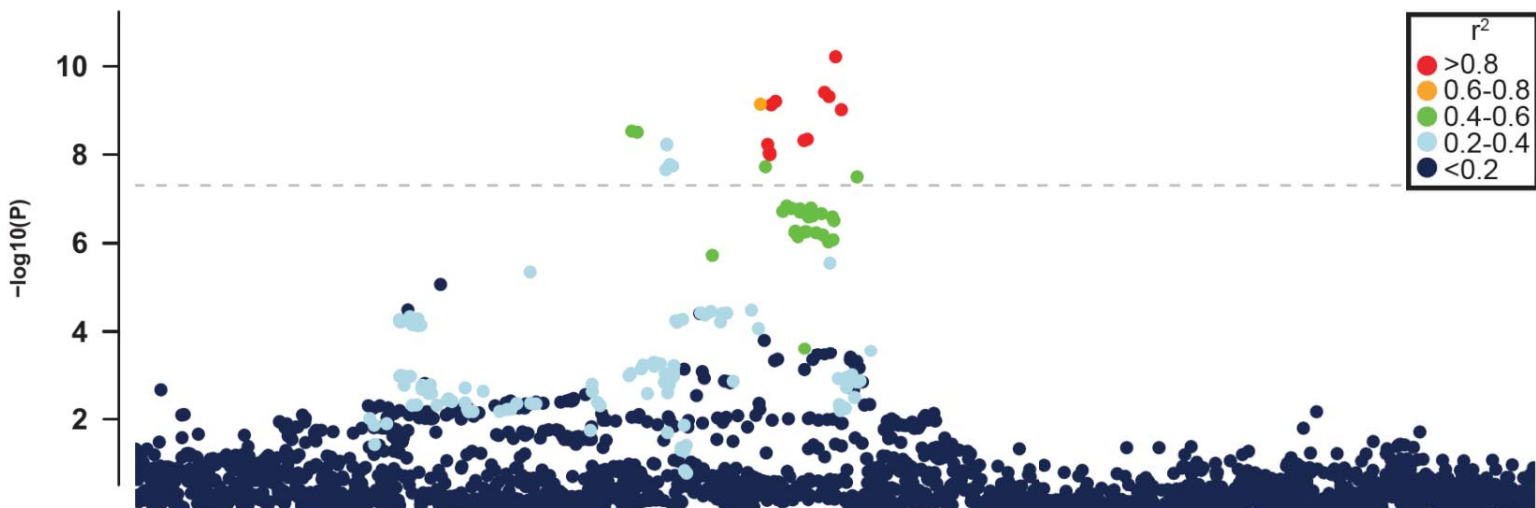
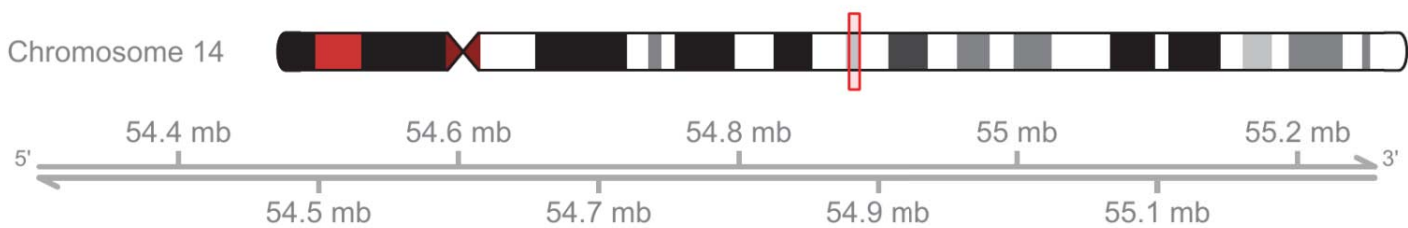


Corrected for total SA

Uncorrected

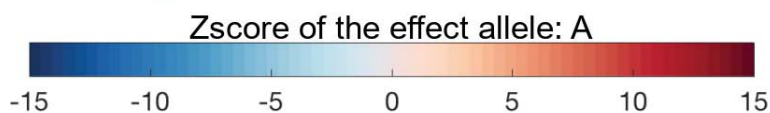
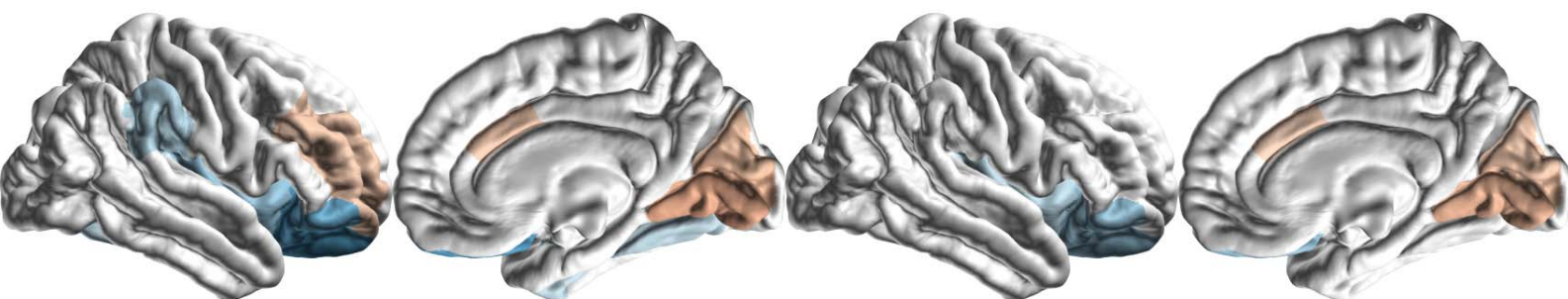


Pars Orbitalis SA: rs7147119

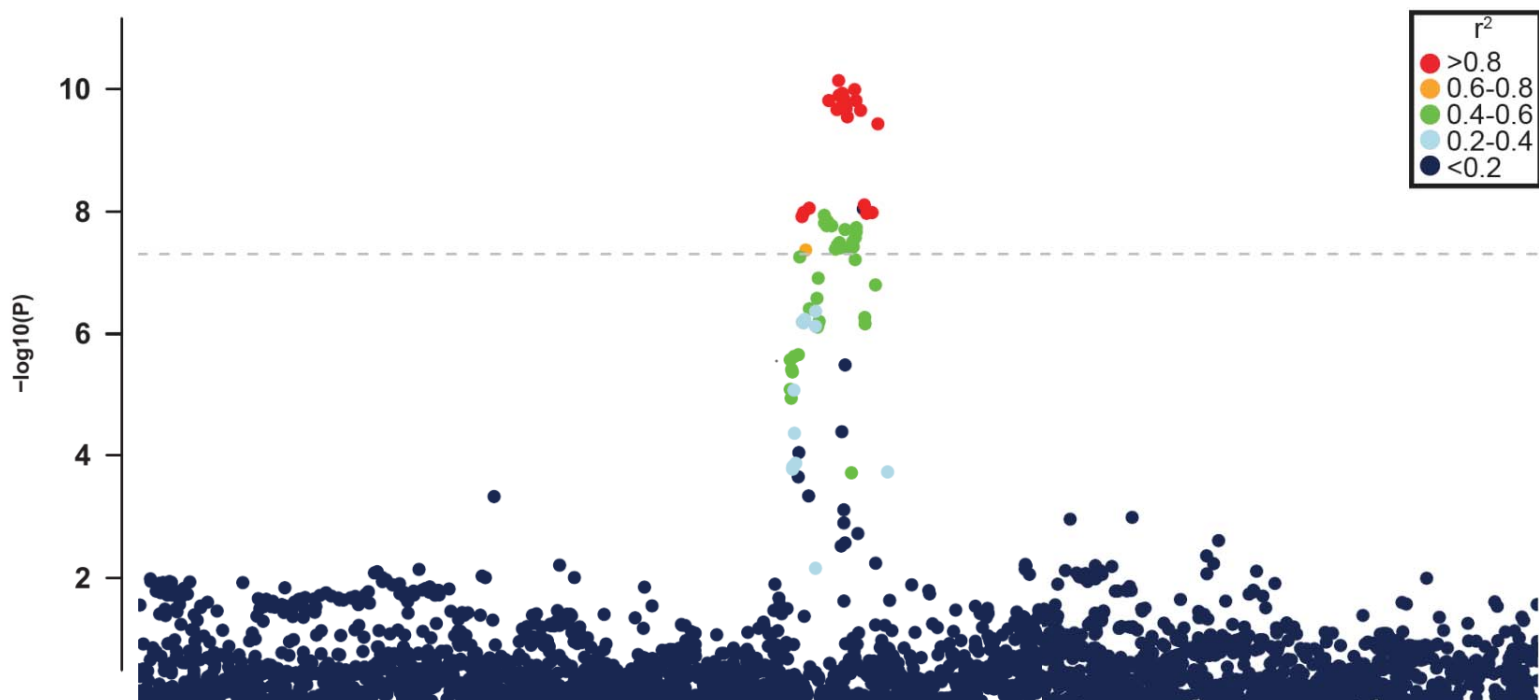
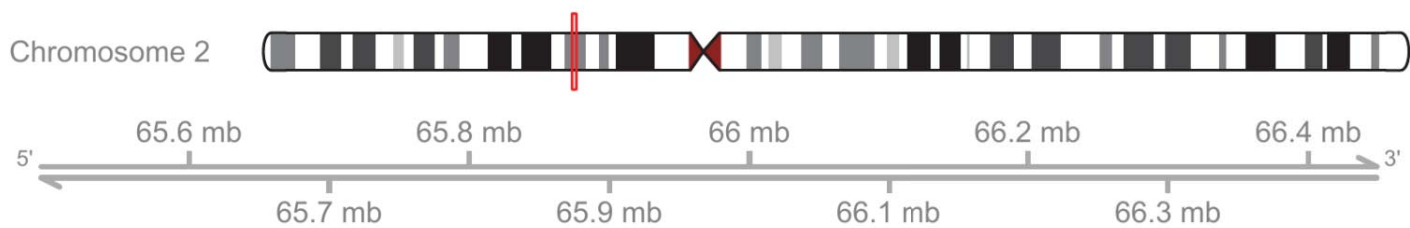


Corrected for total SA

Uncorrected



Pars Triangularis SA: rs2287283



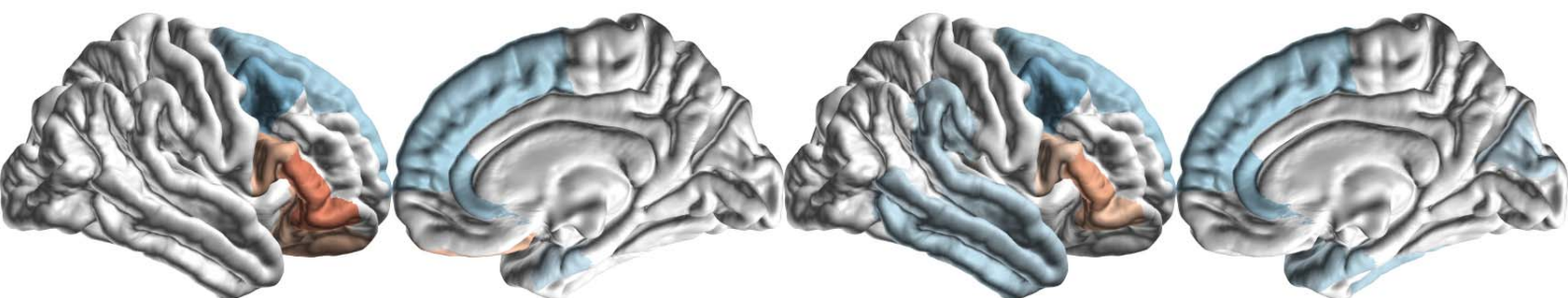
ACTR2 ← SPRED2

GZ>CP

CP>GZ

Corrected for total SA

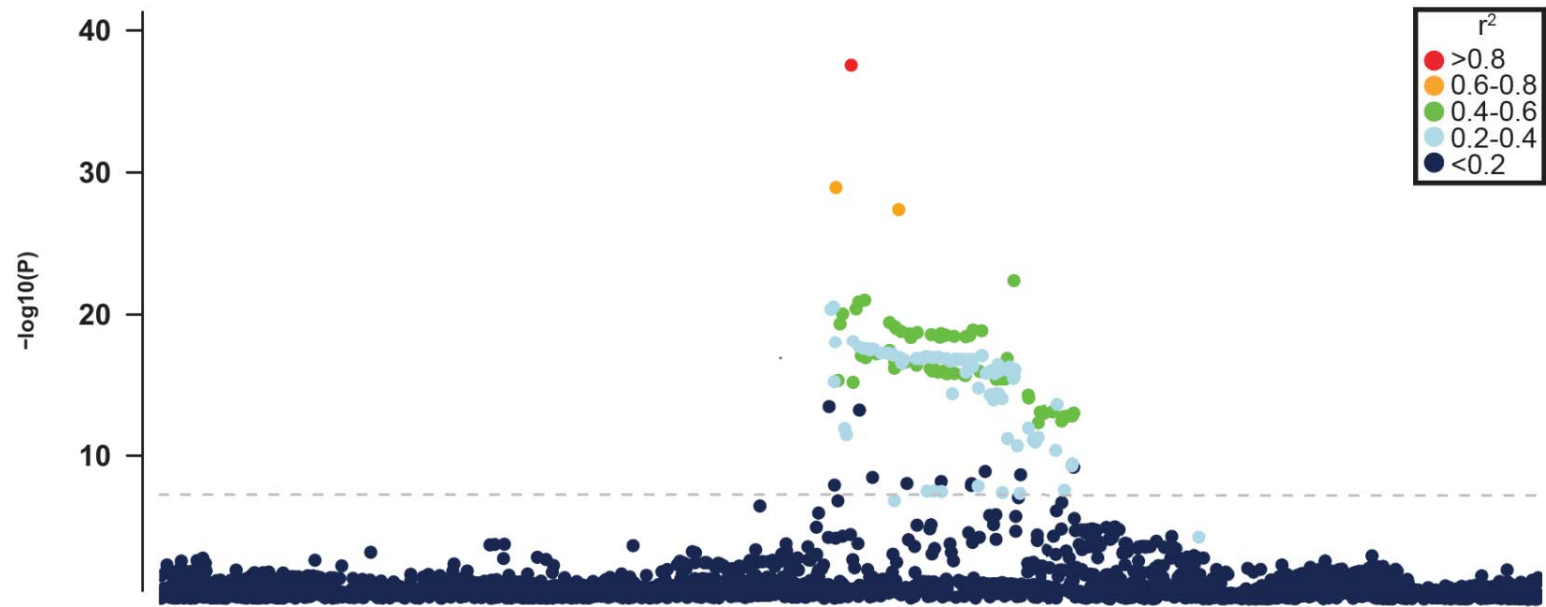
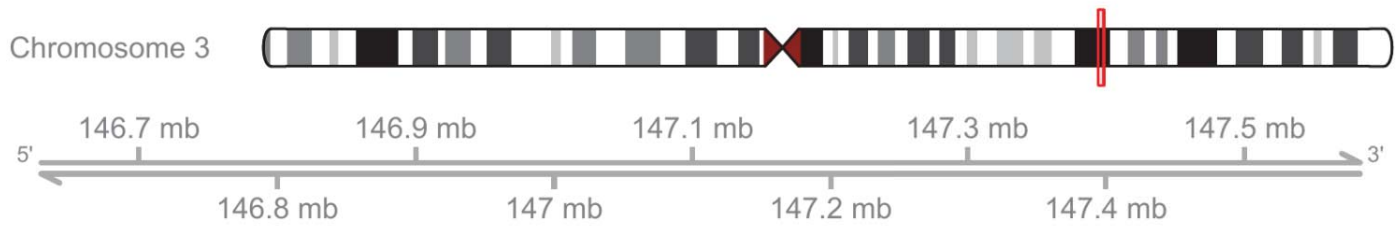
Uncorrected



Zscore of the effect allele: T



Pars Triangularis SA: rs2279829



ZIC4

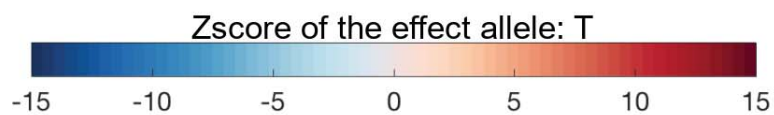
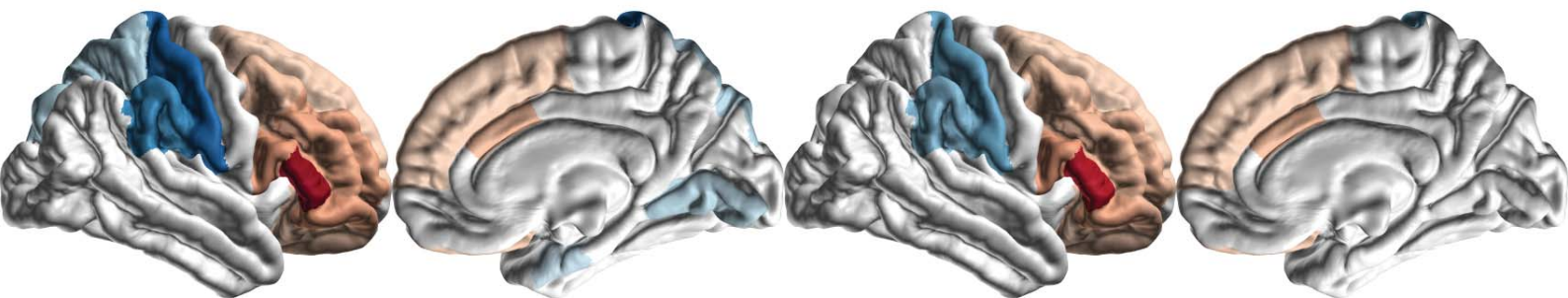
ZIC1

GZ>CP

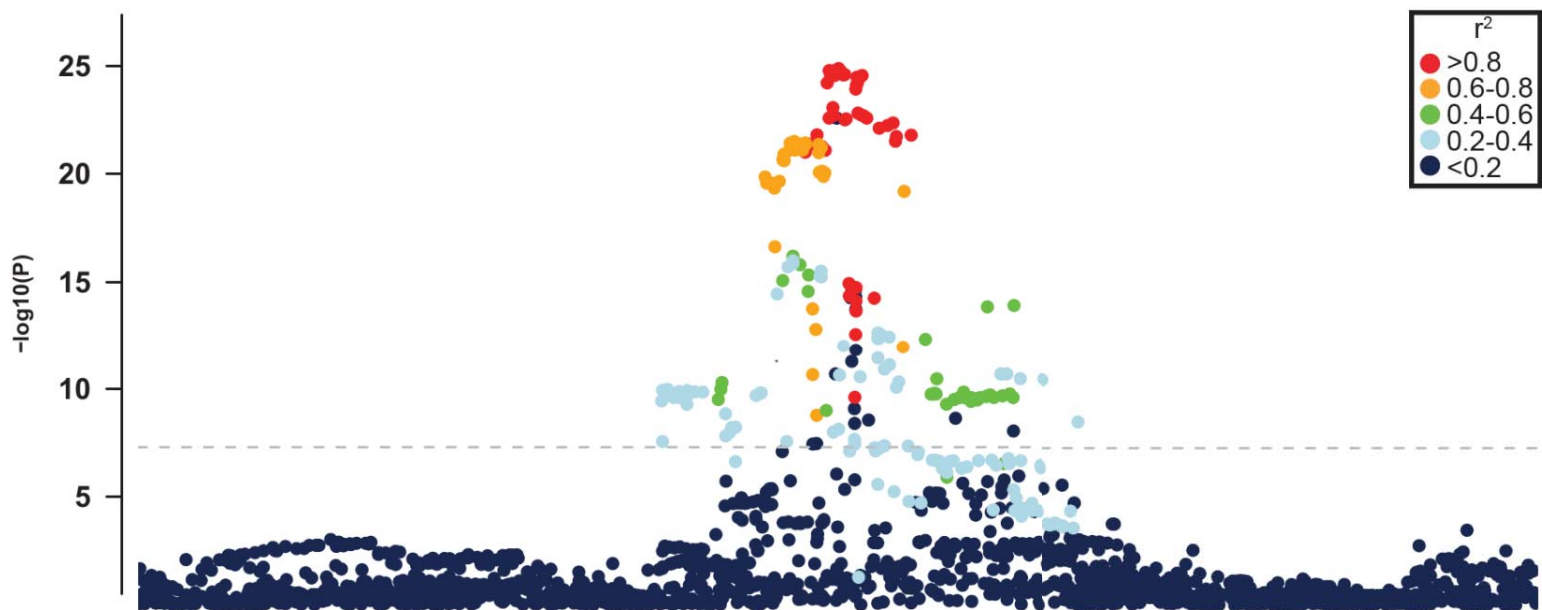
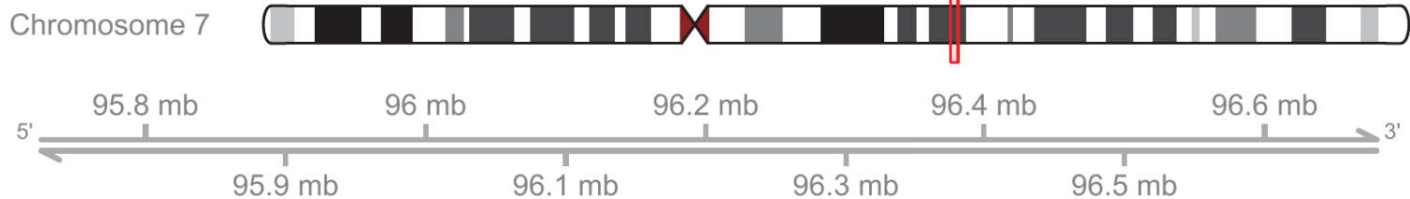
CP>GZ

Corrected for total SA

Uncorrected



Pars Triangularis SA: rs10278627



Genes: DYNC111, C7orf76, SHFM1, DLX5

Gene: SLC25A13, DLX5

GZ>CP

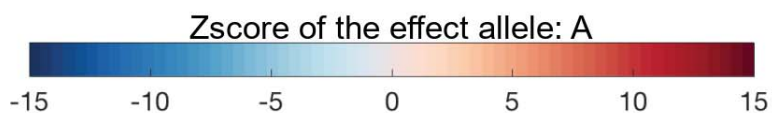
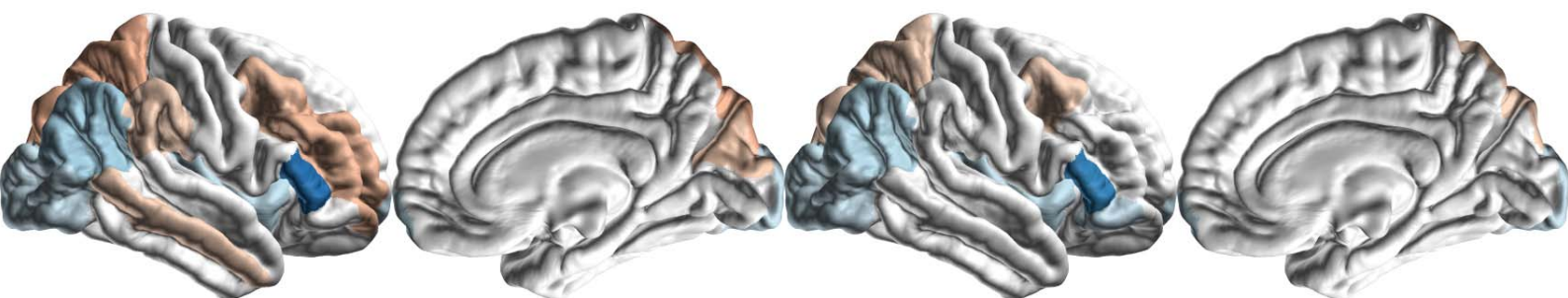


CP>GZ

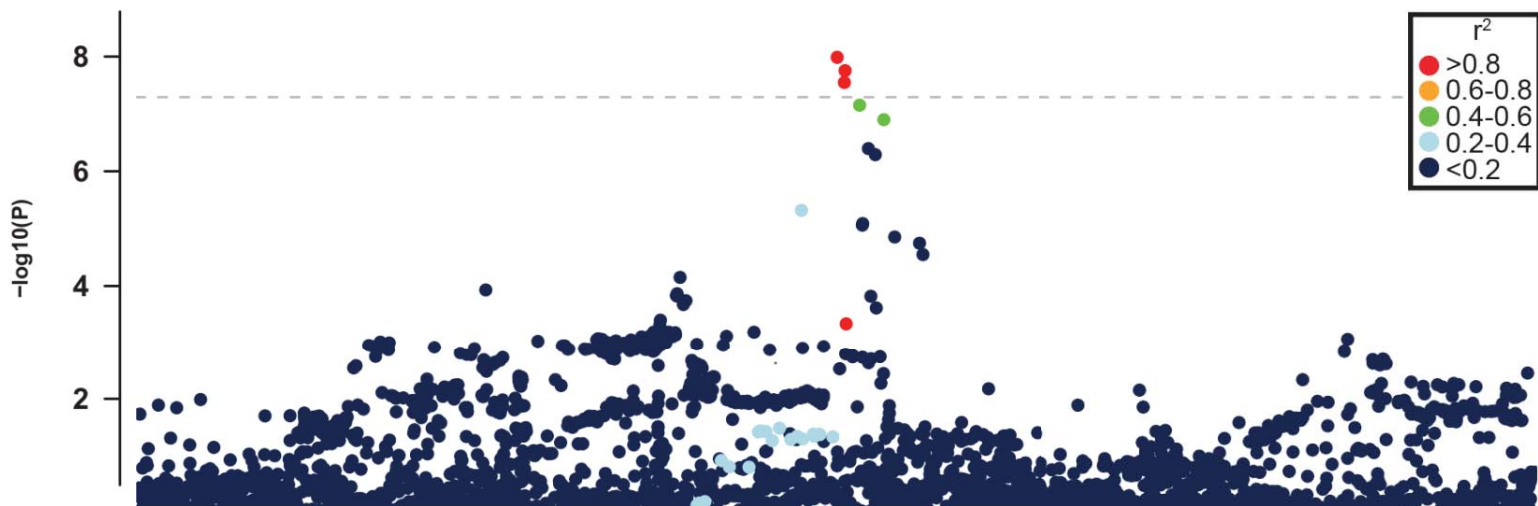
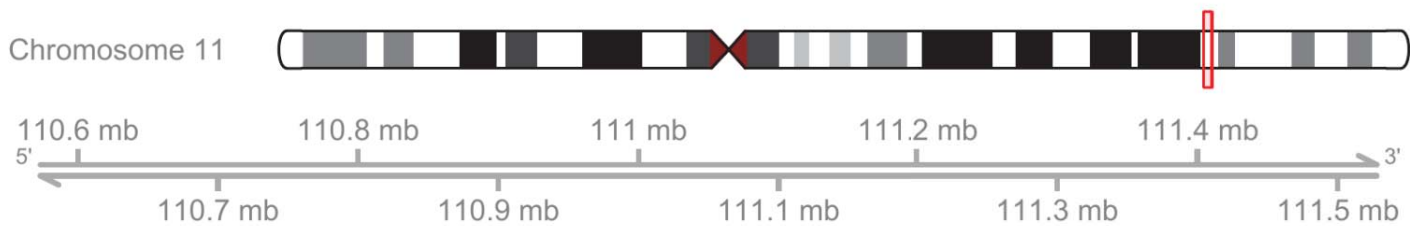


Corrected for total SA

Uncorrected



Pars Triangularis SA: rs59614433



ARHGAP20

C11orf53

C11orf88

COLCA2

LAYN

POU2AF1

→

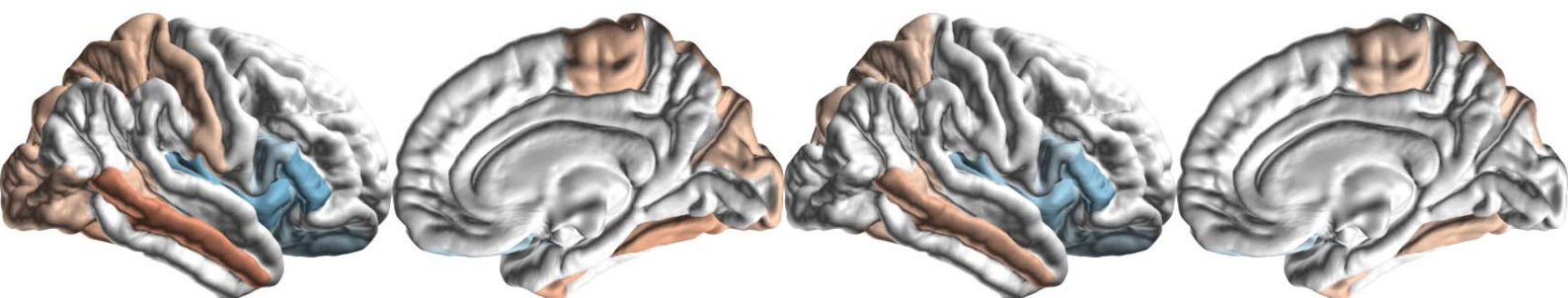
← BTG4

CP>GZ

GZ>CP

Corrected for total SA

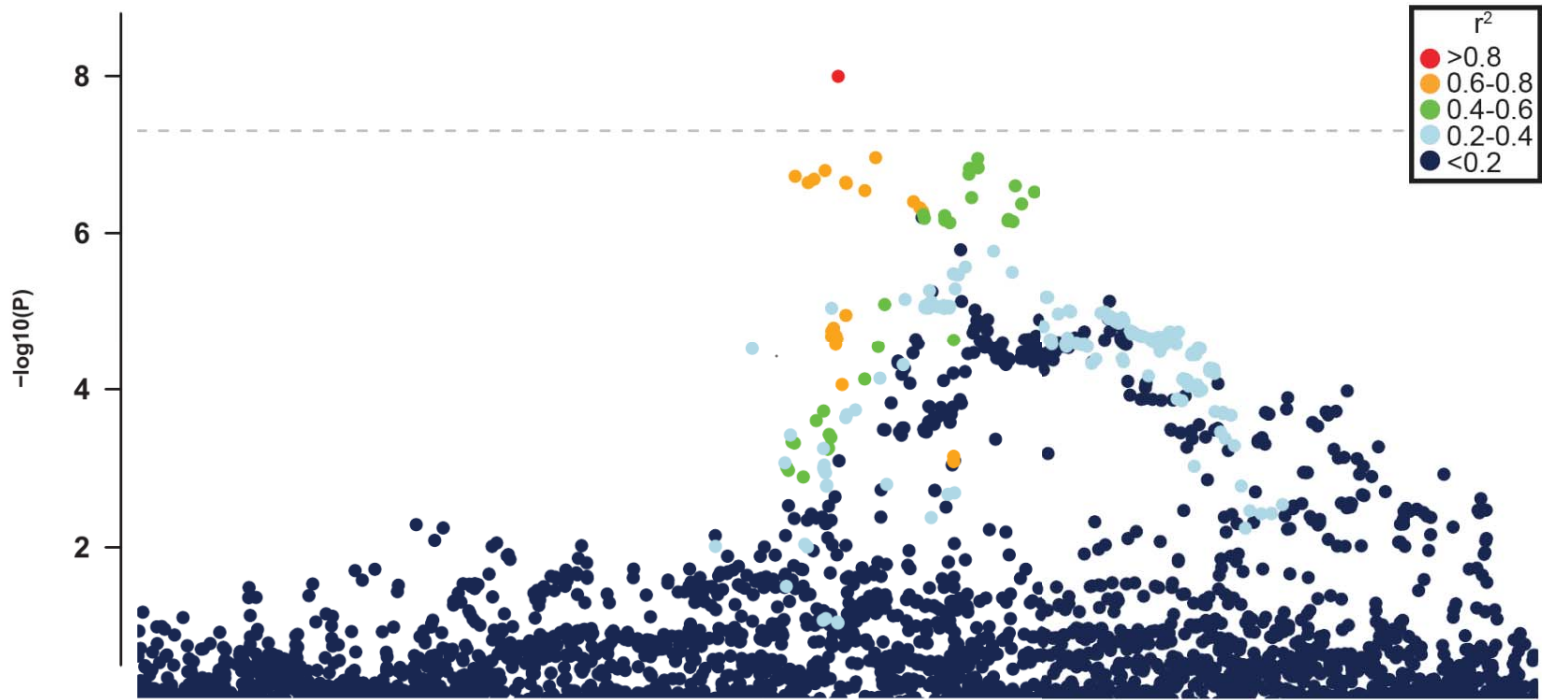
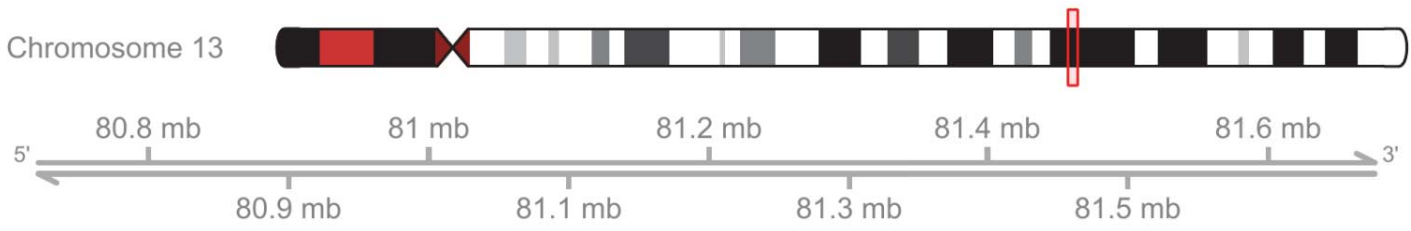
Uncorrected



Zscore of the effect allele: T



Pars Triangularis SA: rs7996803



SPRY2

GZ>CP

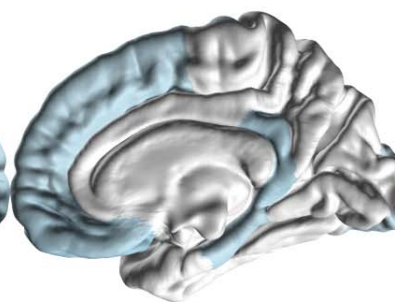
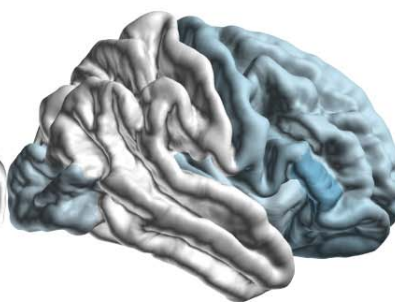
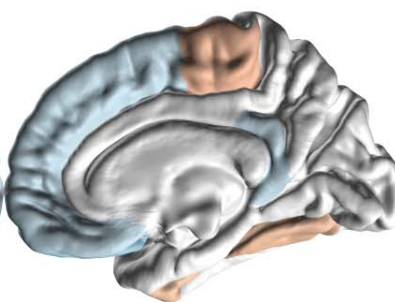
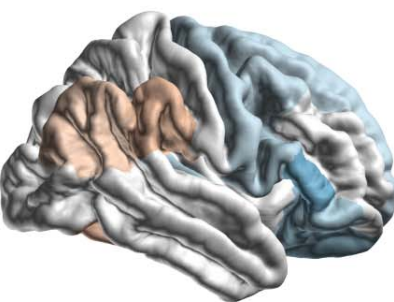


CP>GZ

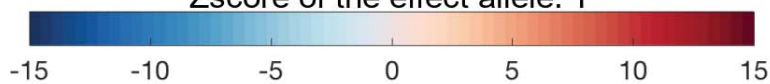


Corrected for total SA

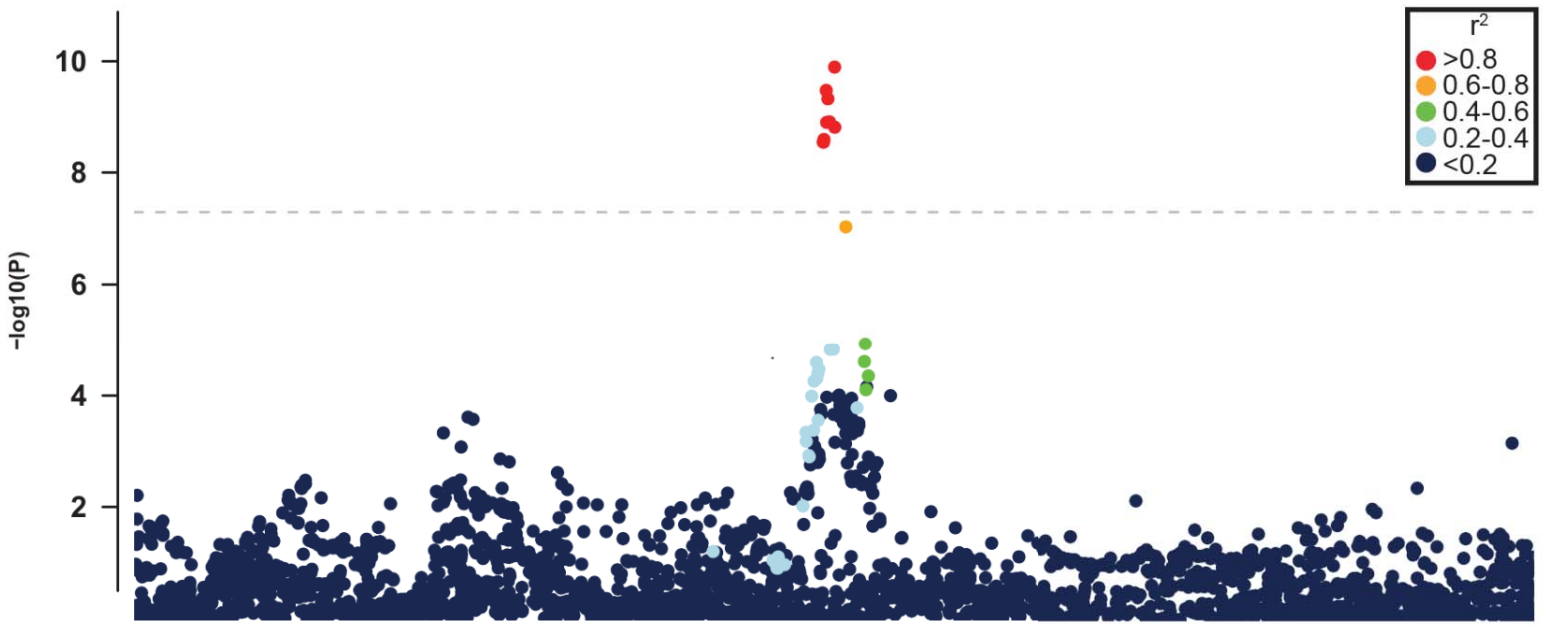
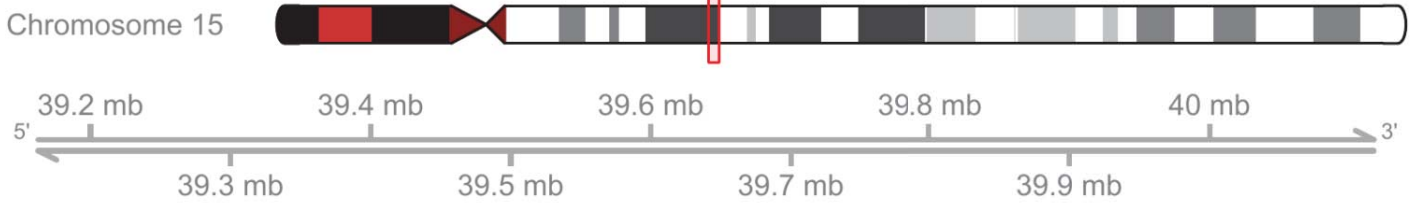
Uncorrected



Zscore of the effect allele: T

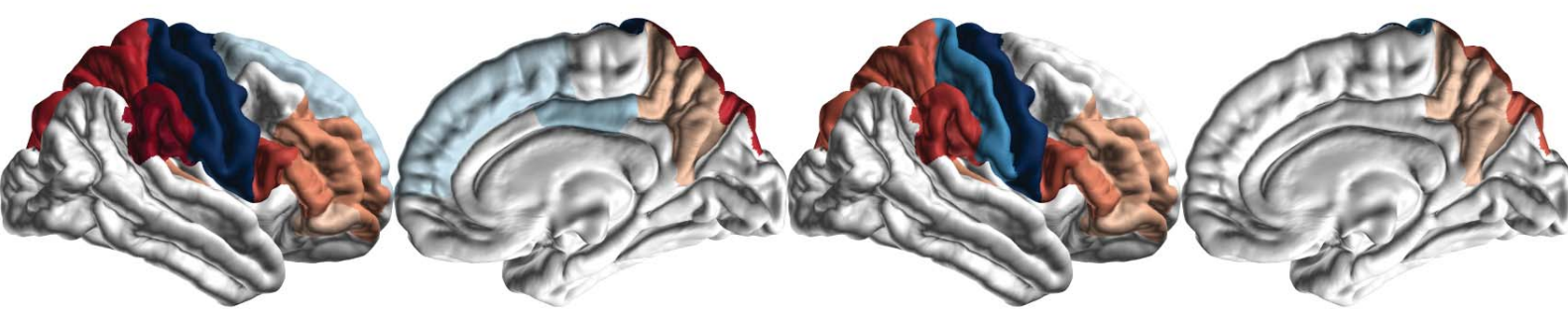


Pars Triangularis SA: rs4924345

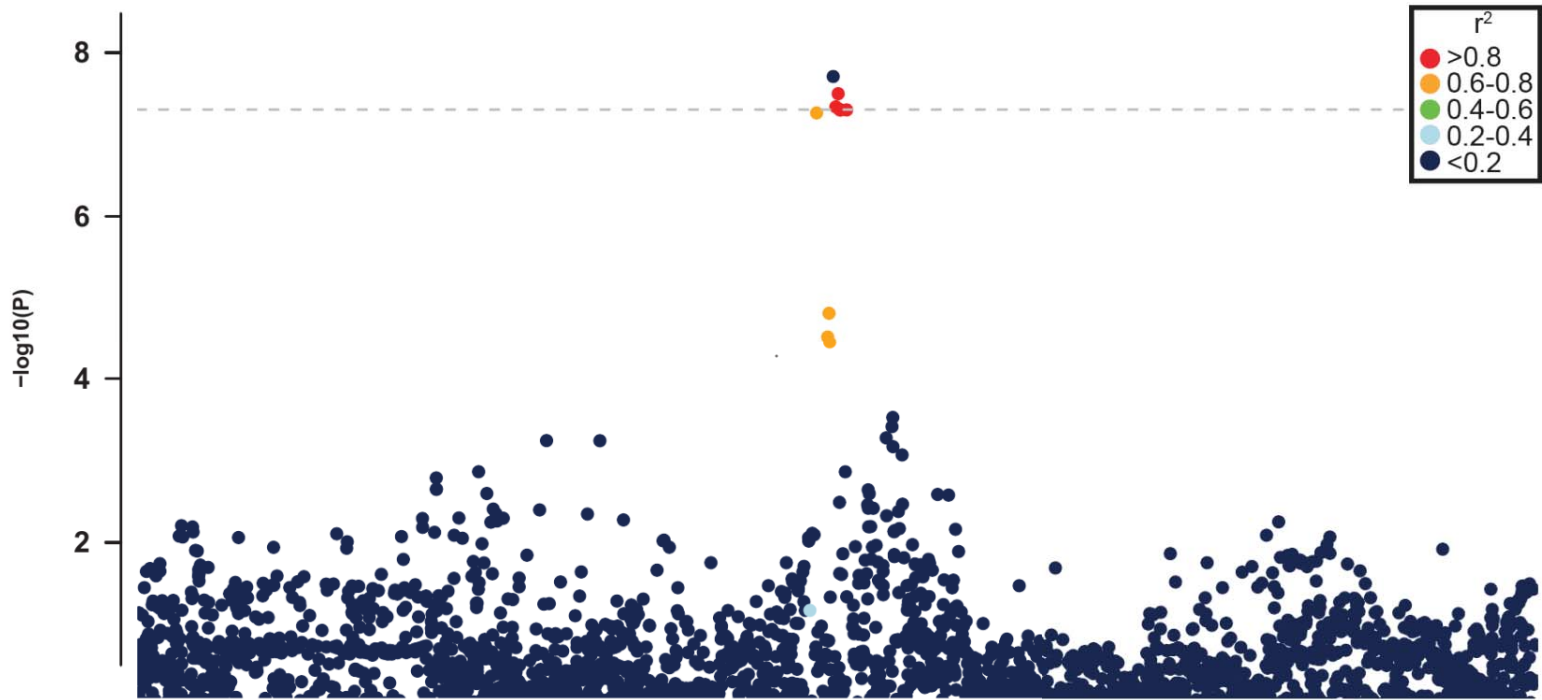
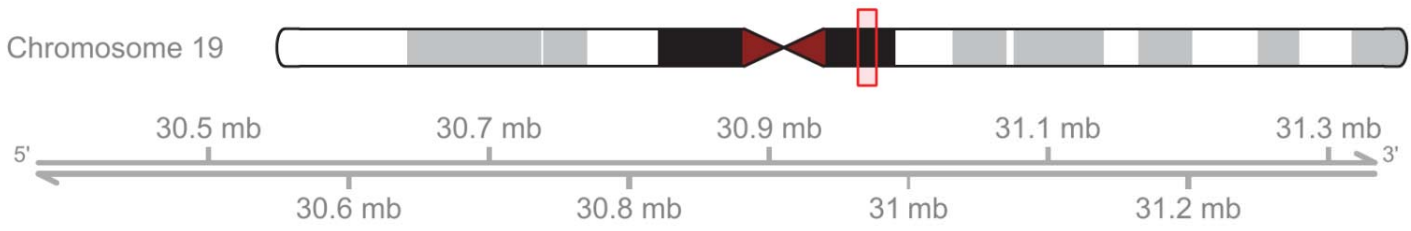


Corrected for total SA

Uncorrected



Pars Triangularis SA: rs1125867



URI1

ZNF536

GZ>CP

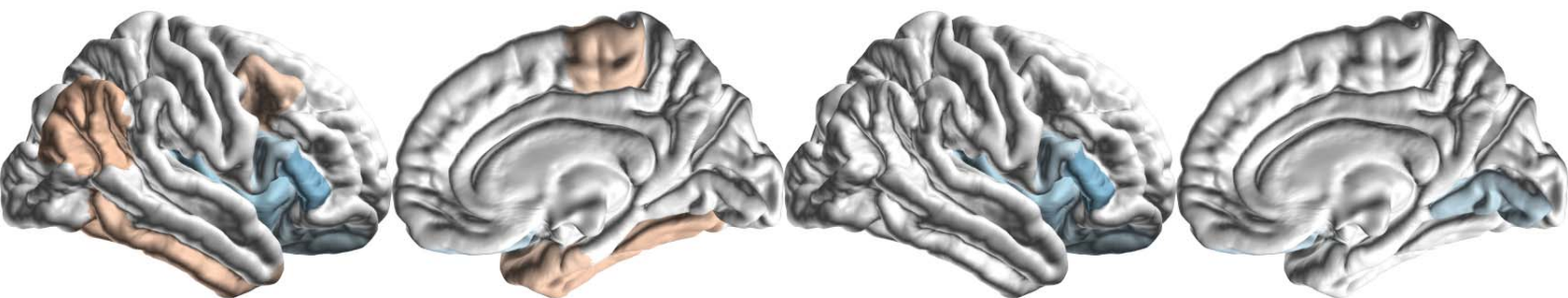


CP>GZ



Corrected for total SA

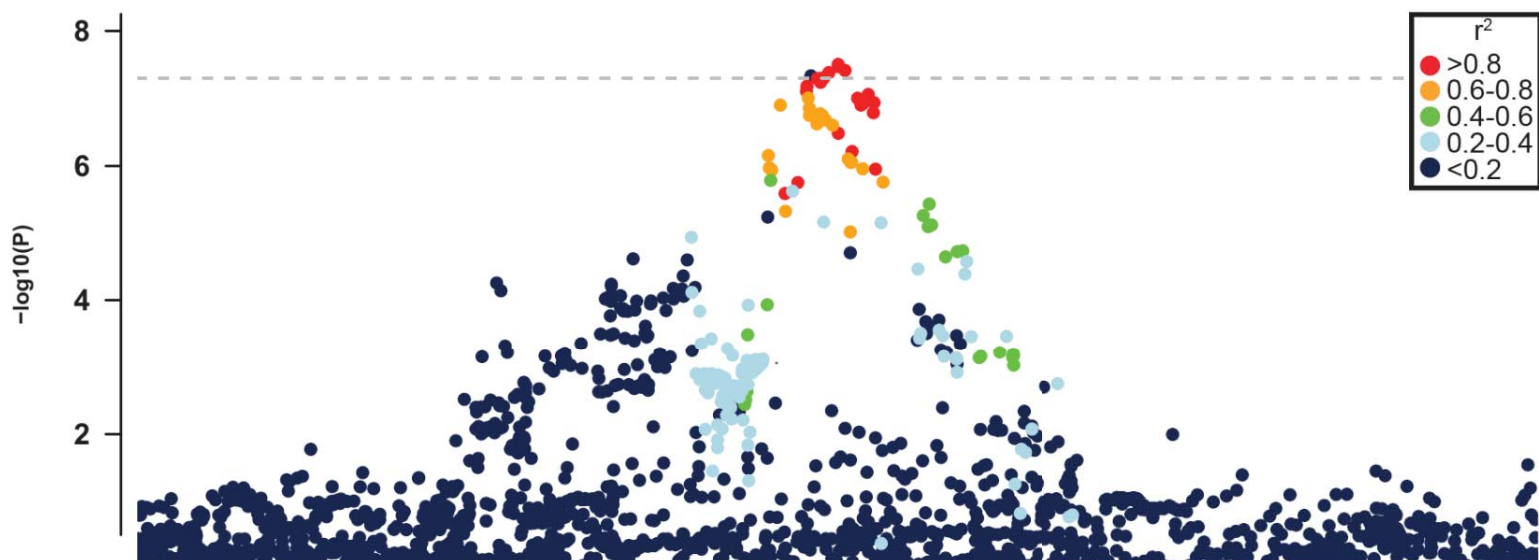
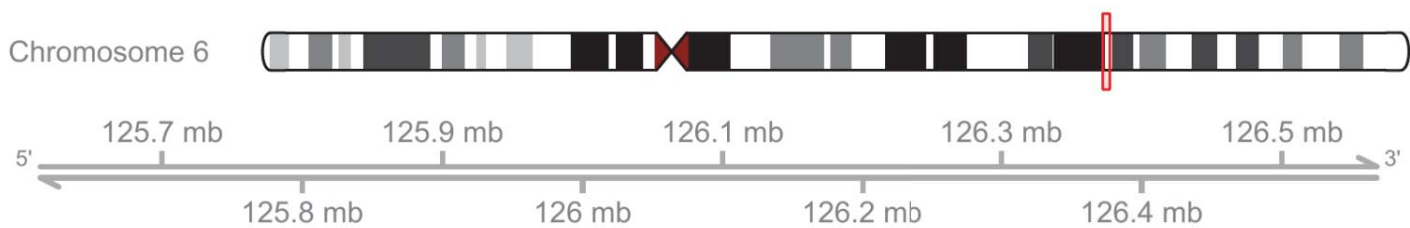
Uncorrected



Zscore of the effect allele: A



Pars Opercularis SA: rs1159974



HDDC2

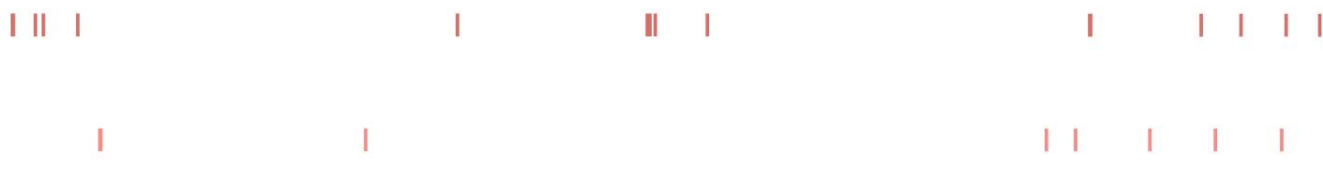
HEY2

HINT3

NCOA7

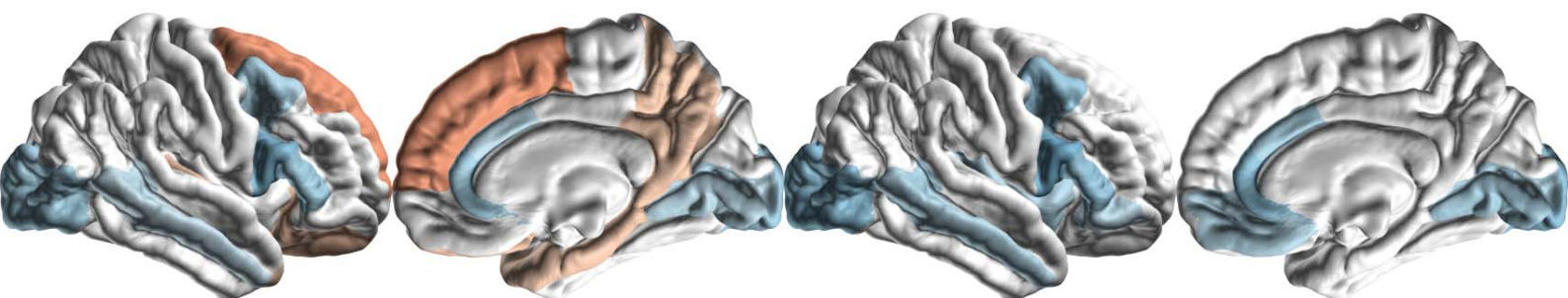
TRMT11

GZ>CP
CP>GZ



Corrected for total SA

Uncorrected

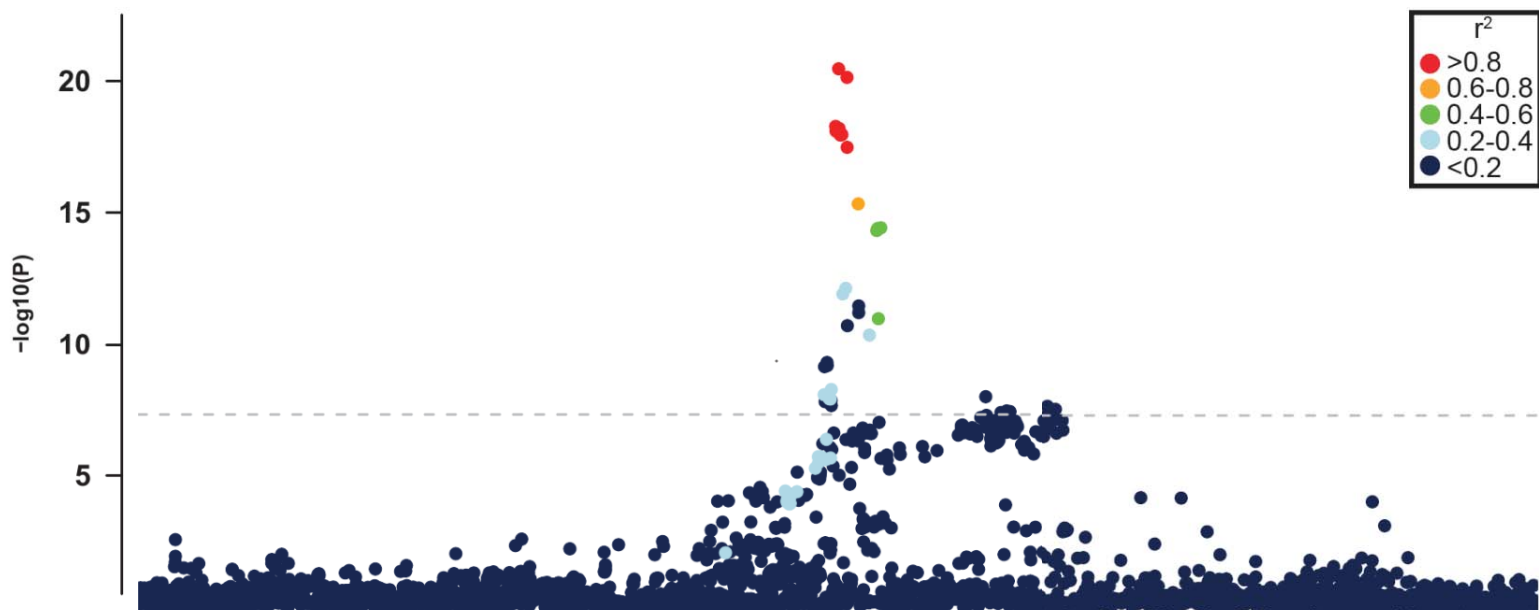
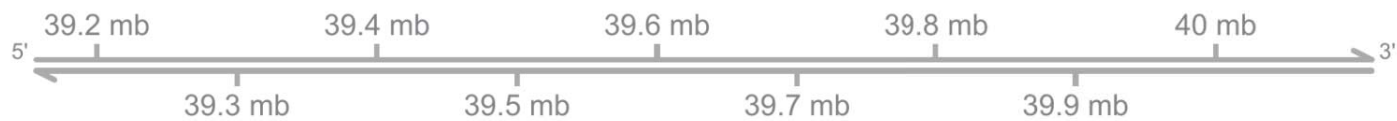


Zscore of the effect allele: T



Pars Opercularis SA: rs2033939

Chromosome 15

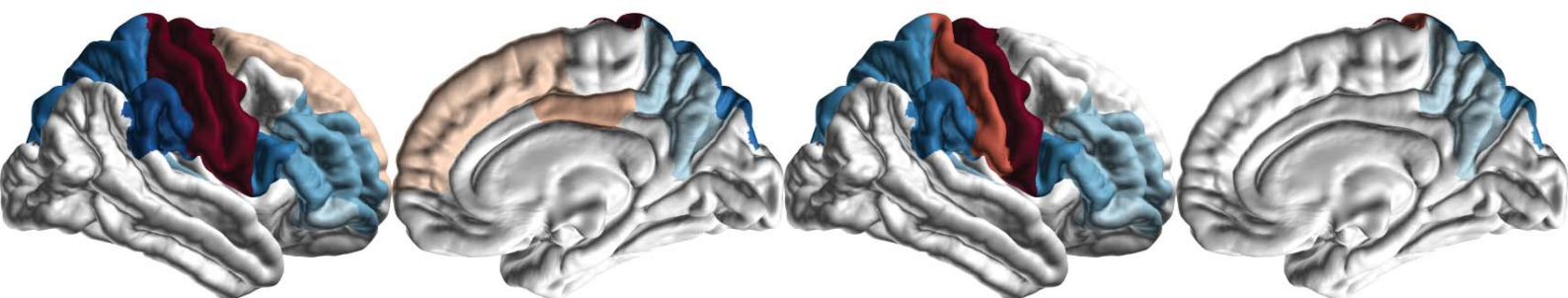


GZ>CP
CP>GZ



Corrected for total SA

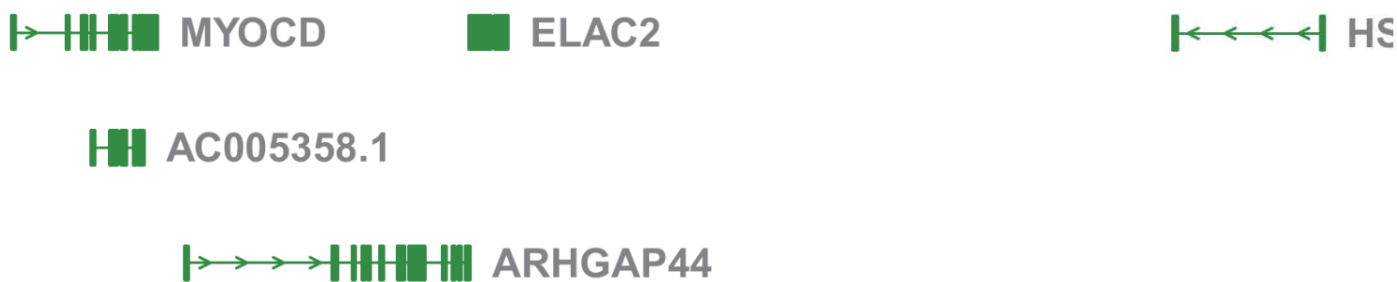
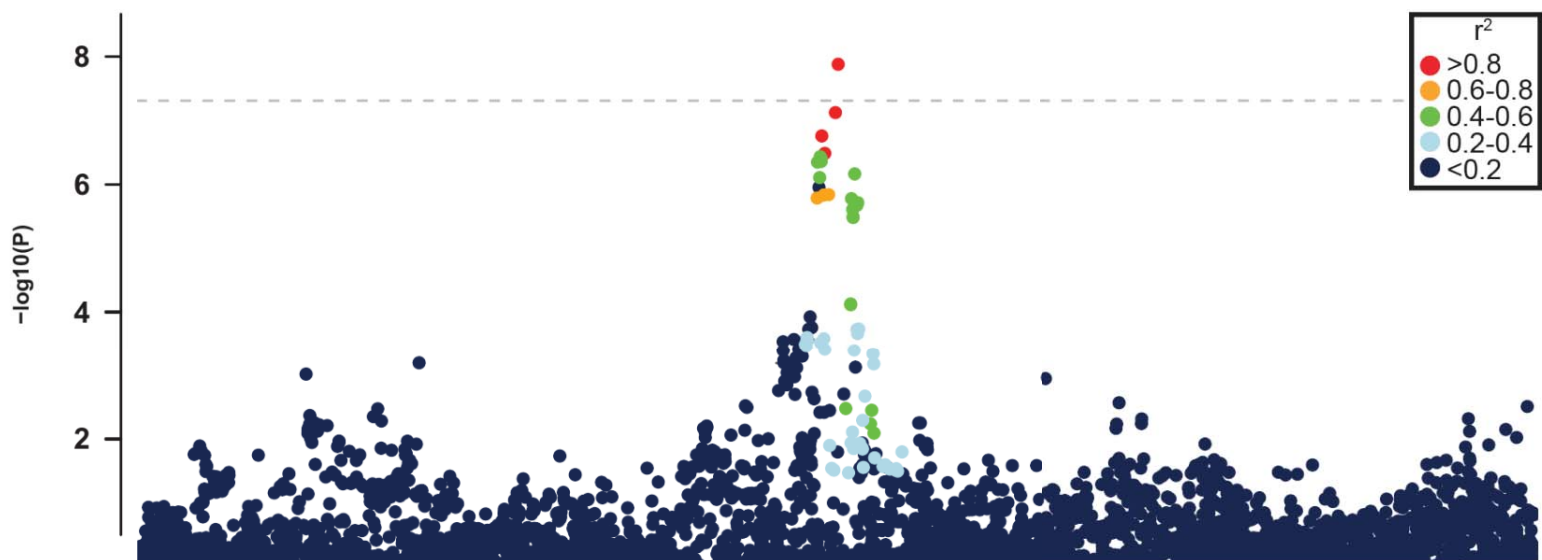
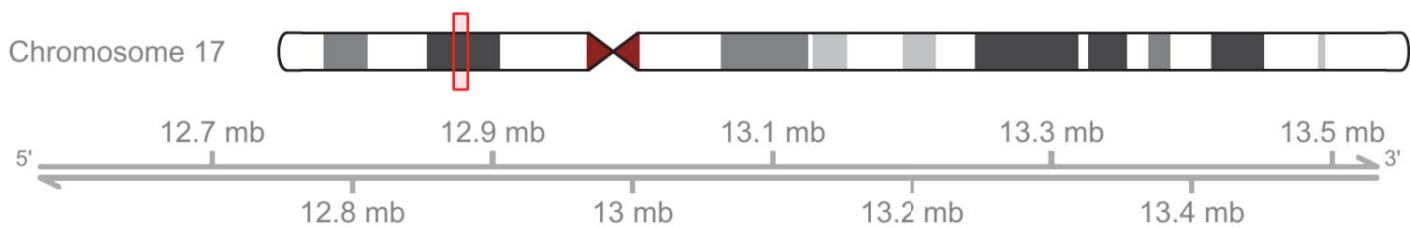
Uncorrected



Zscore of the effect allele: A



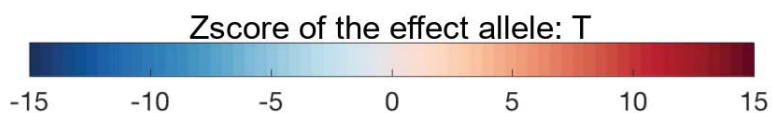
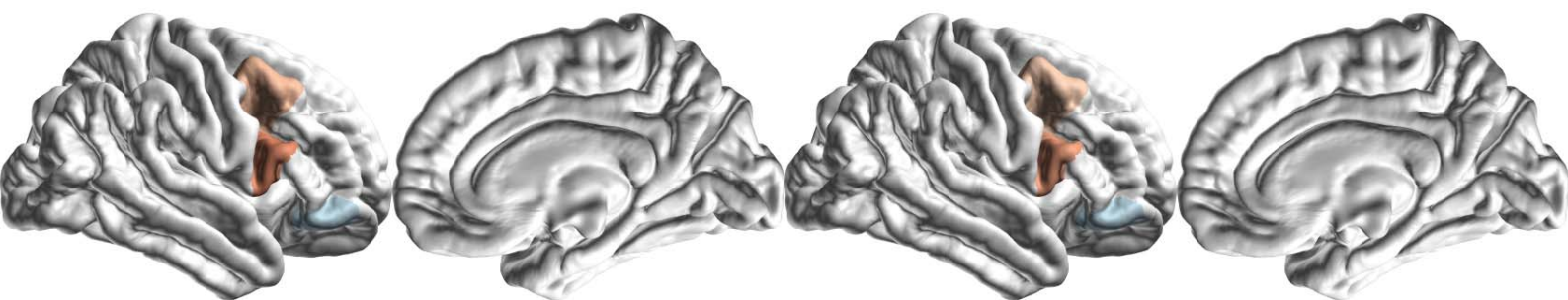
Pars Opercularis SA: rs12938190



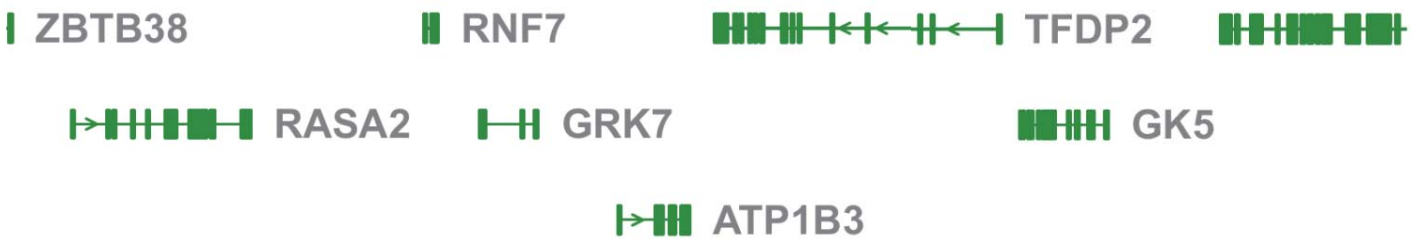
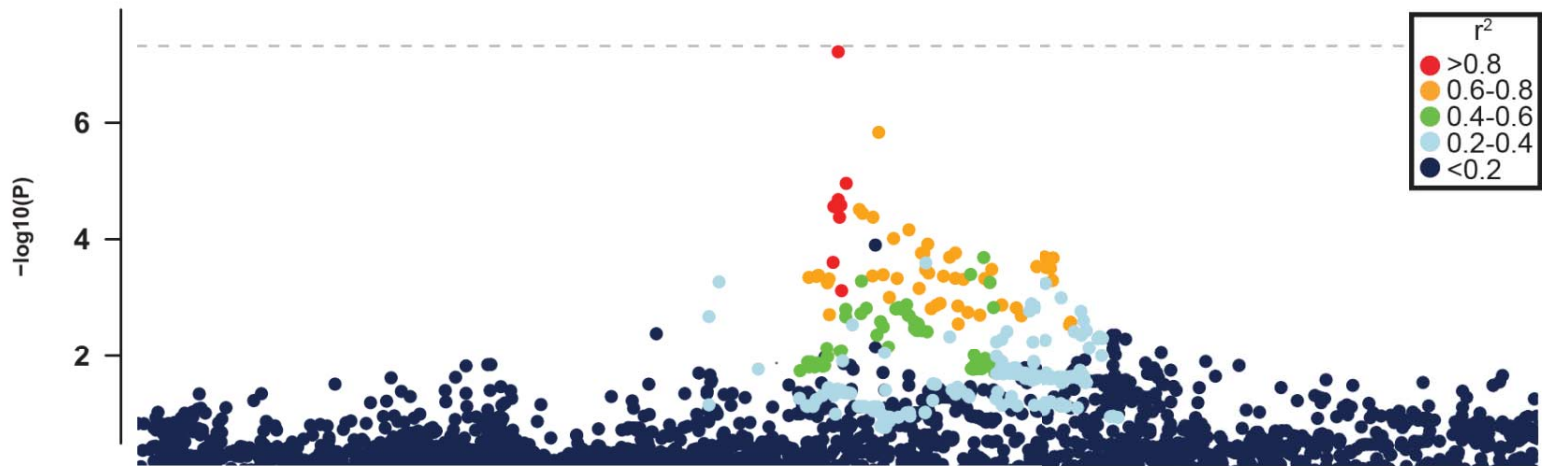
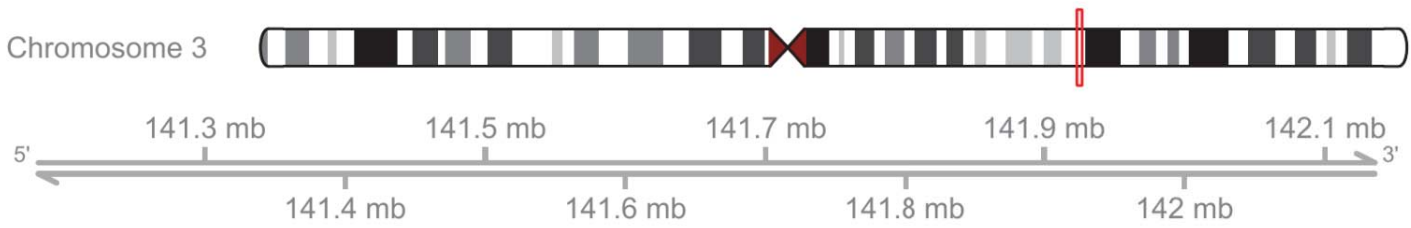
GZ>CP
CP>GZ

Corrected for total SA

Uncorrected

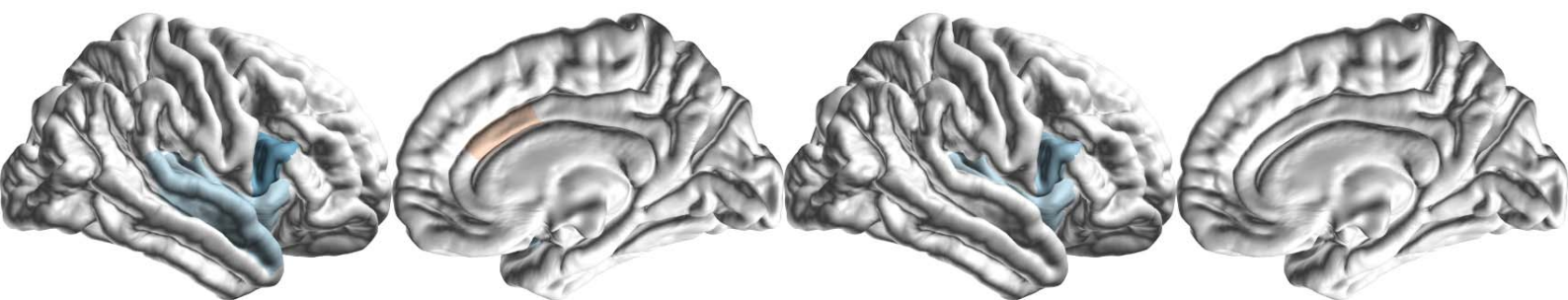


Pars Opercularis TH: rs7628850

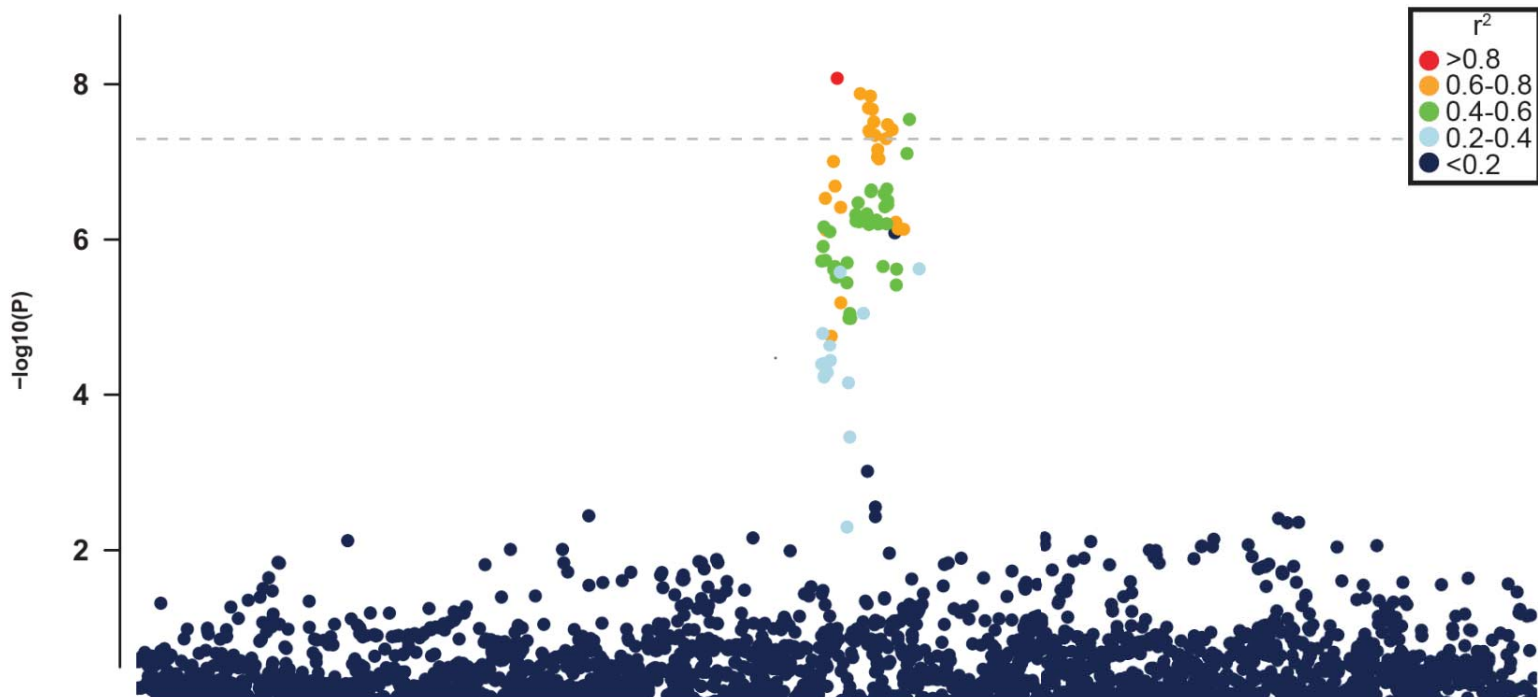
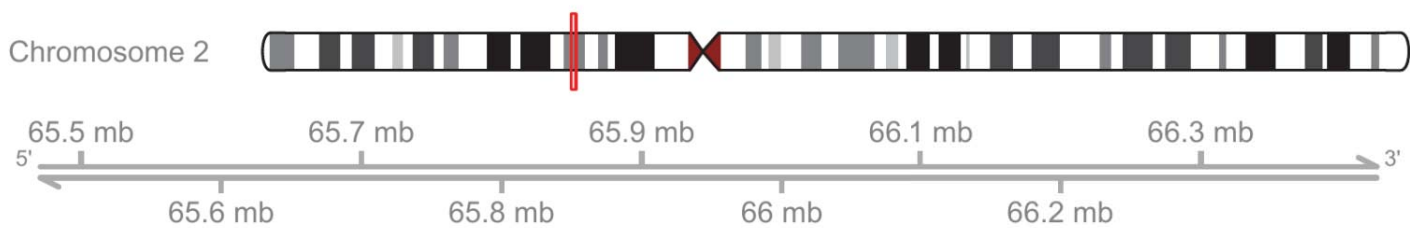


Corrected for average TH

Uncorrected



Caudal Middle Frontal SA: rs7580807



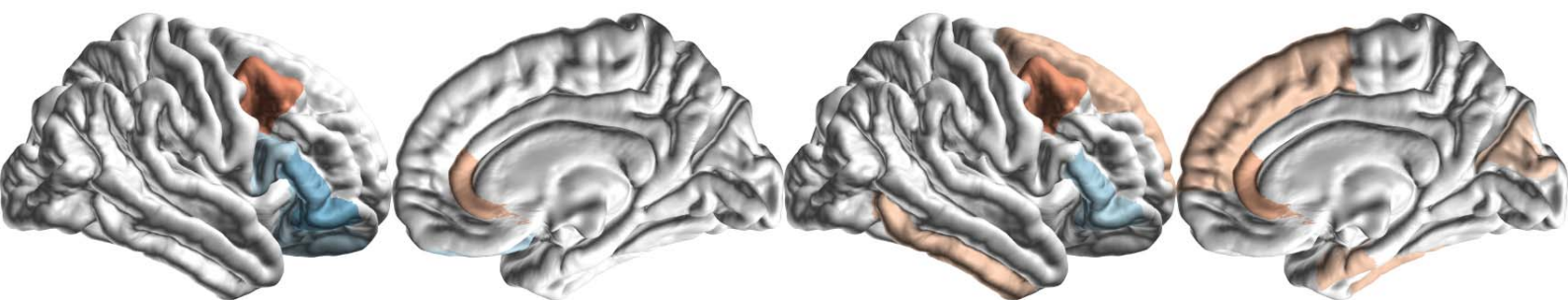
ACTR2 ← SPRED2

GZ>CP

CP>GZ

Corrected for total SA

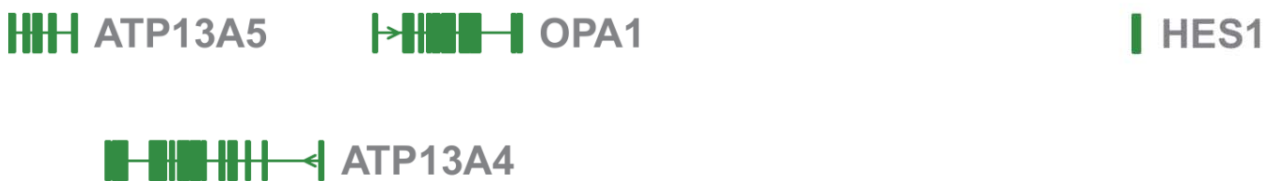
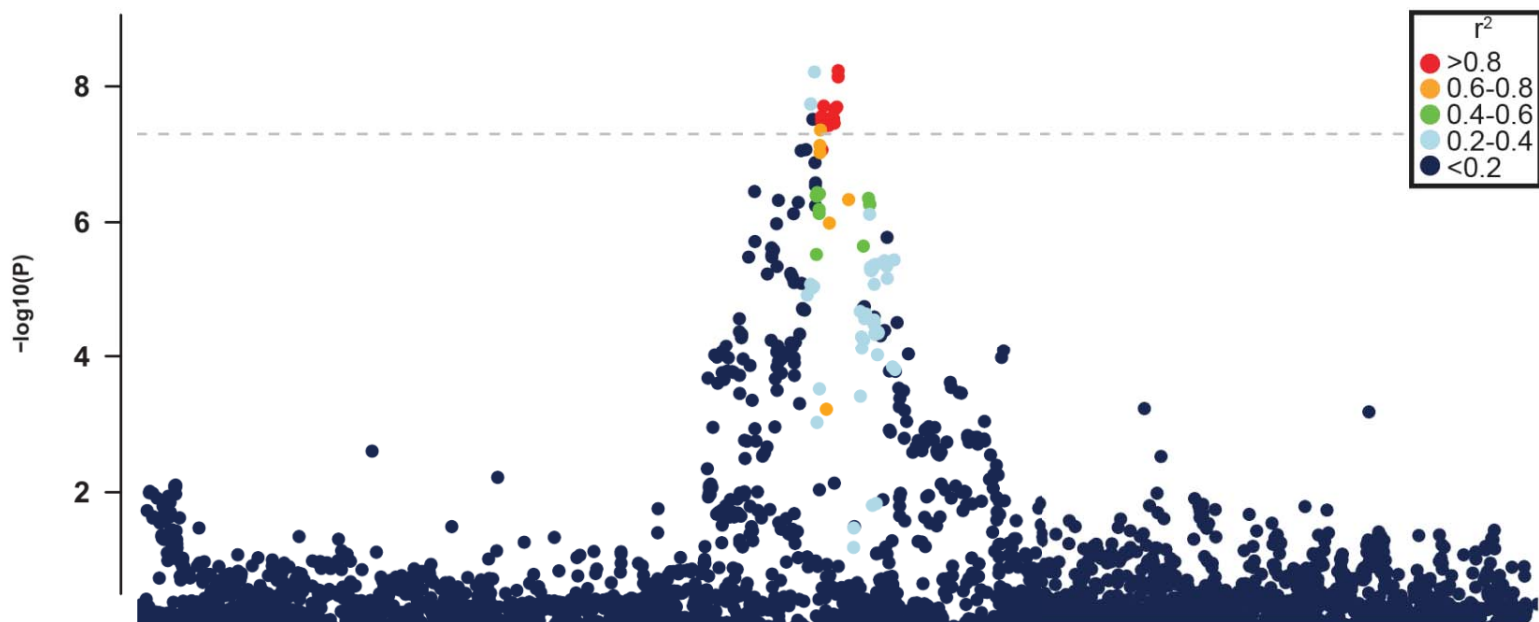
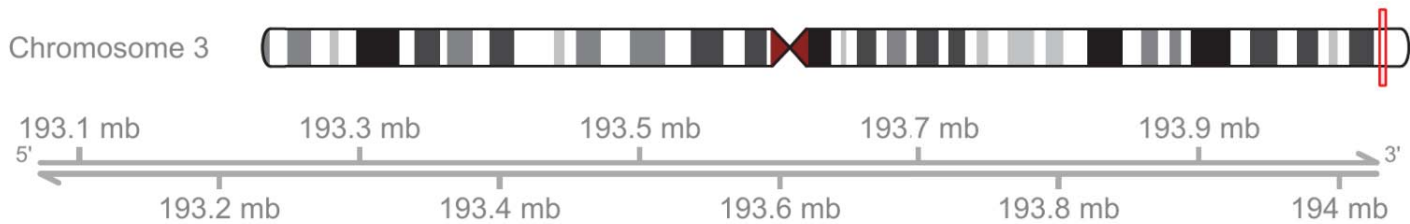
Uncorrected



Zscore of the effect allele: A

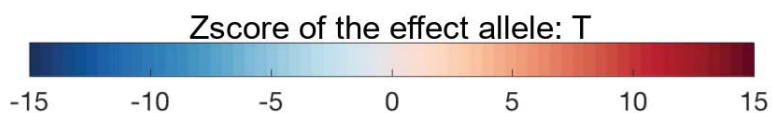
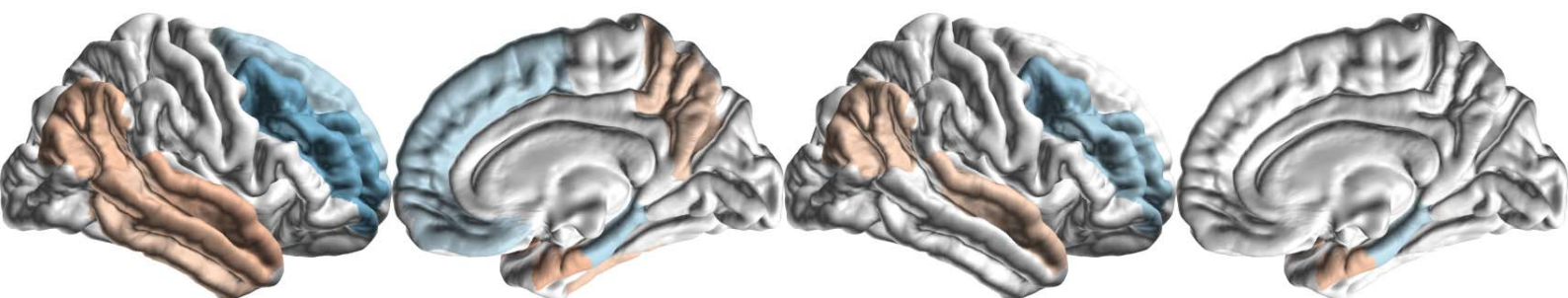


Caudal Middle Frontal SA: rs655045

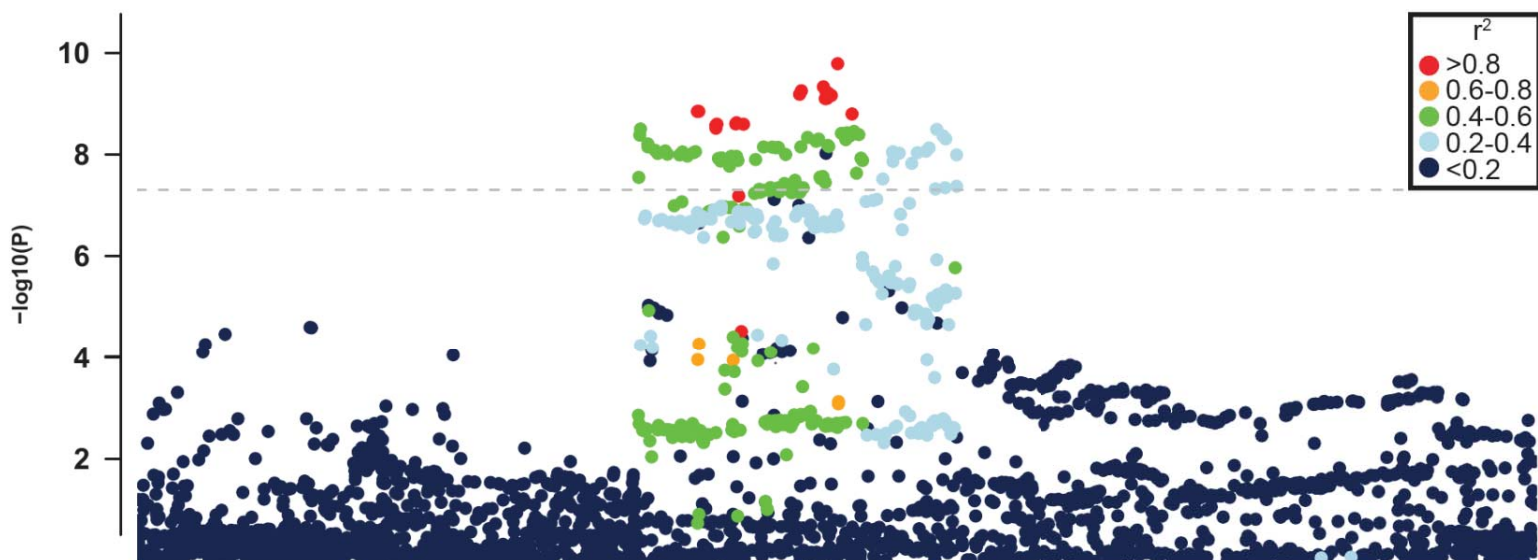
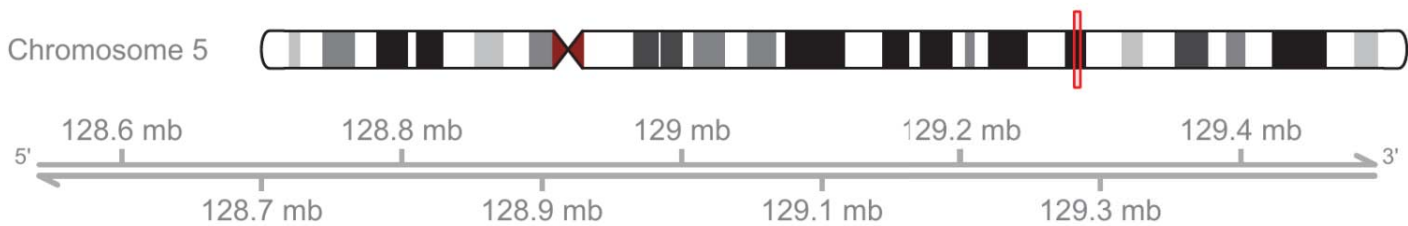


Corrected for total SA

Uncorrected



Caudal Middle Frontal SA: rs30641



ADAMTS19

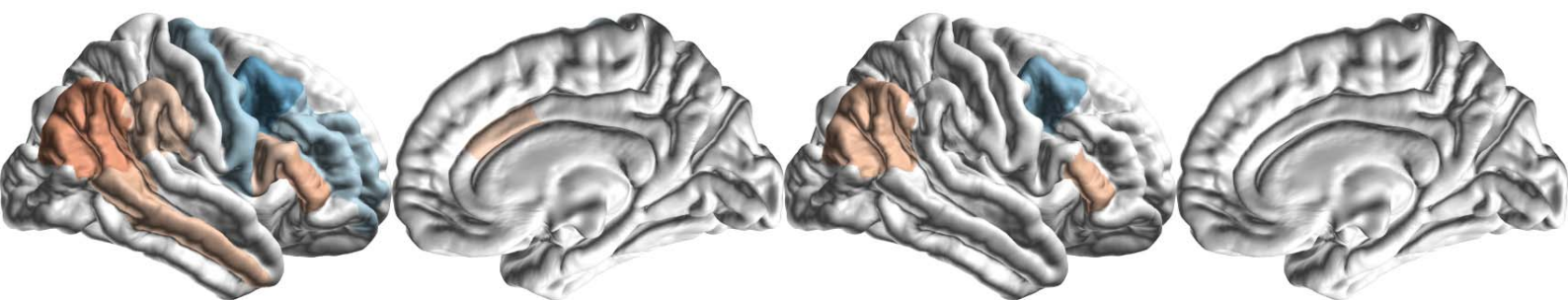
KIAA1024L

GZ>CP

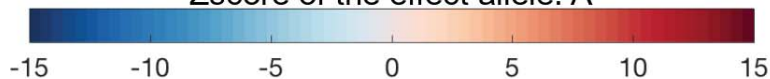
CP>GZ

Corrected for total SA

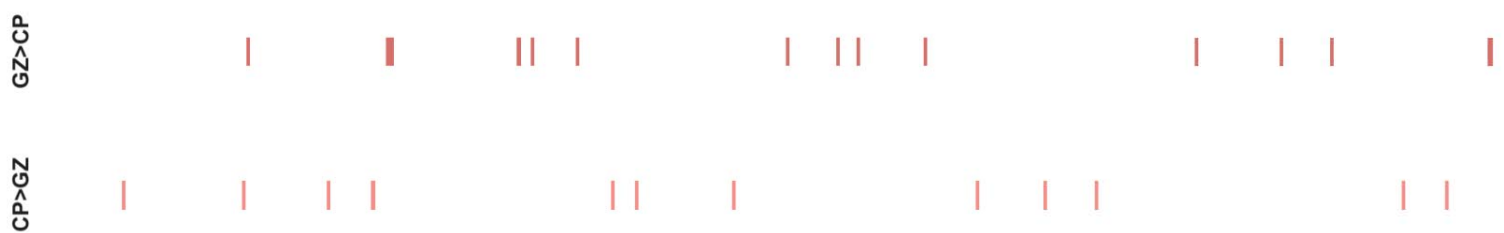
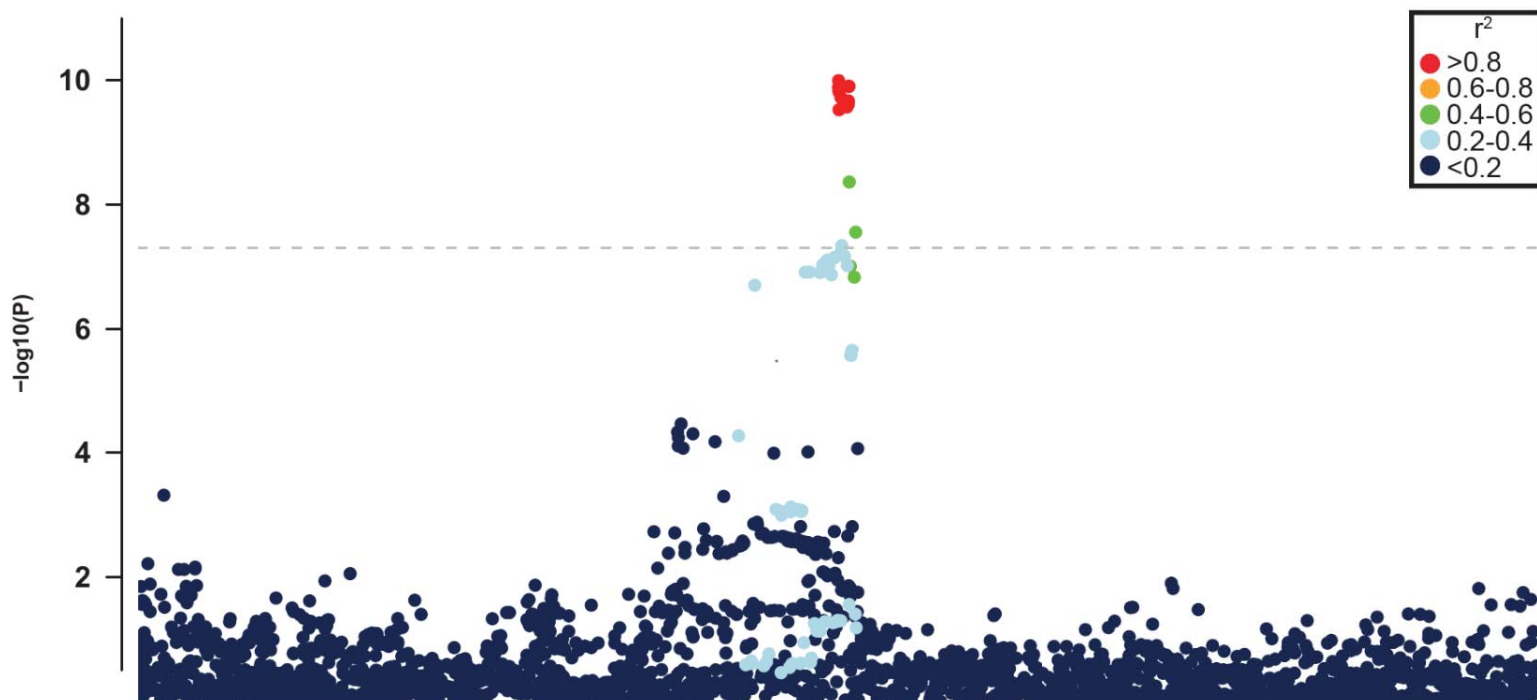
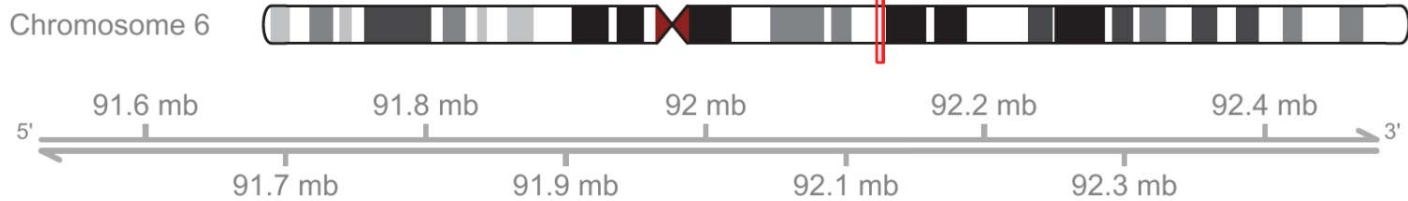
Uncorrected



Zscore of the effect allele: A

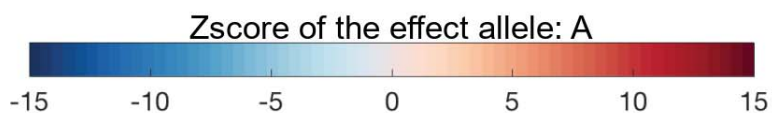
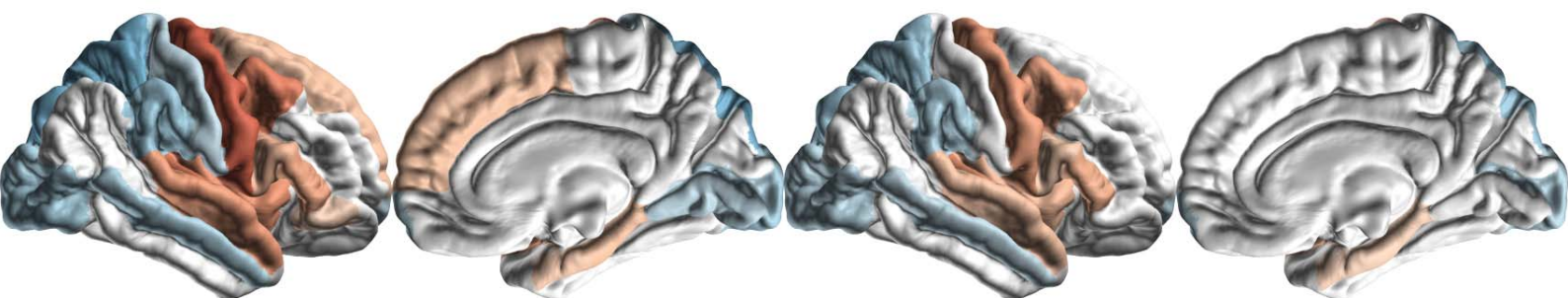


Caudal Middle Frontal SA: rs9345125

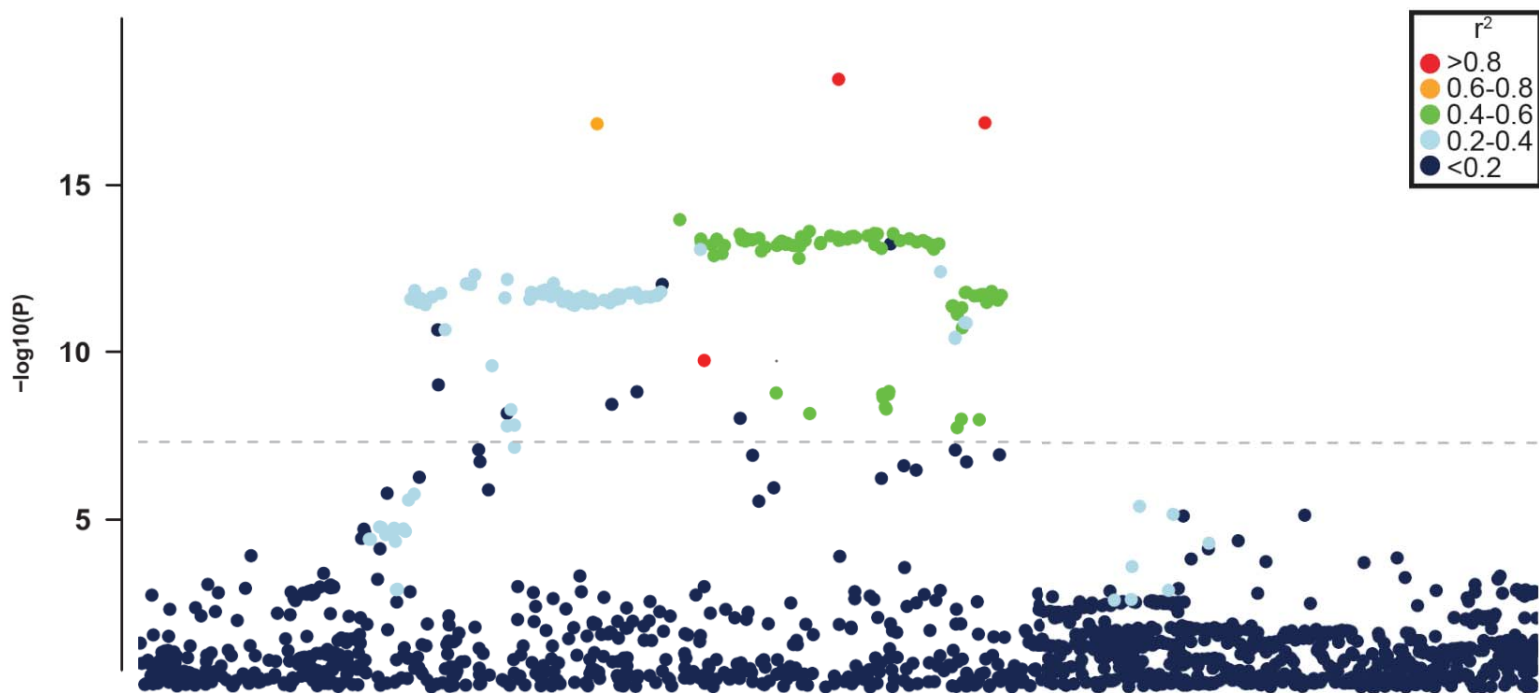
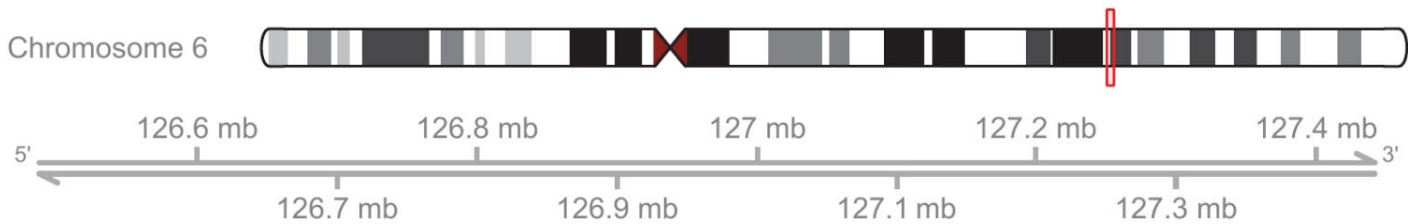


Corrected for total SA

Uncorrected



Caudal Middle Frontal SA: rs4273712



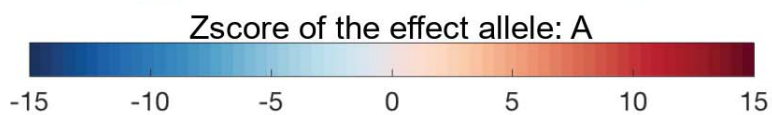
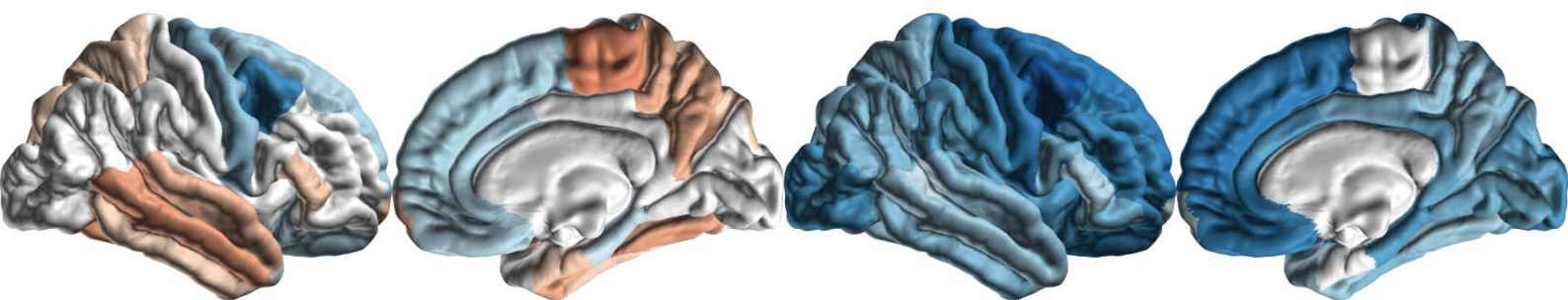
CENPW

GZ>CP

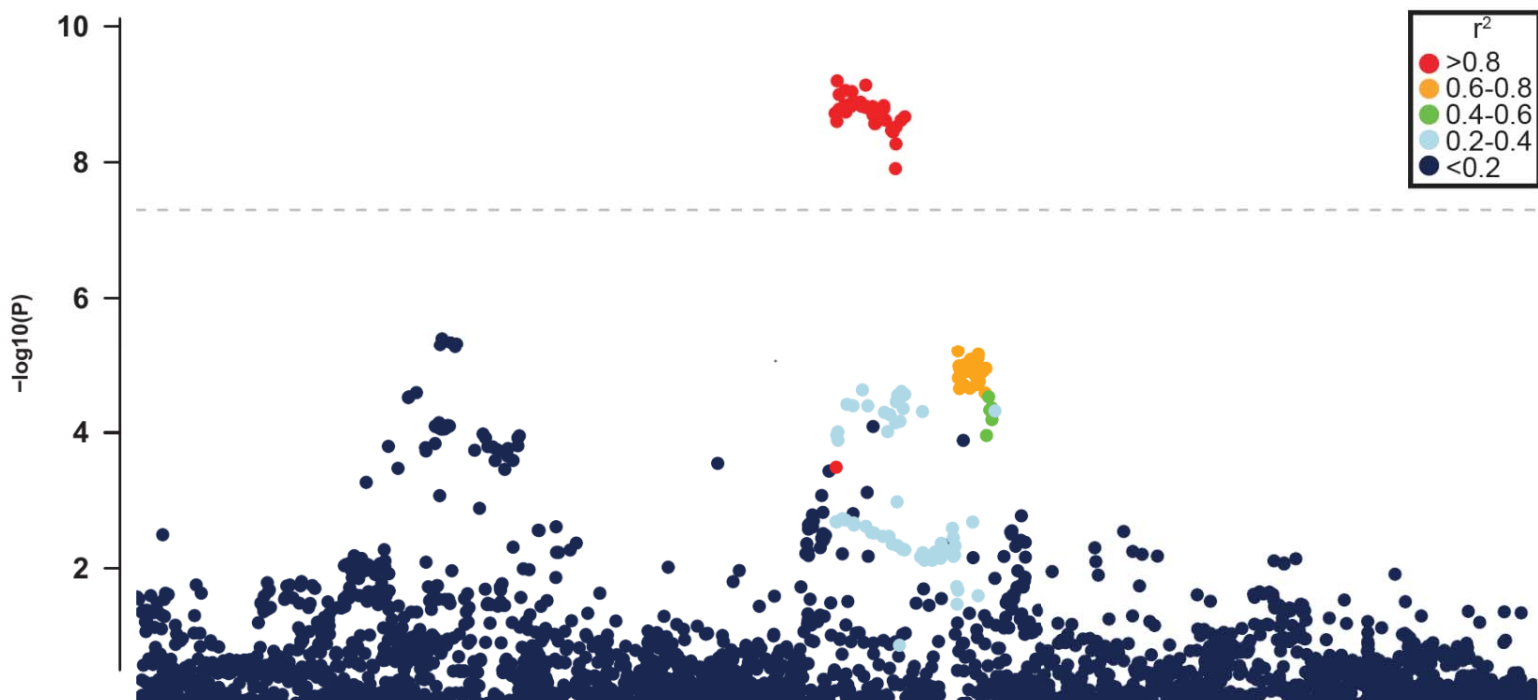
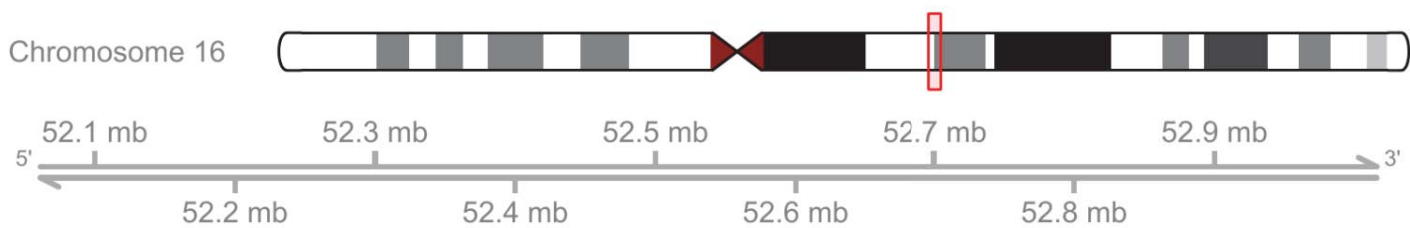
CP>GZ

Corrected for total SA

Uncorrected



Caudal Middle Frontal SA: rs7184835



C16orf97

TOX3

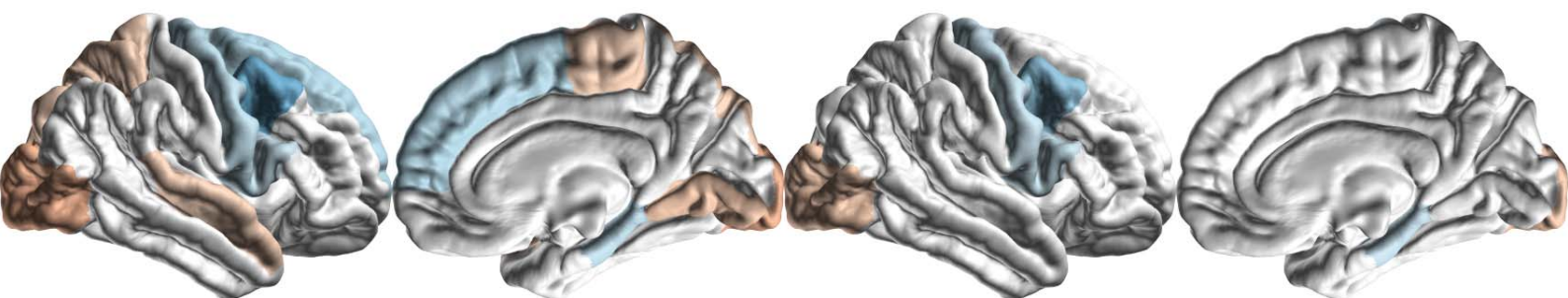
GZ>CP

CP>GZ

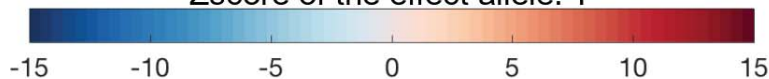


Corrected for total SA

Uncorrected

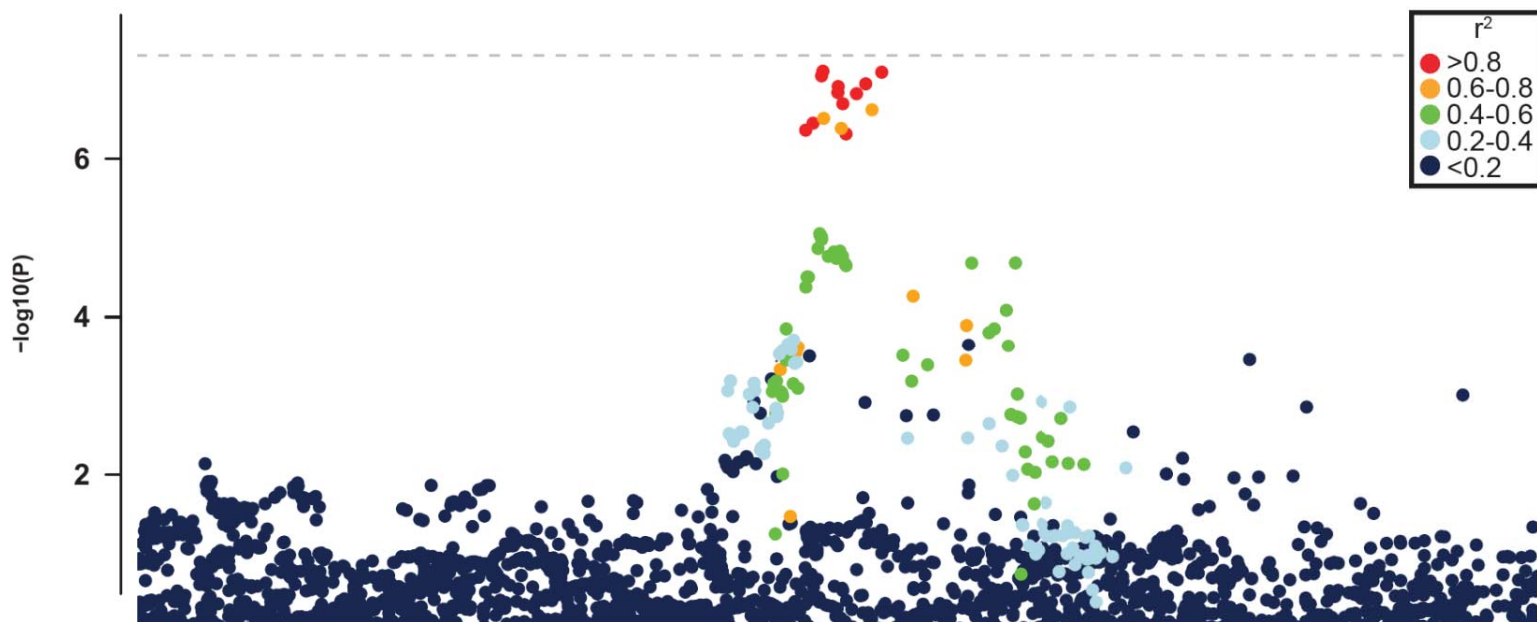


Zscore of the effect allele: T



Caudal Middle Frontal SA: rs1893430

Chromosome 18



RAB27B

TCF4

CCDC68

GZ>CP

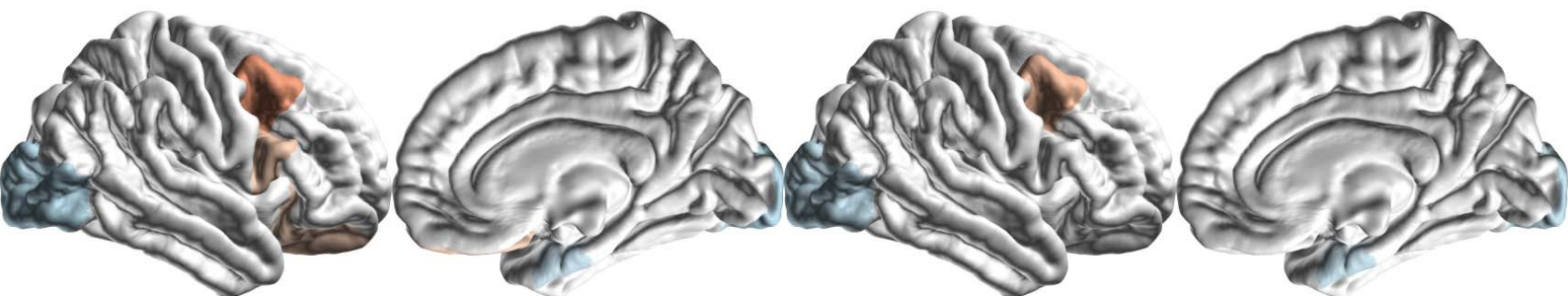


CP>GZ



Corrected for total SA

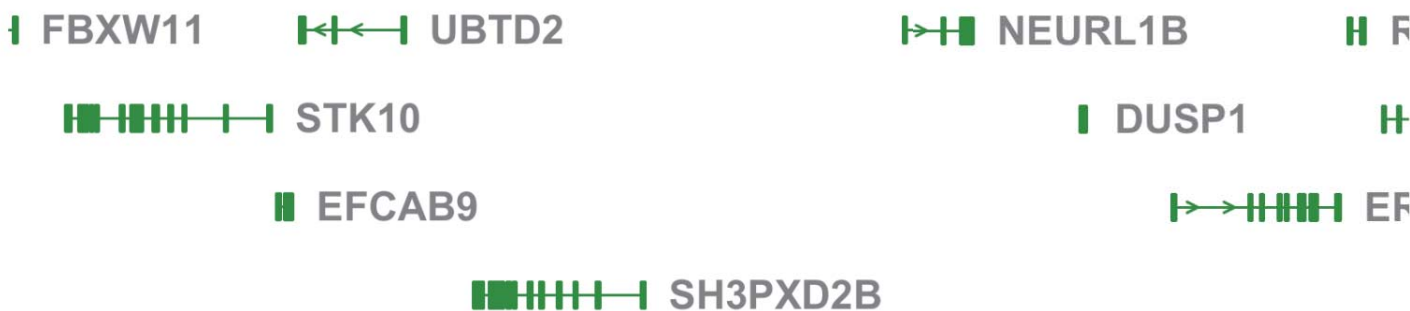
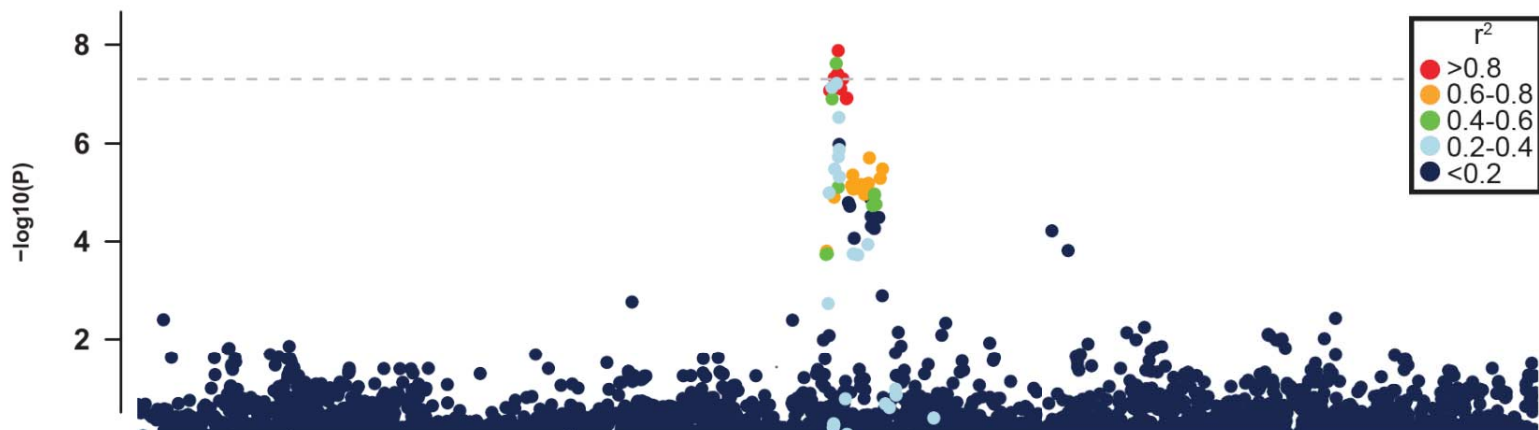
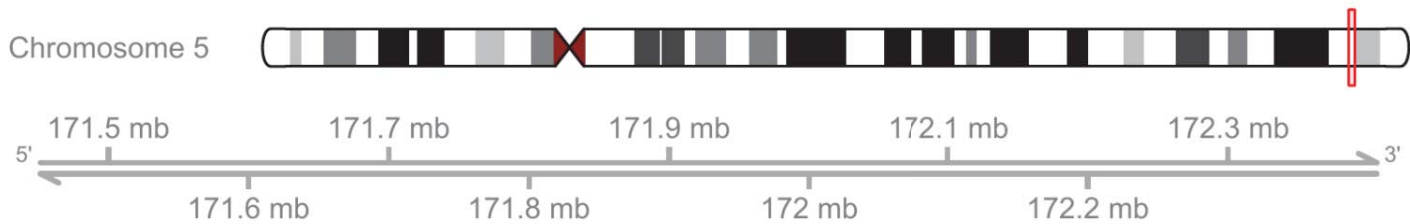
Uncorrected



Zscore of the effect allele: A

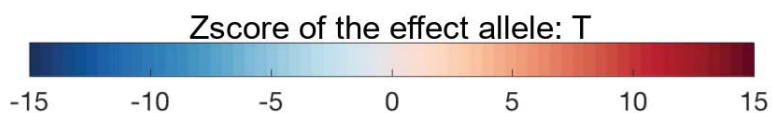
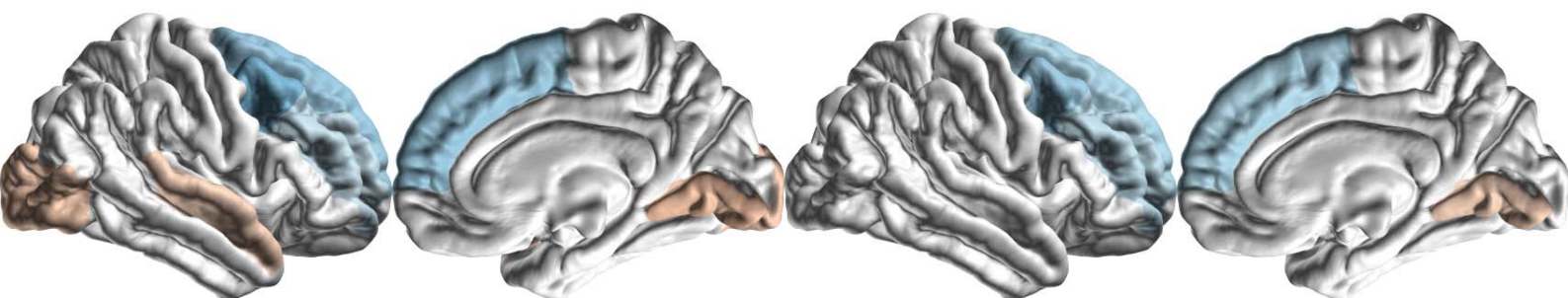


Caudal Middle Frontal TH: rs62385486

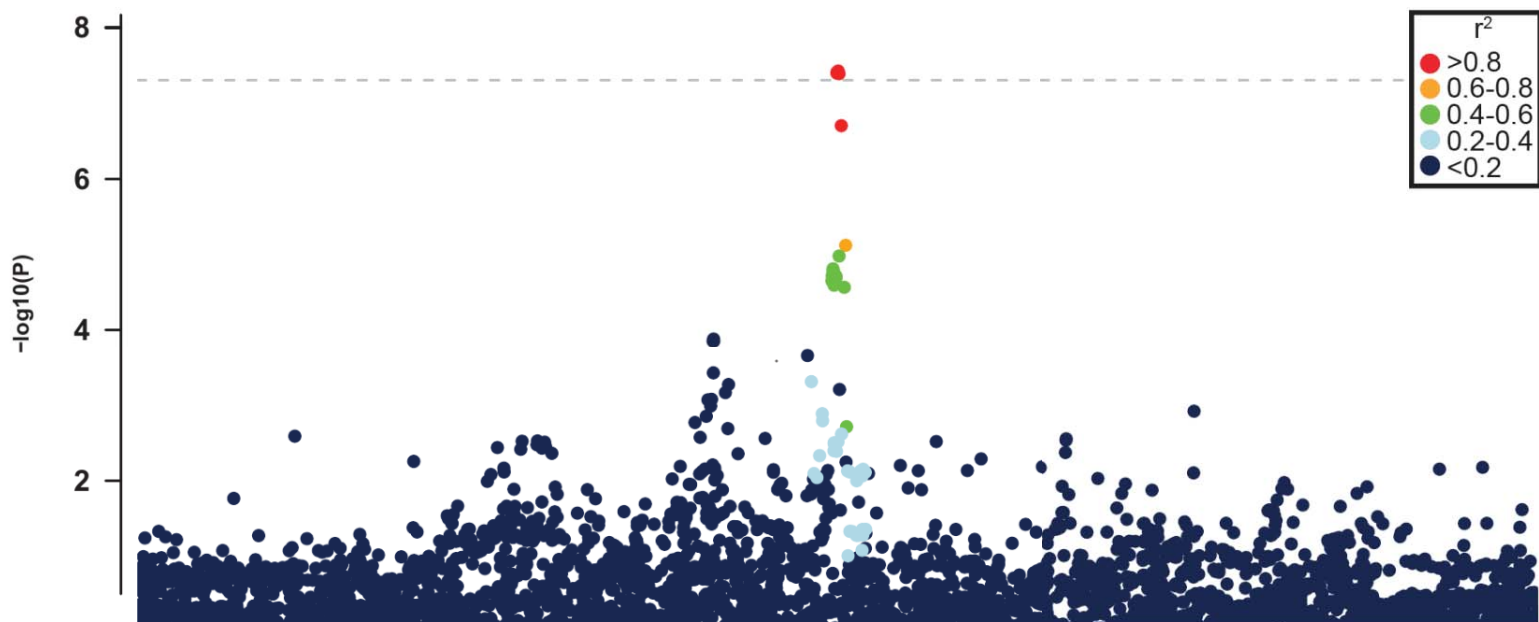
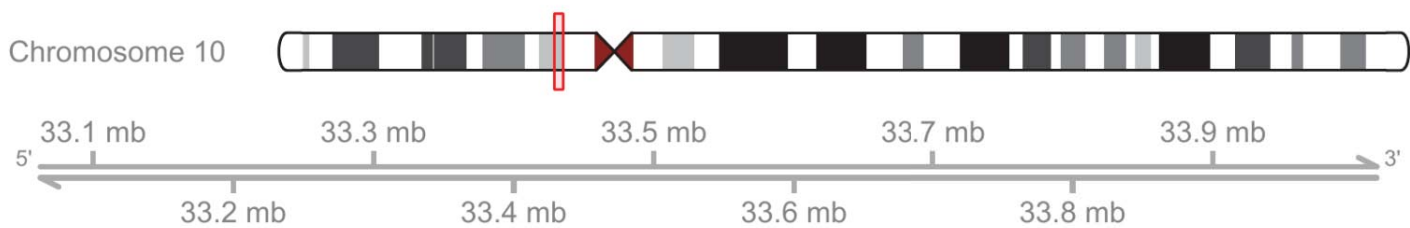


Corrected for average TH

Uncorrected



Paracentral SA: rs2269084



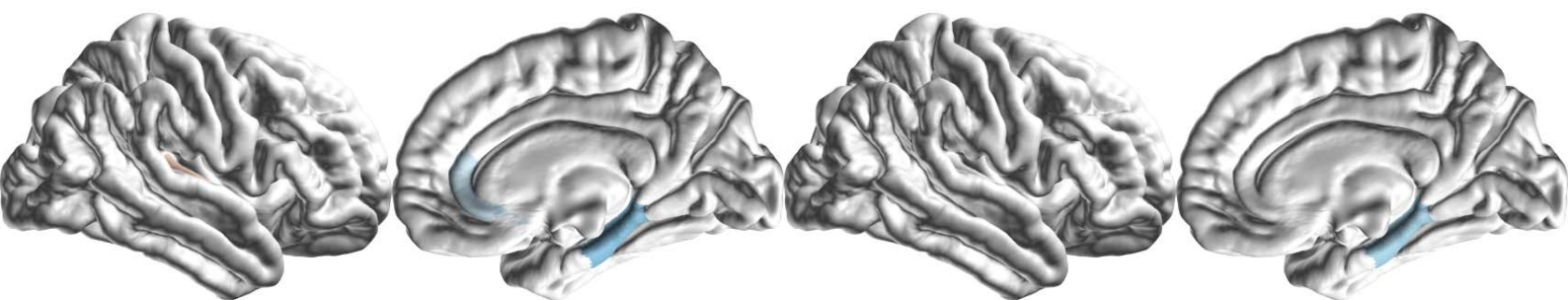
GZ>CP

CP>GZ

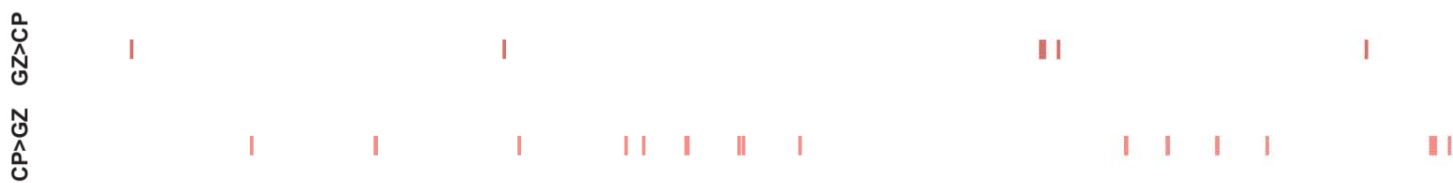
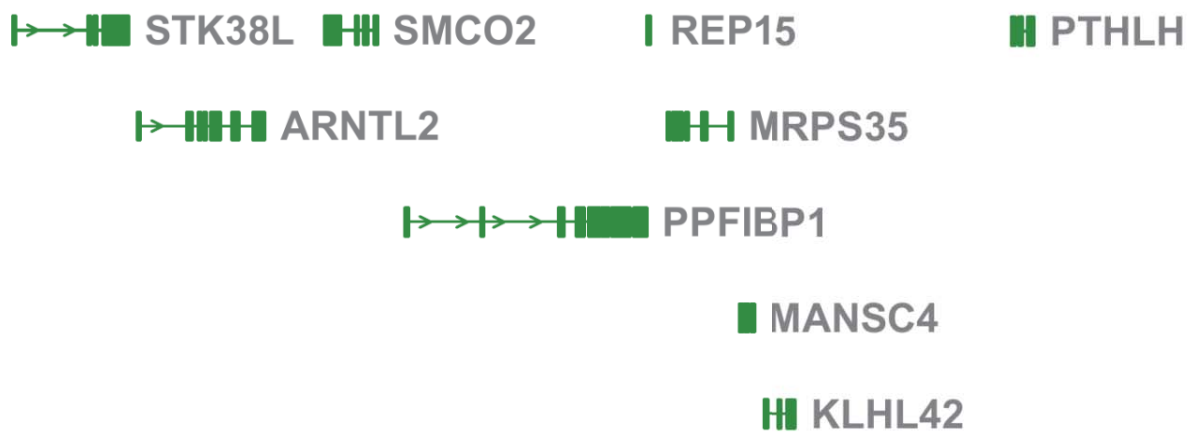
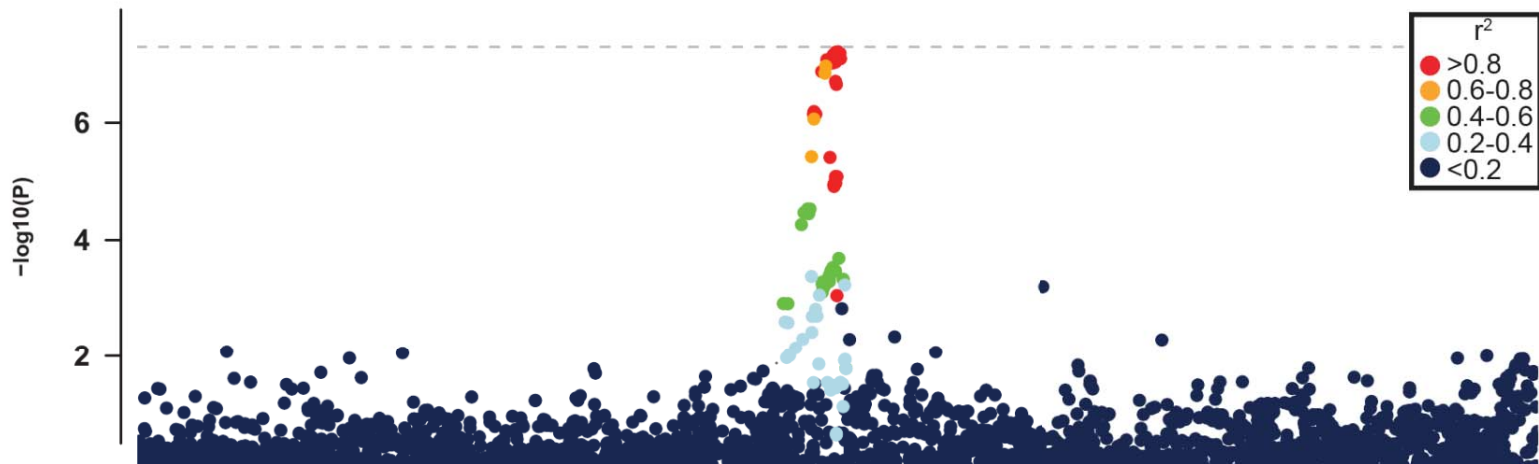


Corrected for total SA

Uncorrected

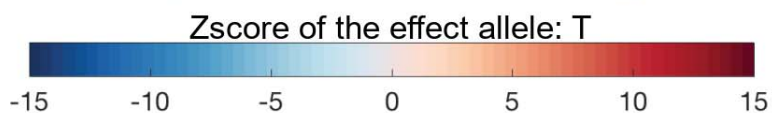
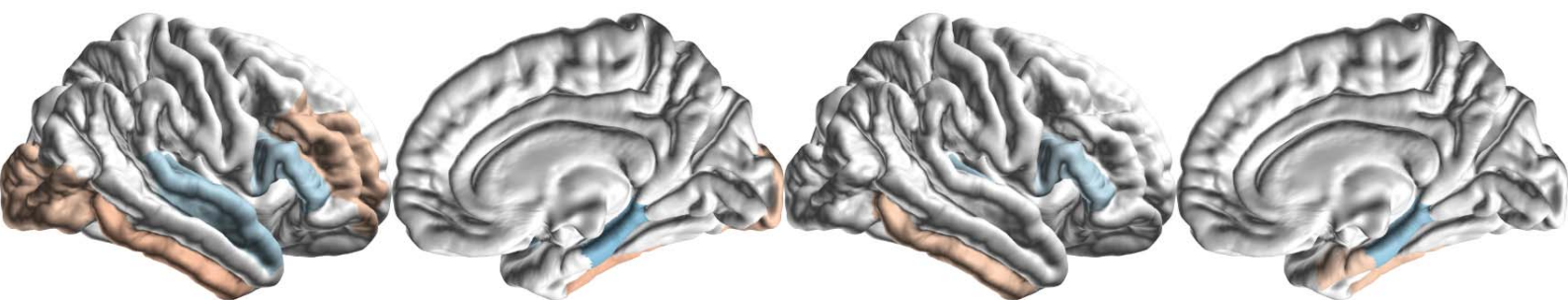


Paracentral SA: rs7303526

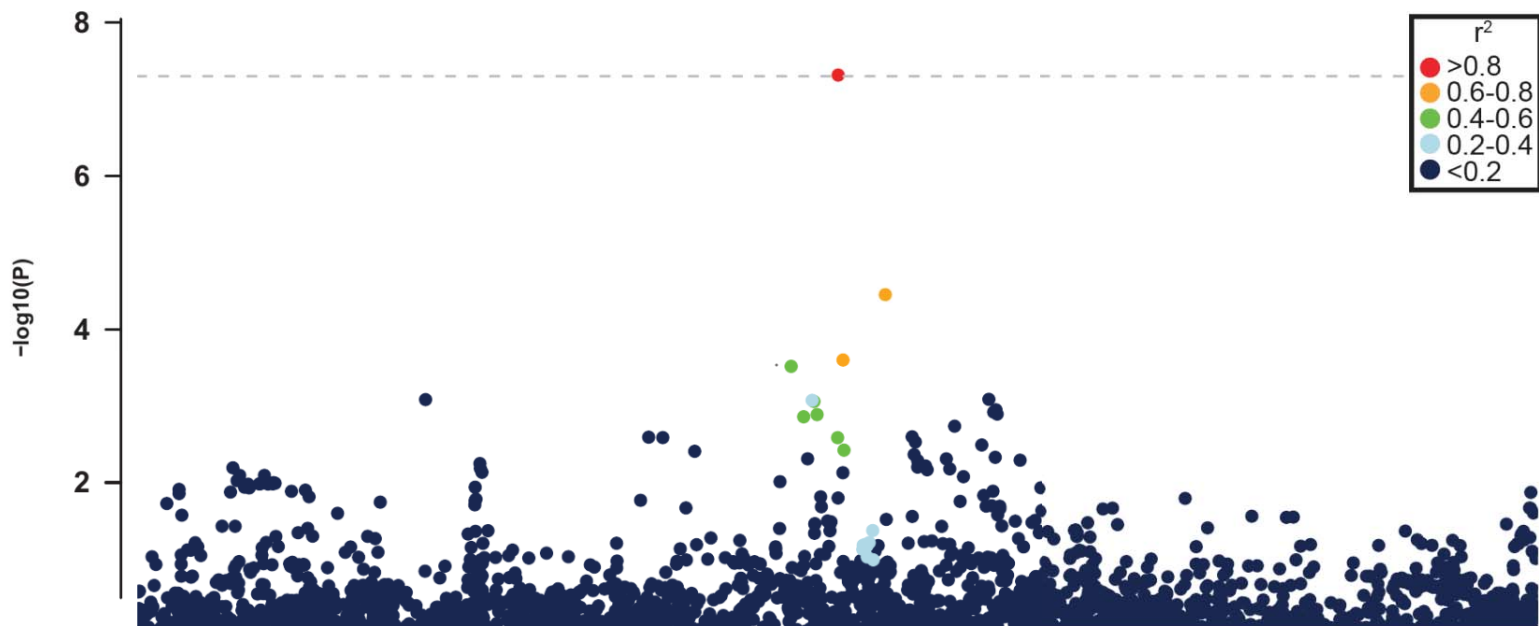
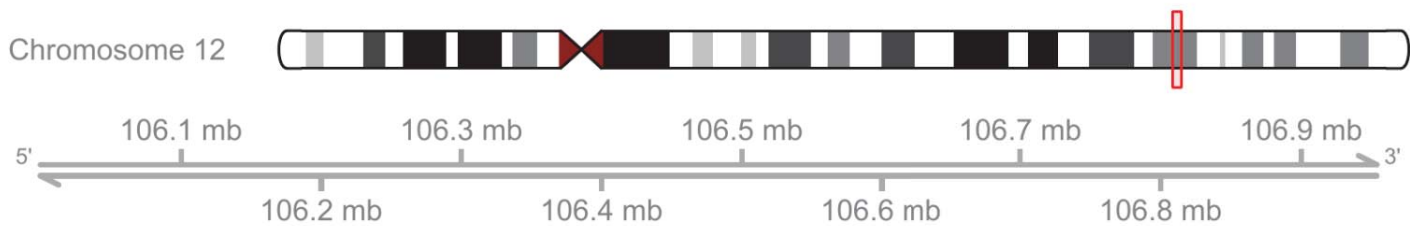


Corrected for total SA

Uncorrected

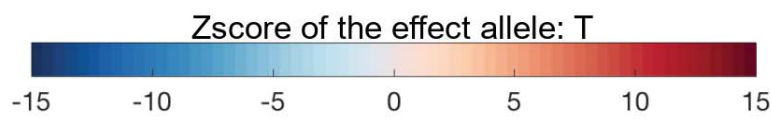
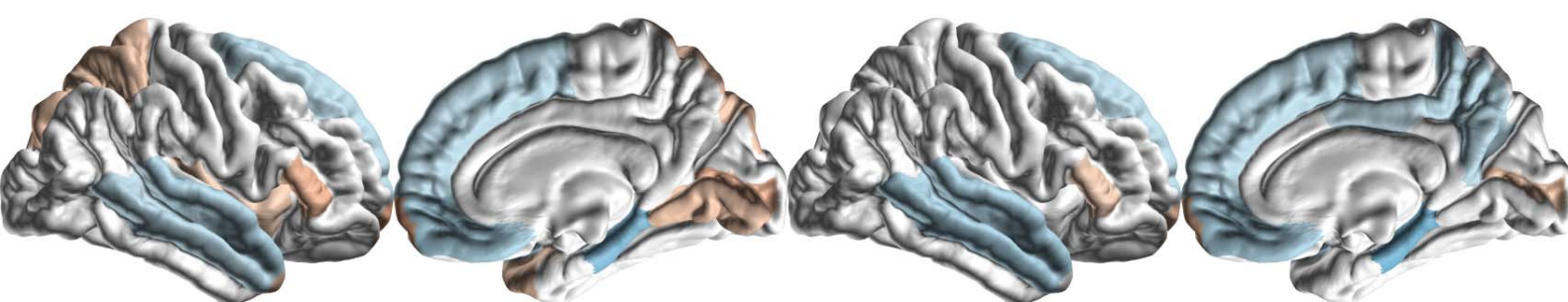


Paracentral SA: rs12146713



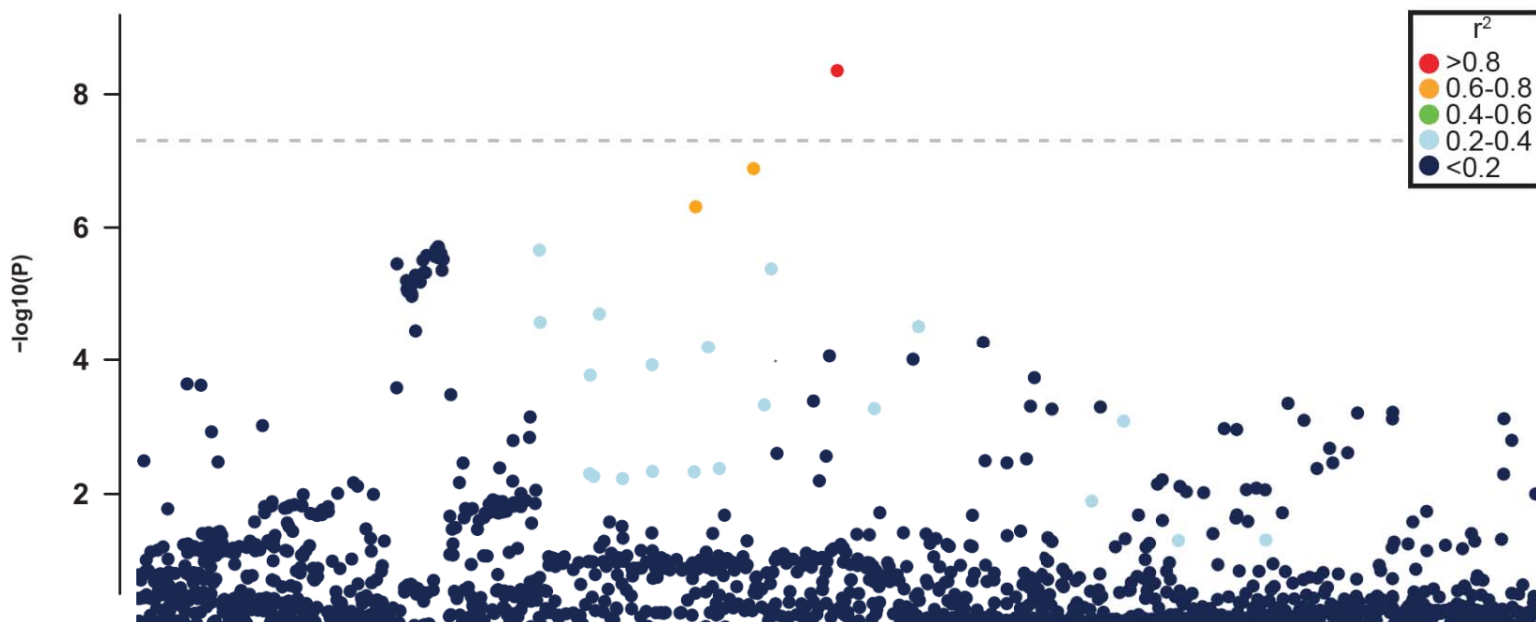
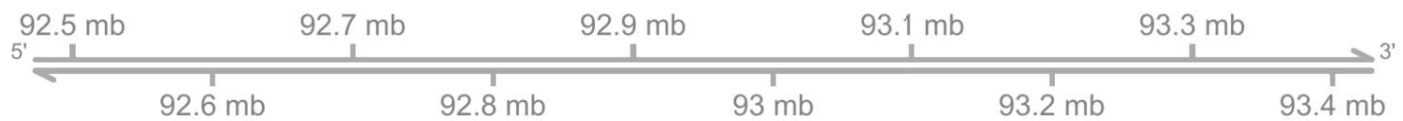
Corrected for total SA

Uncorrected



Precentral SA: rs10064431

Chromosome 5



NR2F1 POU5F2



GZ>CP

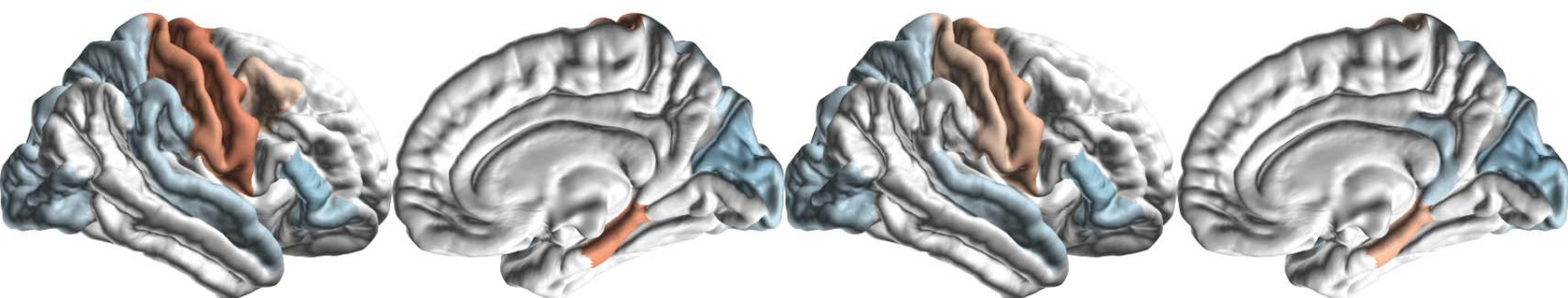


CP>GZ

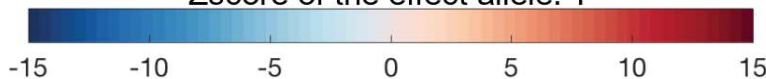


Corrected for total SA

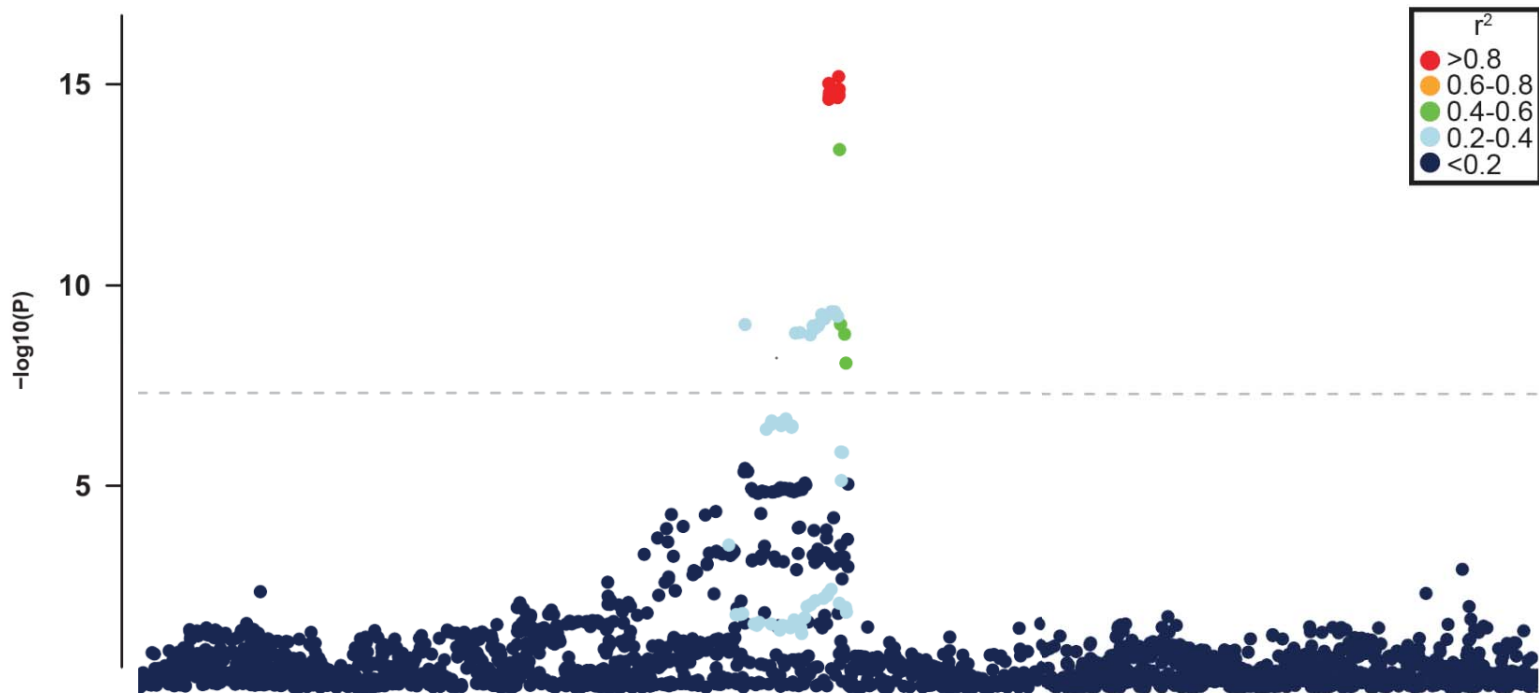
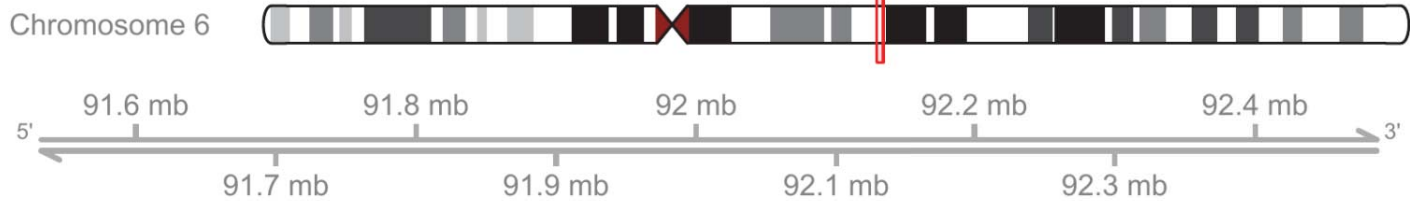
Uncorrected



Zscore of the effect allele: T



Precentral SA: rs4706392

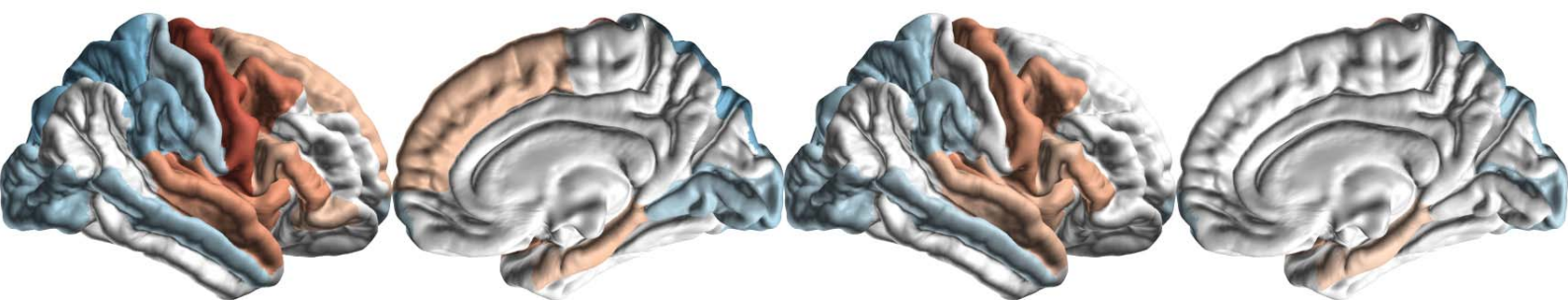


GZ>CP

CP>GZ

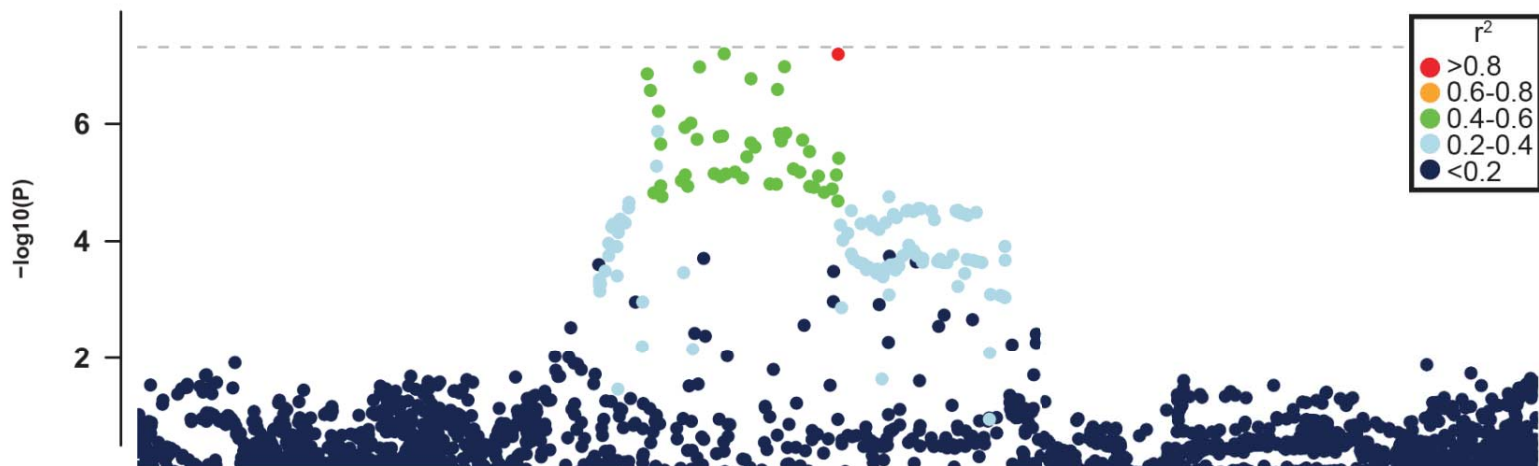
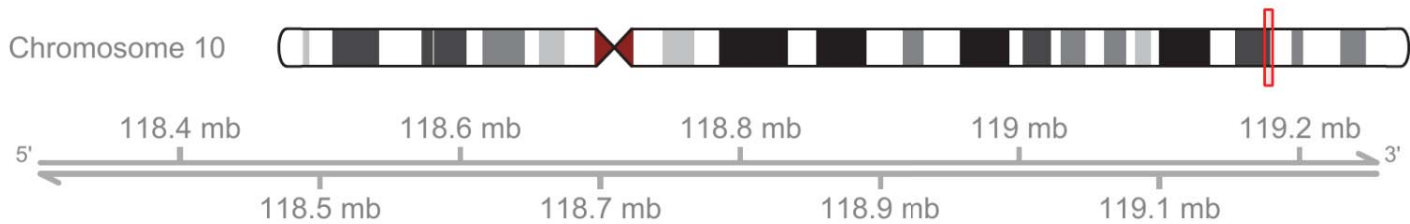
Corrected for total SA

Uncorrected



-15 -10 -5 0 5 10 15

Precentral SA: rs11197866

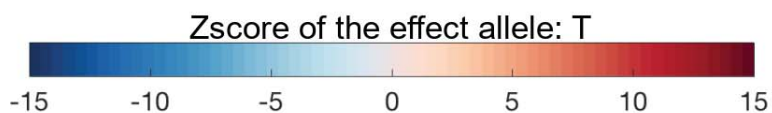
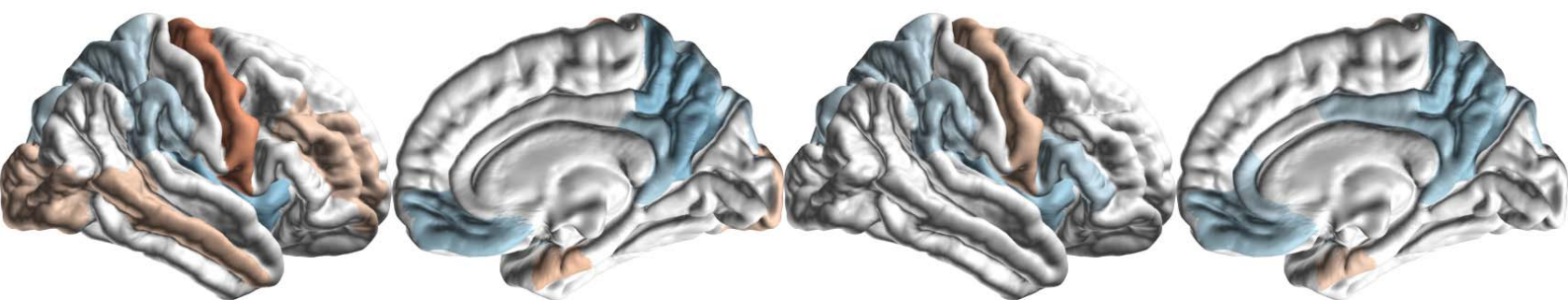


CP>GZ GZ>CP



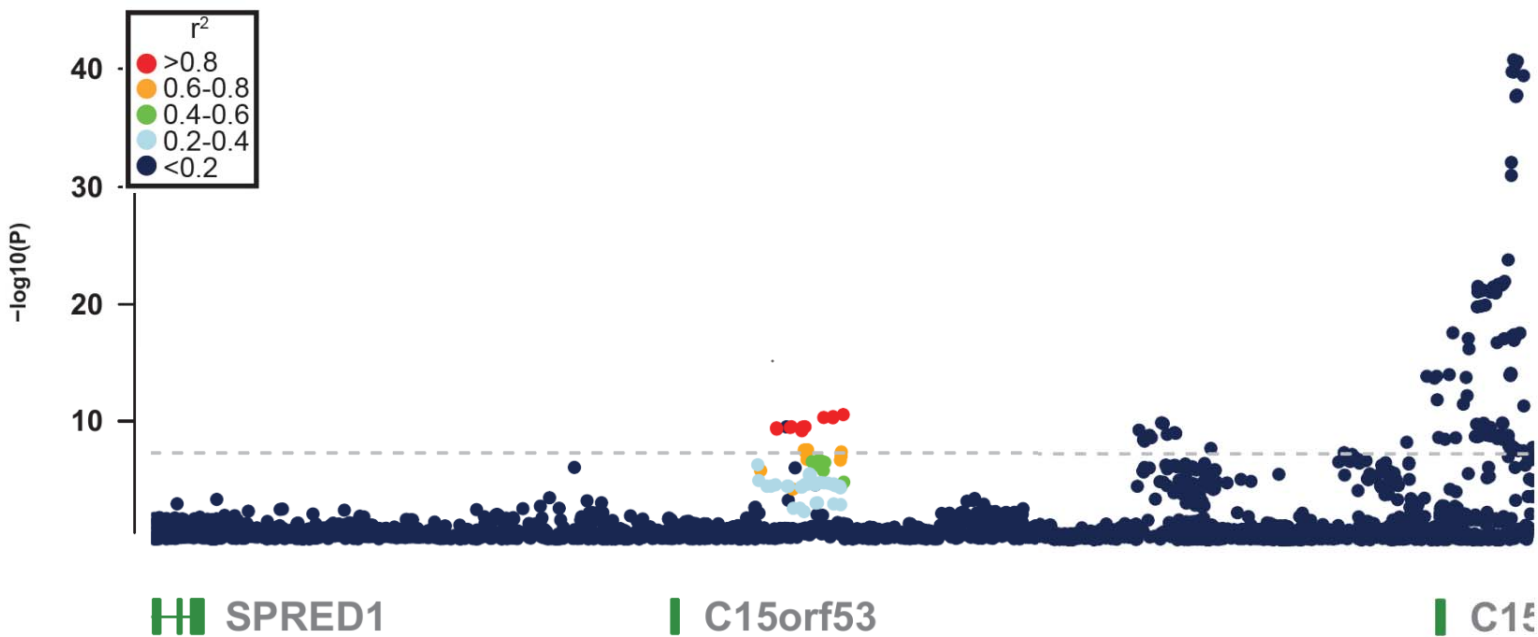
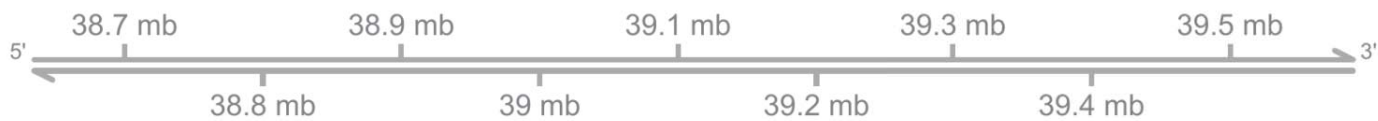
Corrected for total SA

Uncorrected



Precentral SA: rs17601140

Chromosome 15



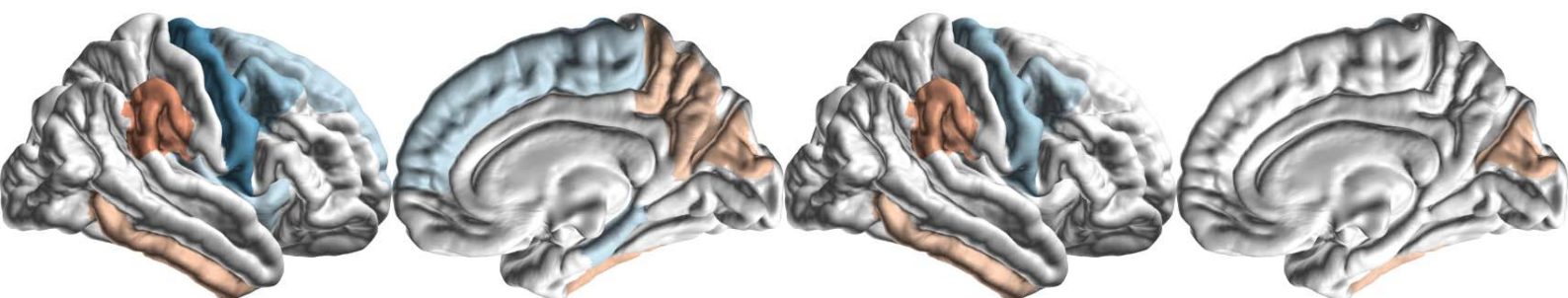
FAM98B

RASGRP1

GZ>CP
CP>GZ

Corrected for total SA

Uncorrected

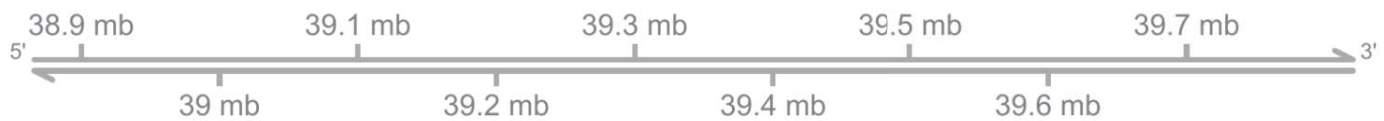


Zscore of the effect allele: C



Precentral SA: rs11070172

Chromosome 15



RASGRP1

C15orf54

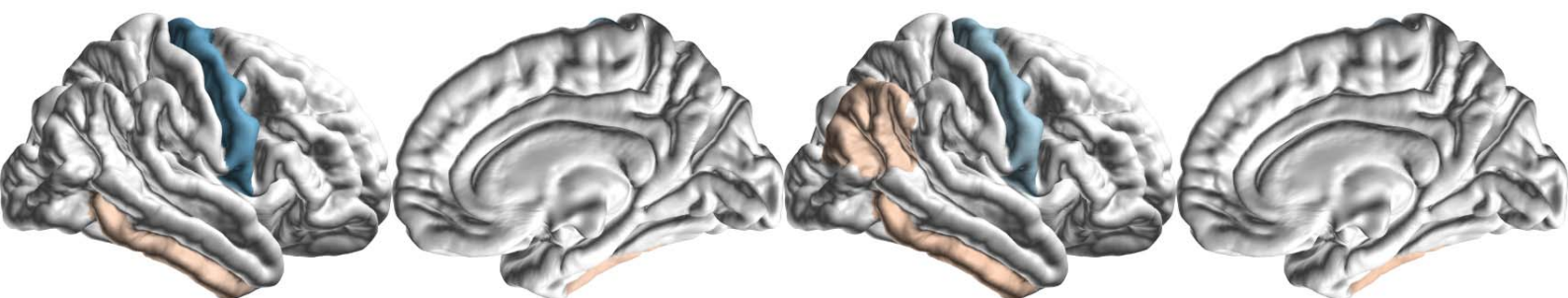
C15orf53

GZ>CP

CP>GZ

Corrected for total SA

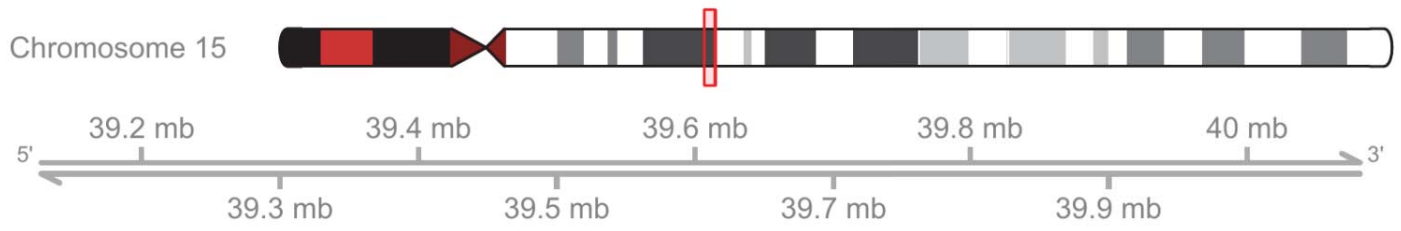
Uncorrected



Zscore of the effect allele: T

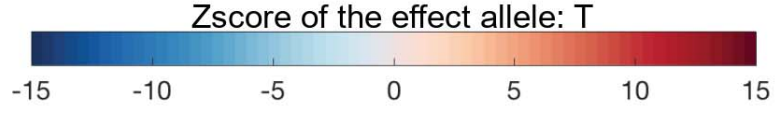
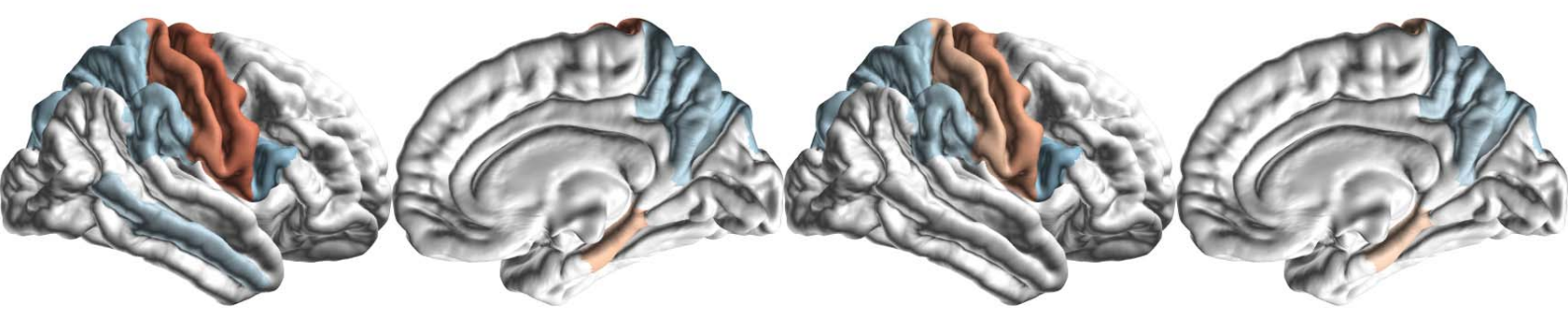


Precentral SA: rs3862145



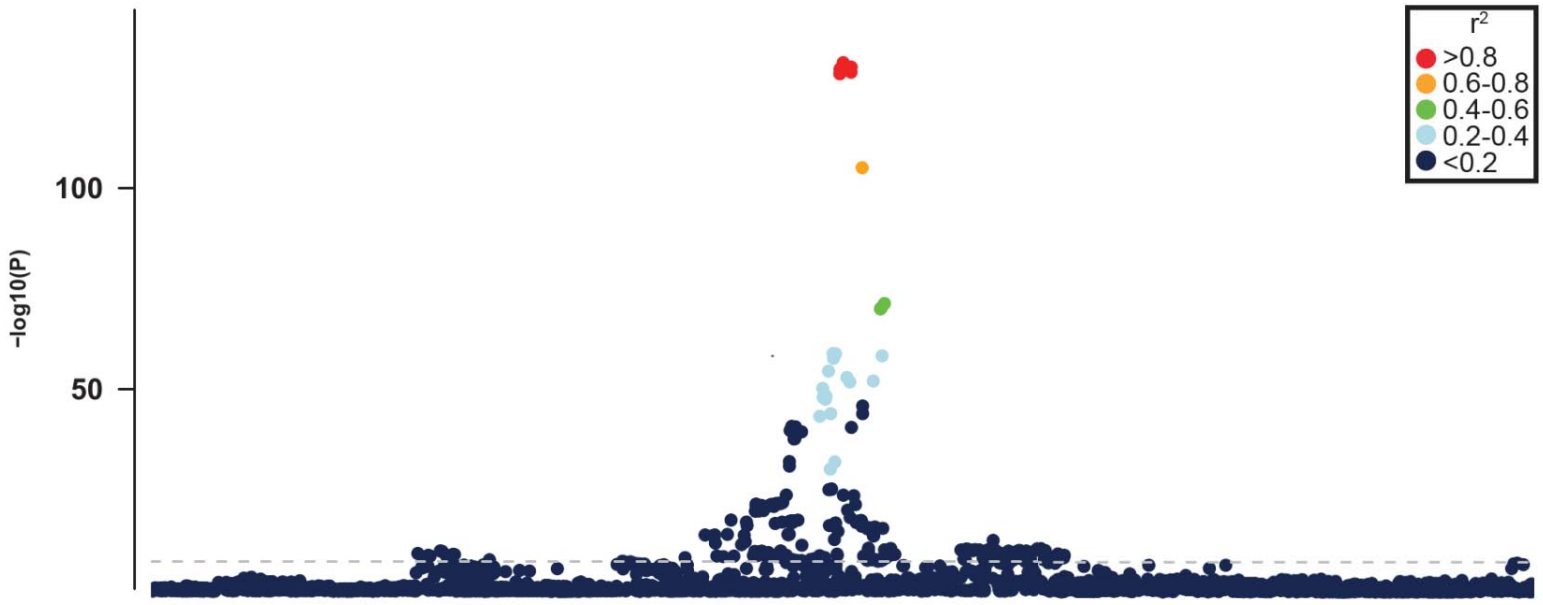
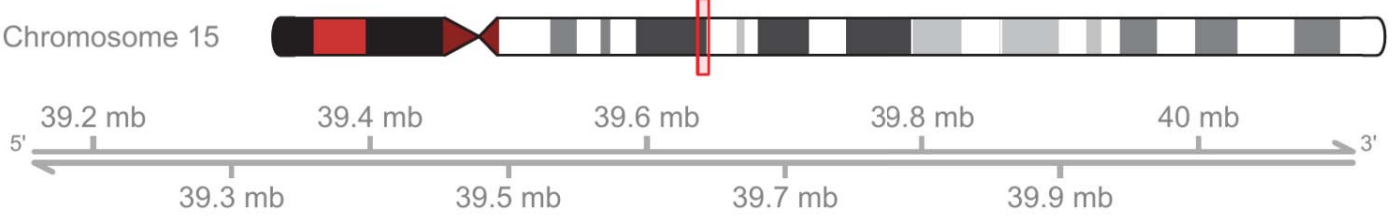
Corrected for total SA

Uncorrected



Precentral SA: rs1080066

Chromosome 15



C15orf54

THBS1

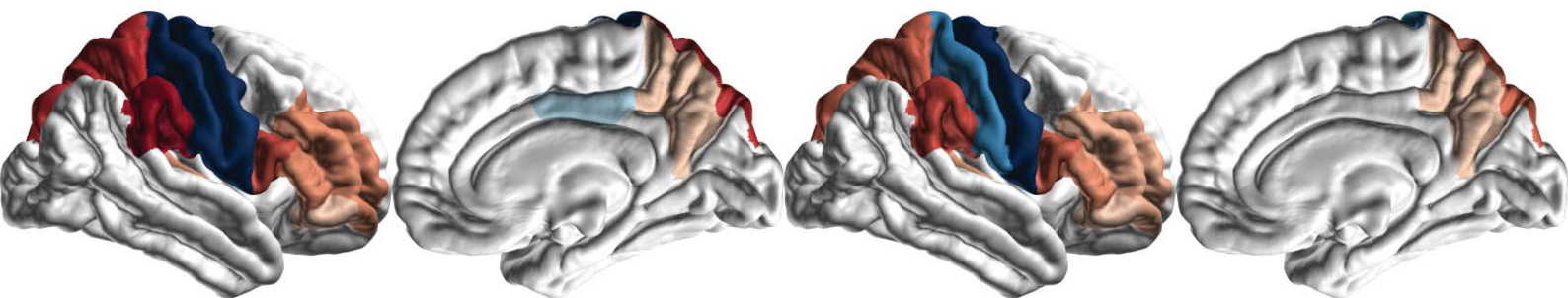
FSII

GZ>CP

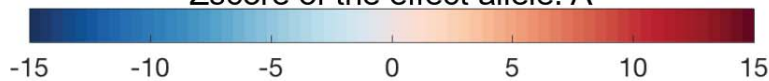
CP>GZ

Corrected for total SA

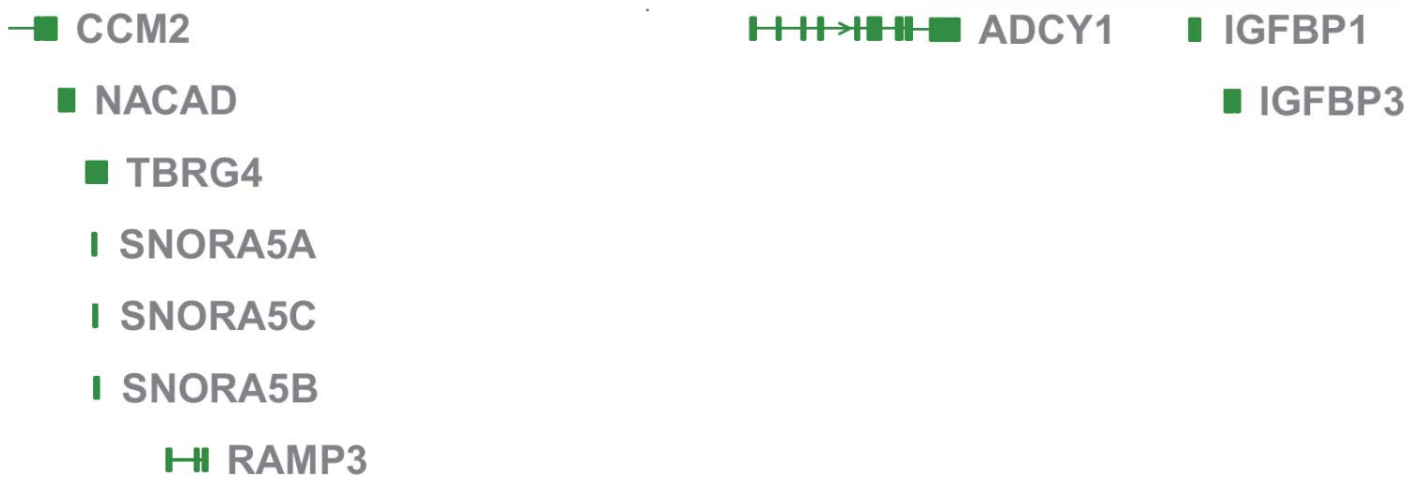
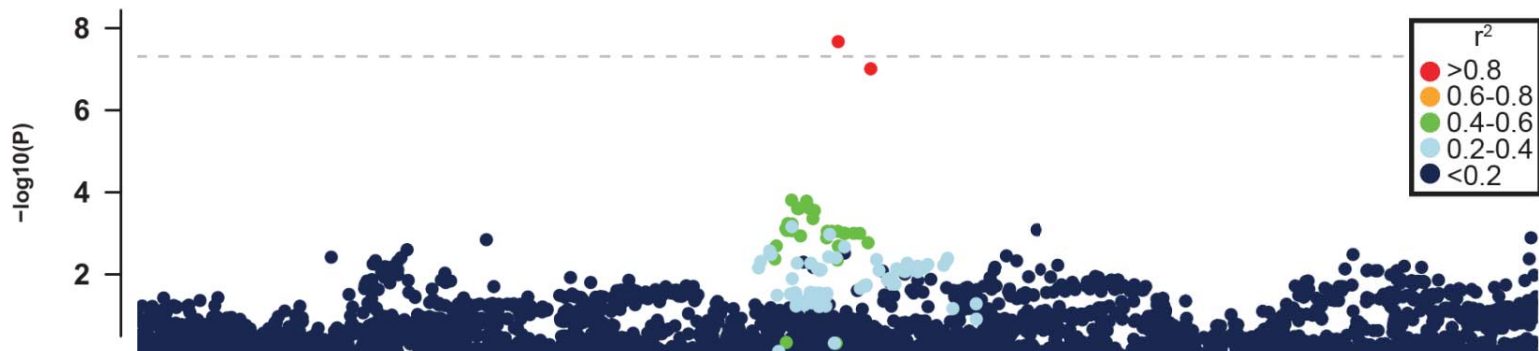
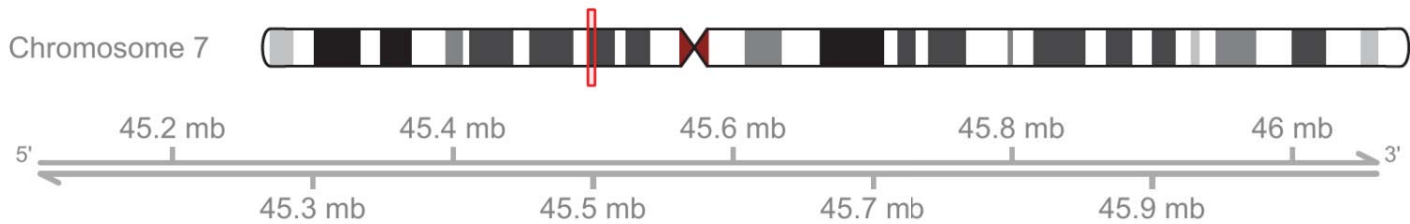
Uncorrected



Zscore of the effect allele: A

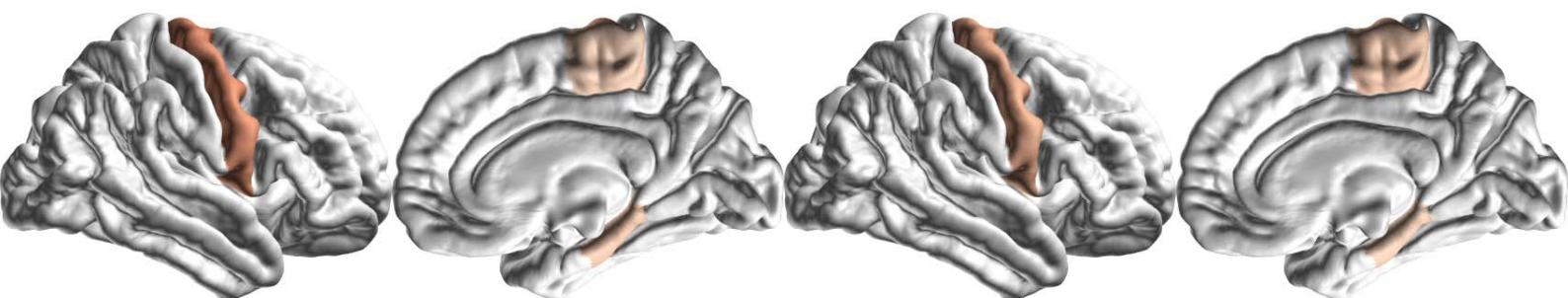


Precentral TH: rs4724413



Corrected for average TH

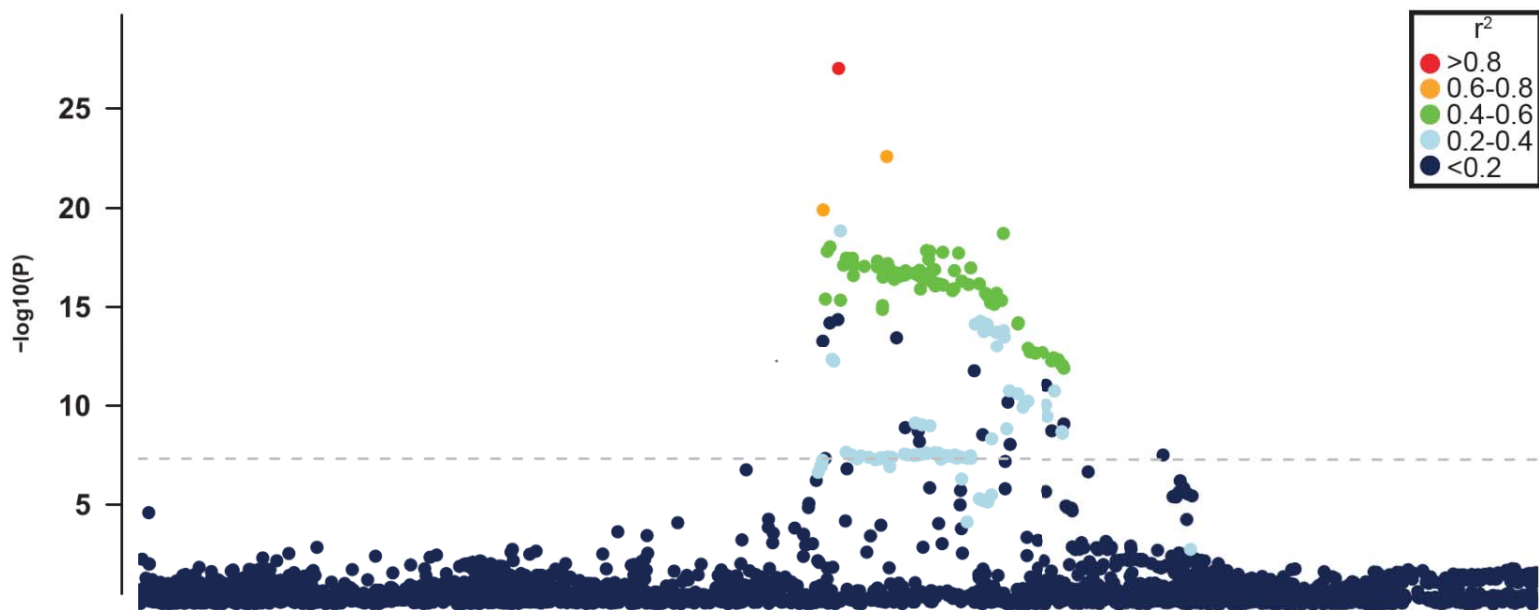
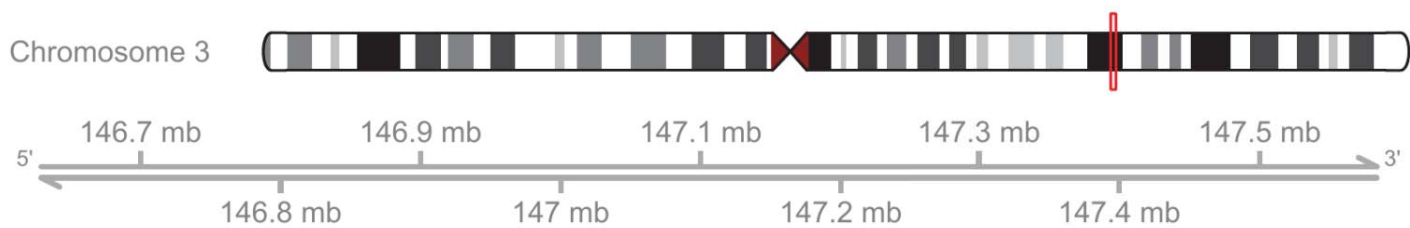
Uncorrected



Zscore of the effect allele: T



Postcentral SA: rs2279829



ZIC4

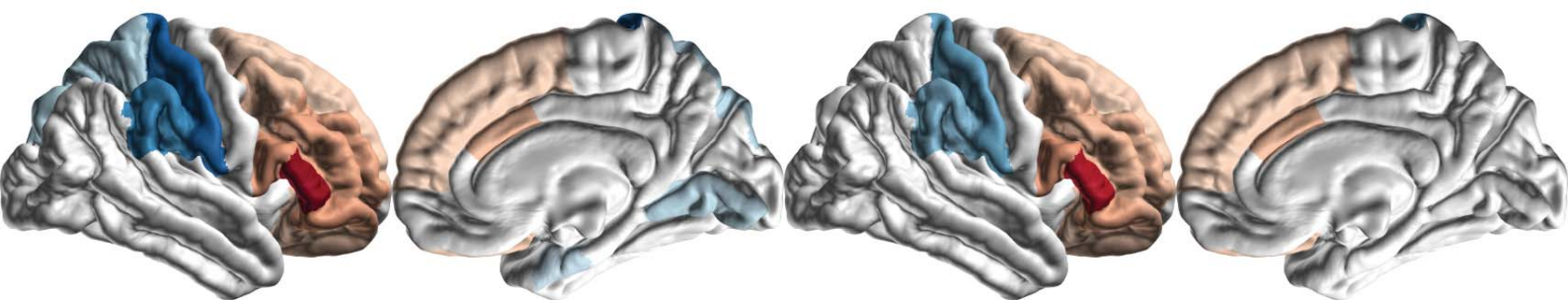
ZIC1

GZ>CP

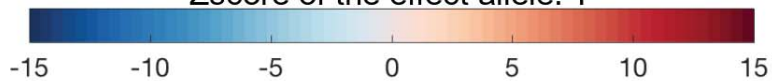
CP>GZ

Corrected for total SA

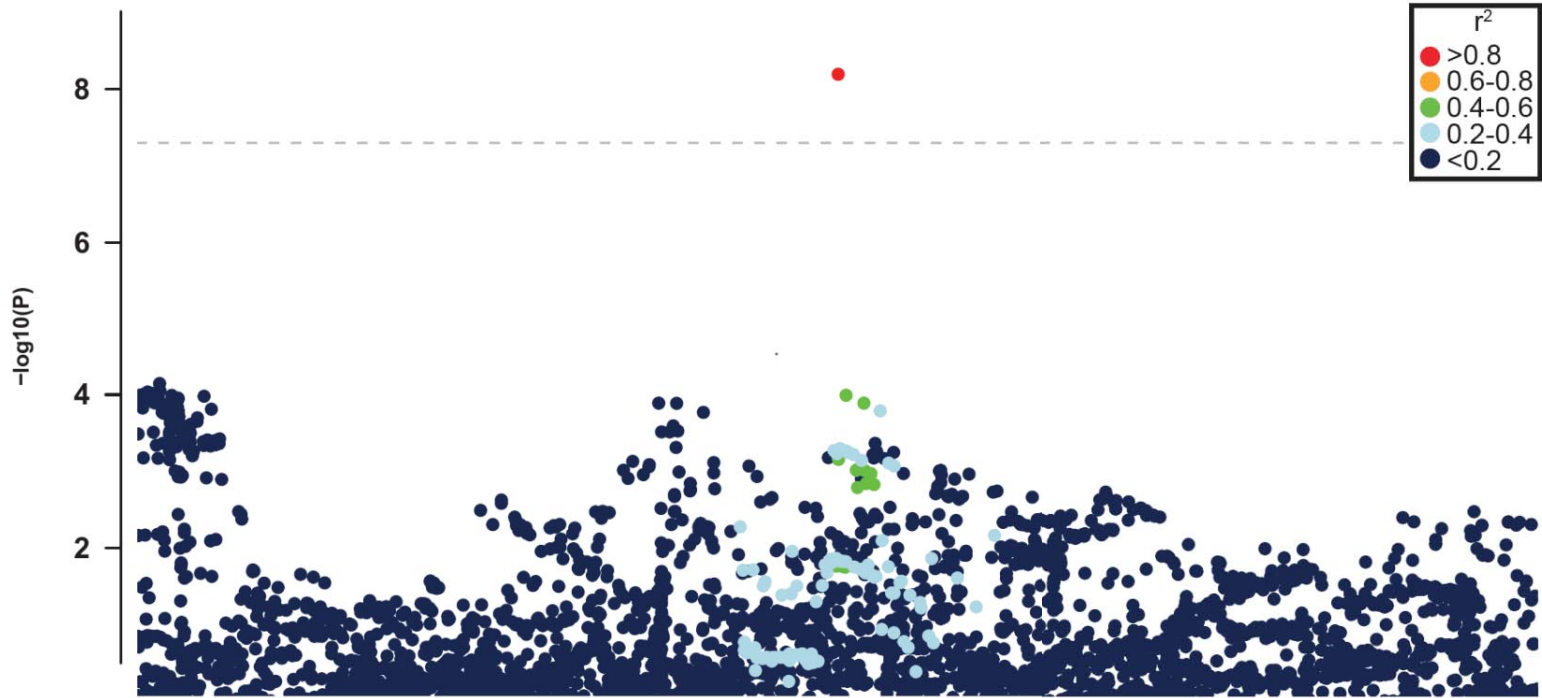
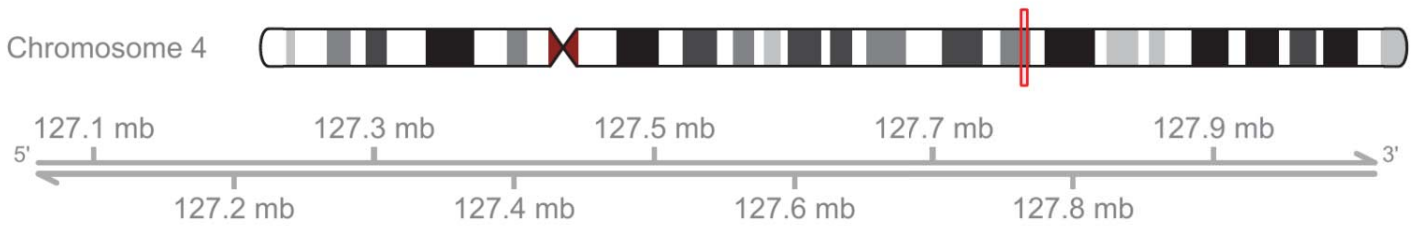
Uncorrected



Zscore of the effect allele: T



Postcentral SA: rs313135



GZ>CP

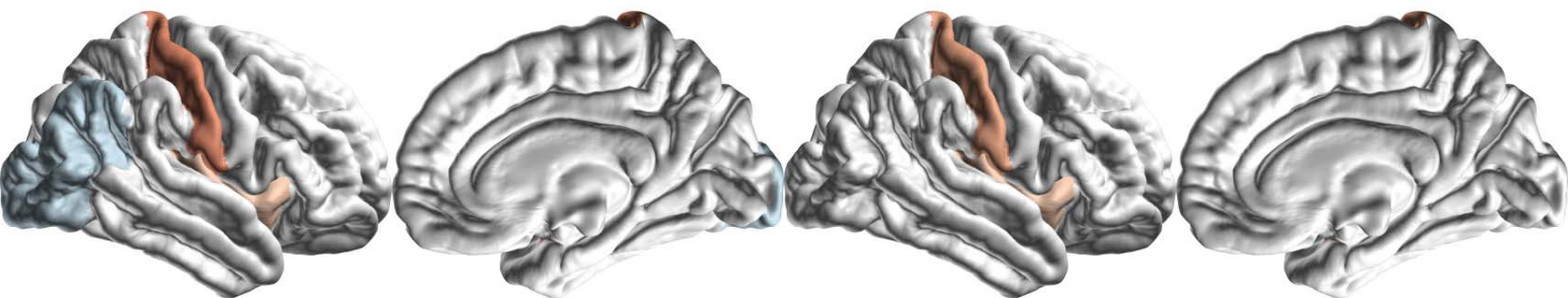


CP>GZ

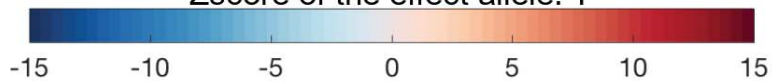


Corrected for total SA

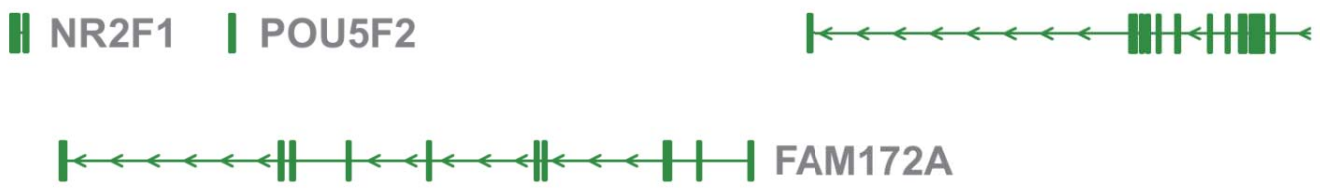
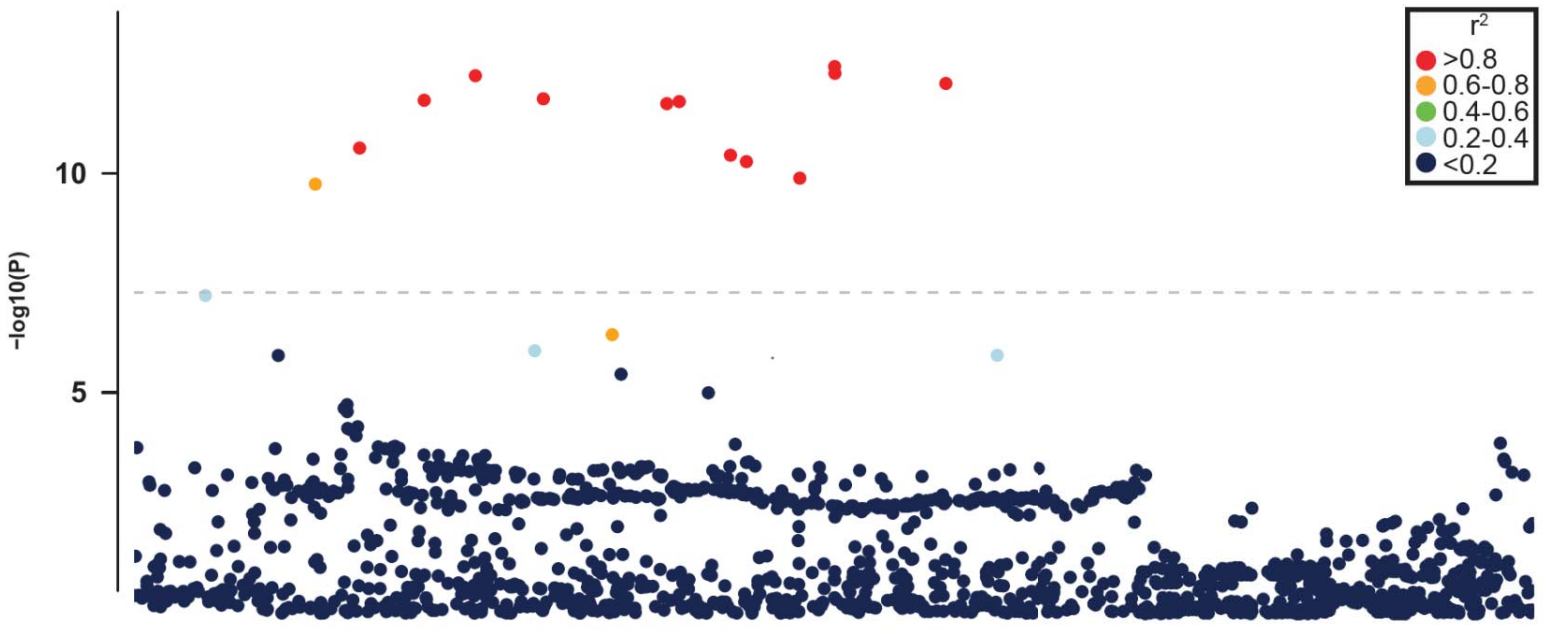
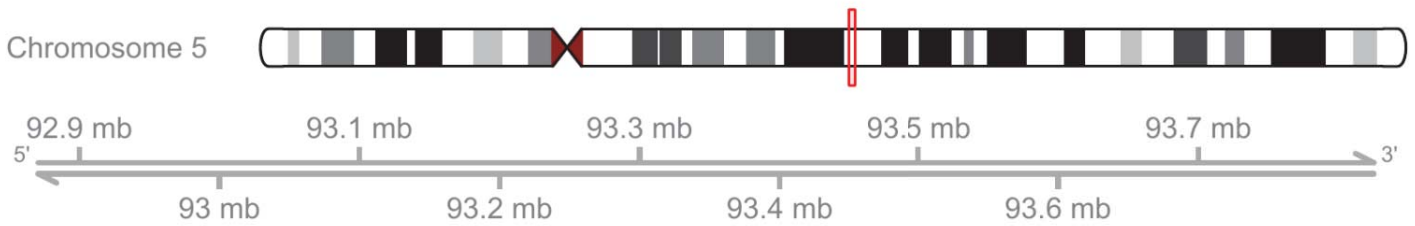
Uncorrected



Zscore of the effect allele: T

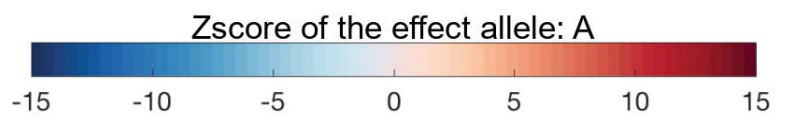
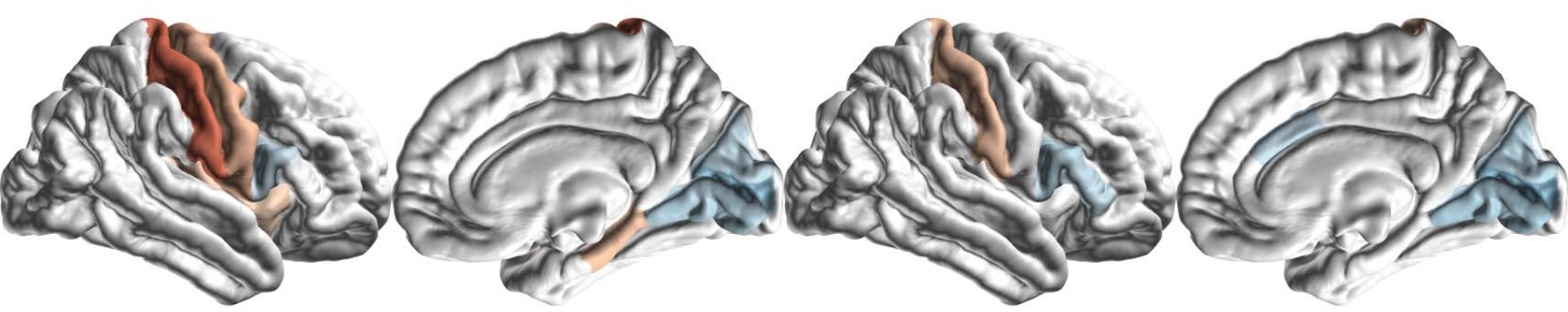


Postcentral SA: rs34322452



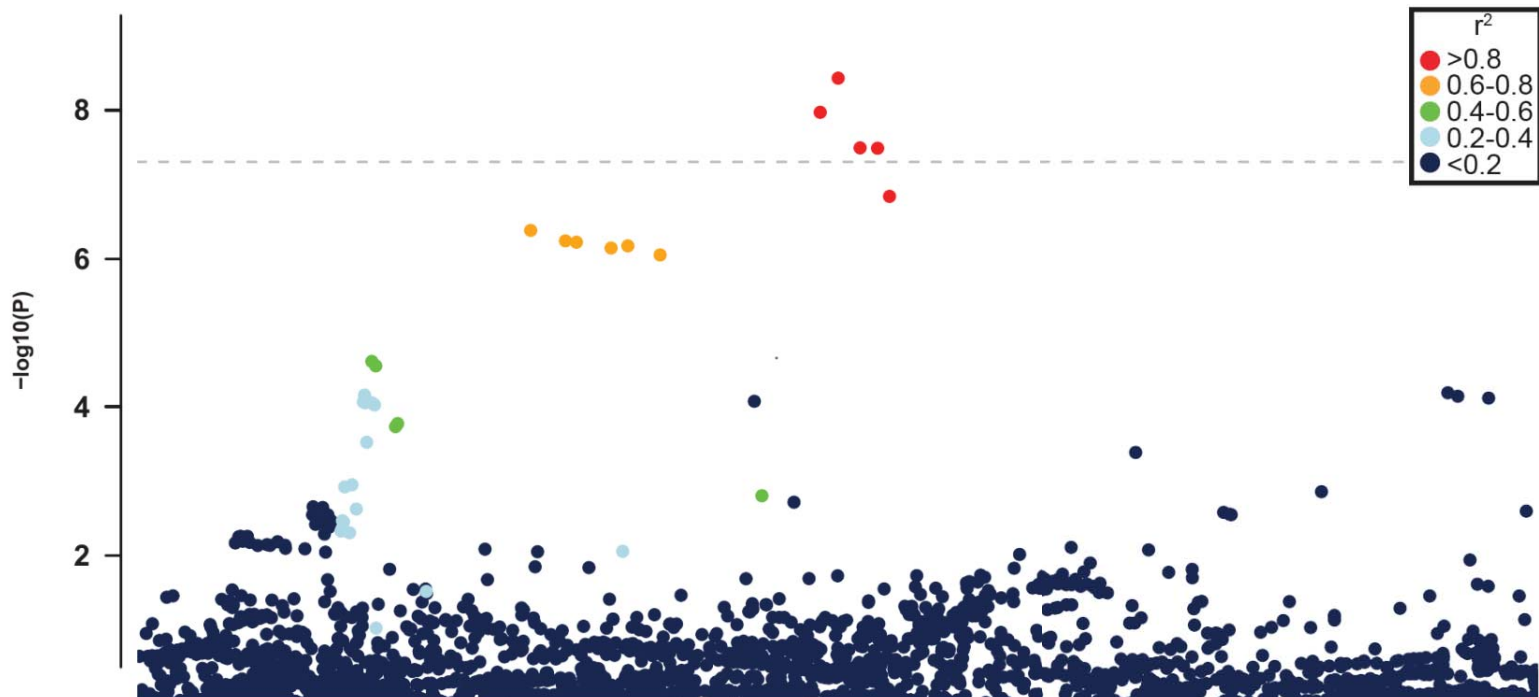
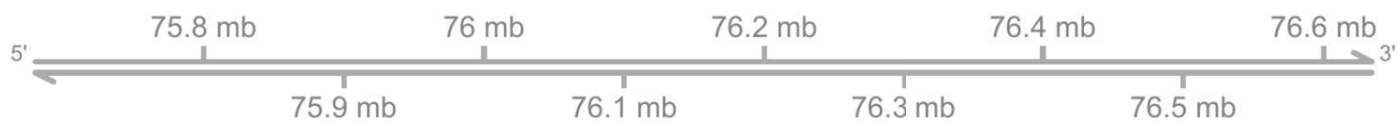
Corrected for total SA

Uncorrected



Postcentral SA: rs11789773

Chromosome 9



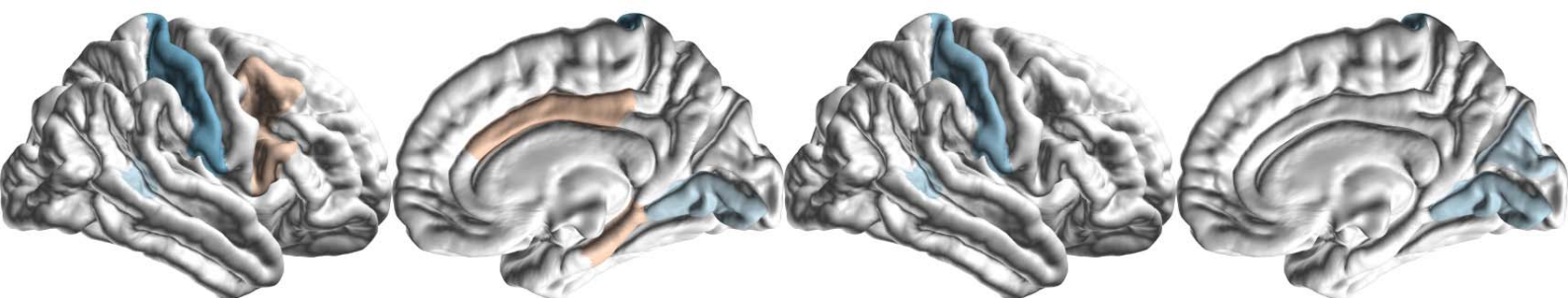
ANXA1

GZ>CP

CP>GZ

Corrected for total SA

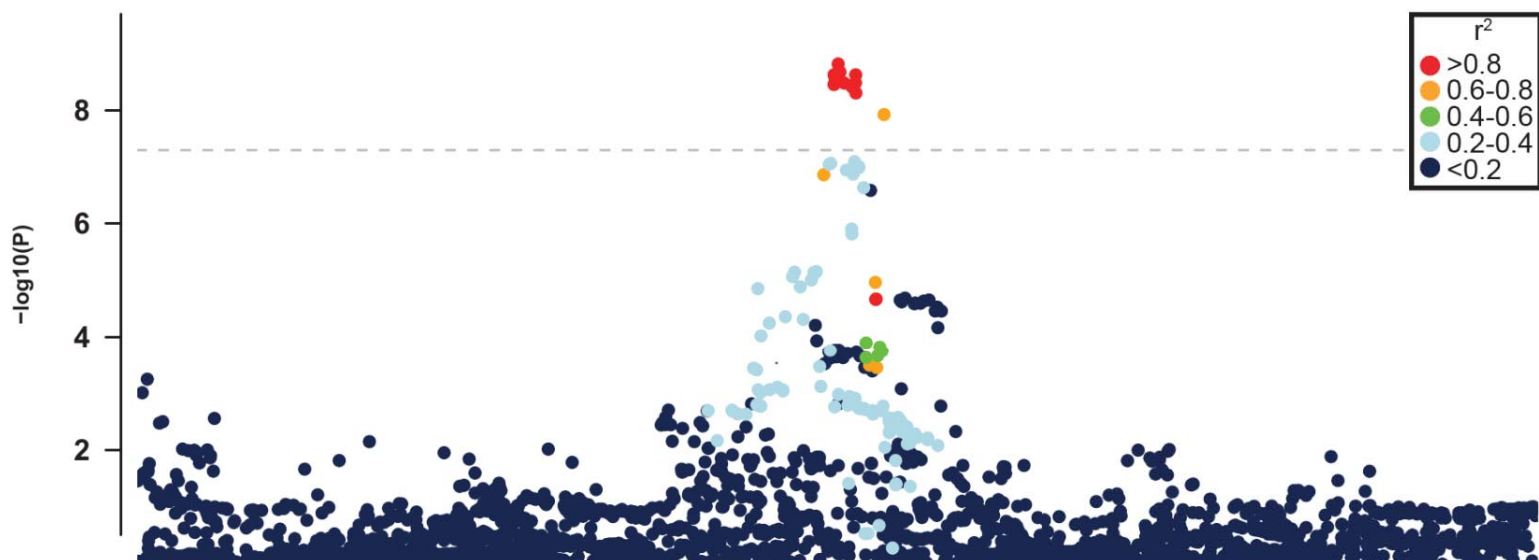
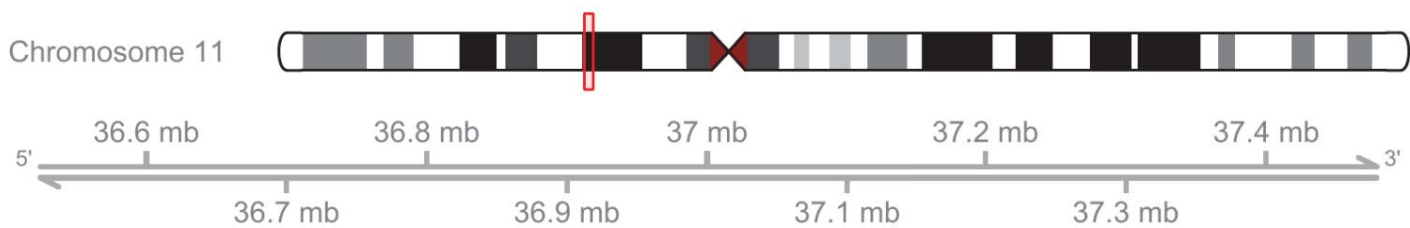
Uncorrected



Zscore of the effect allele: A



Postcentral SA: rs11033898



TRA1 RAG1

RAG2

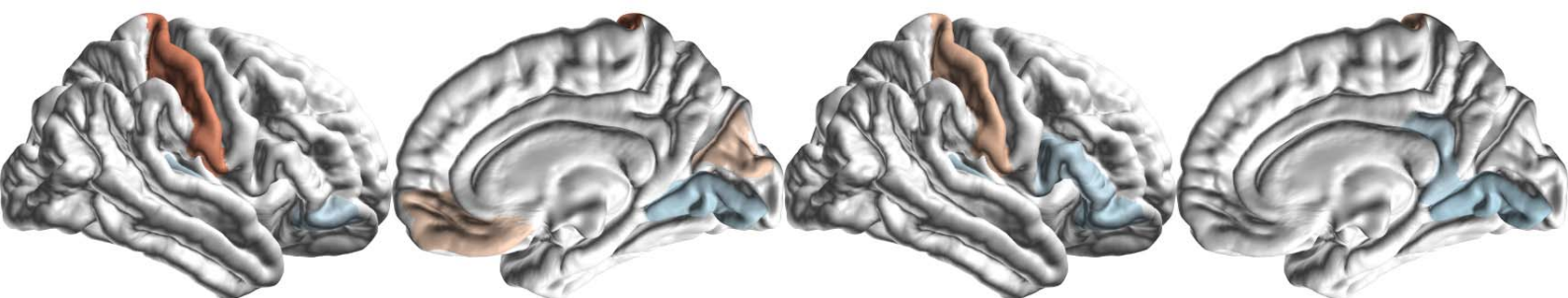
C11orf74

GZ>CP

CP>GZ

Corrected for total SA

Uncorrected

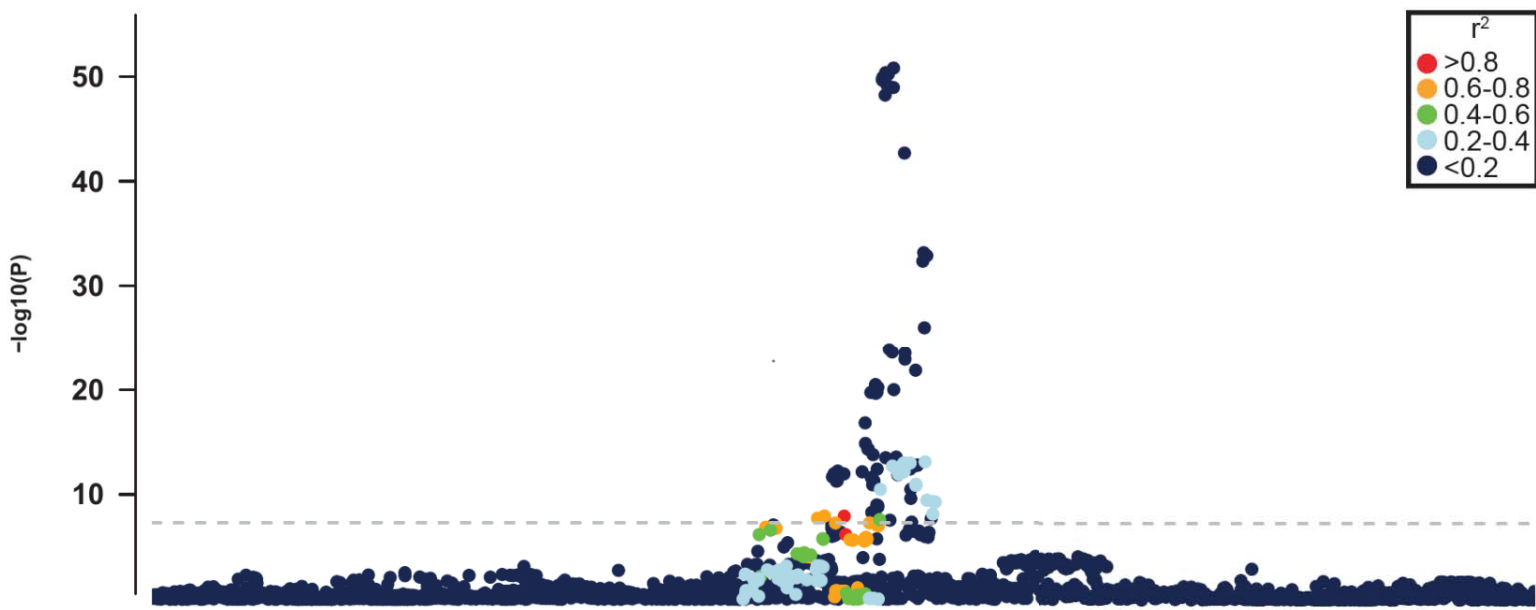


Zscore of the effect allele: C



Postcentral SA: rs3862145

Chromosome 15



C15orf54

THBS1

F

GZ>CP

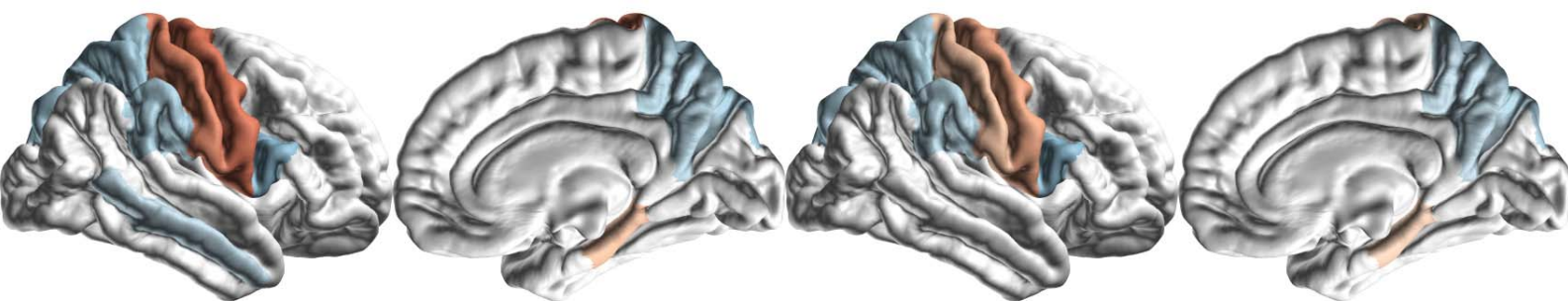


CP>GZ

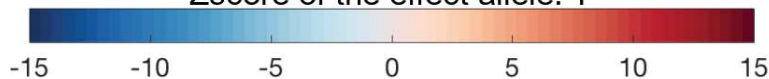


Corrected for total SA

Uncorrected

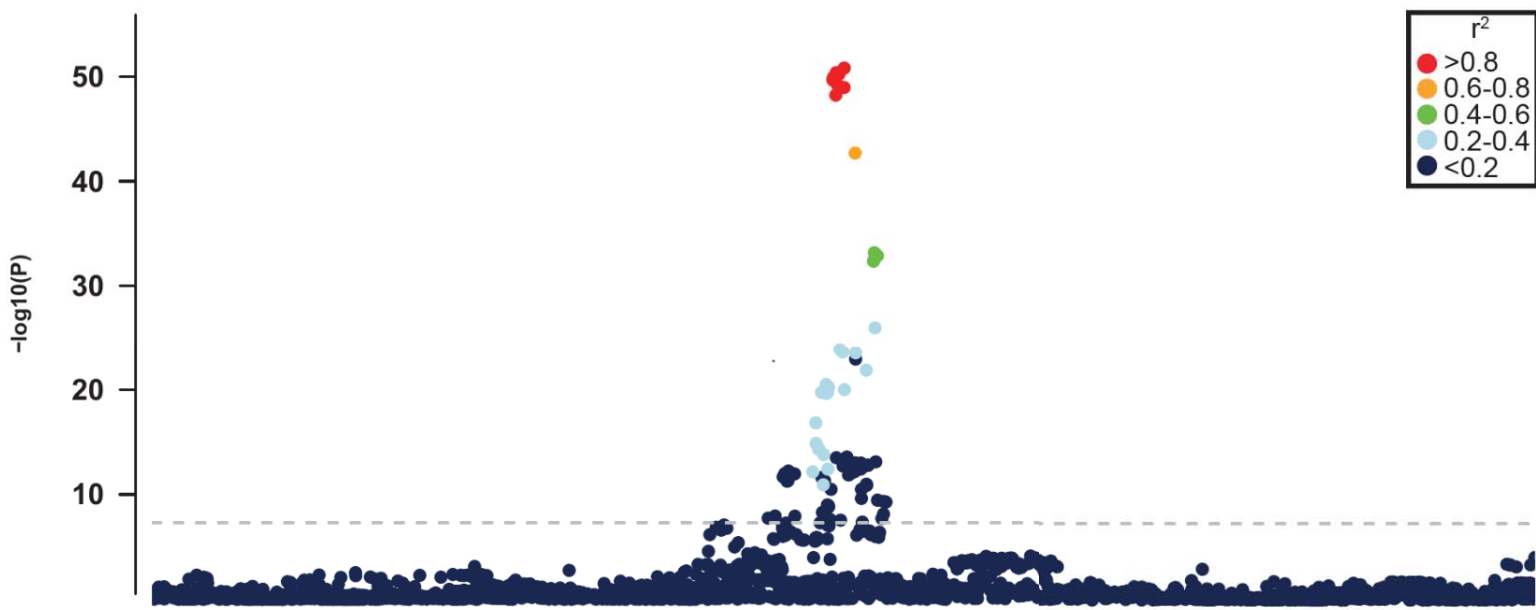
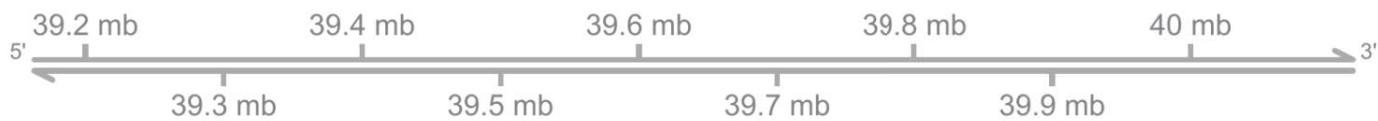


Zscore of the effect allele: T



Postcentral SA: rs4924346

Chromosome 15



C15orf54

THBS1



FSIF

GZ>CP

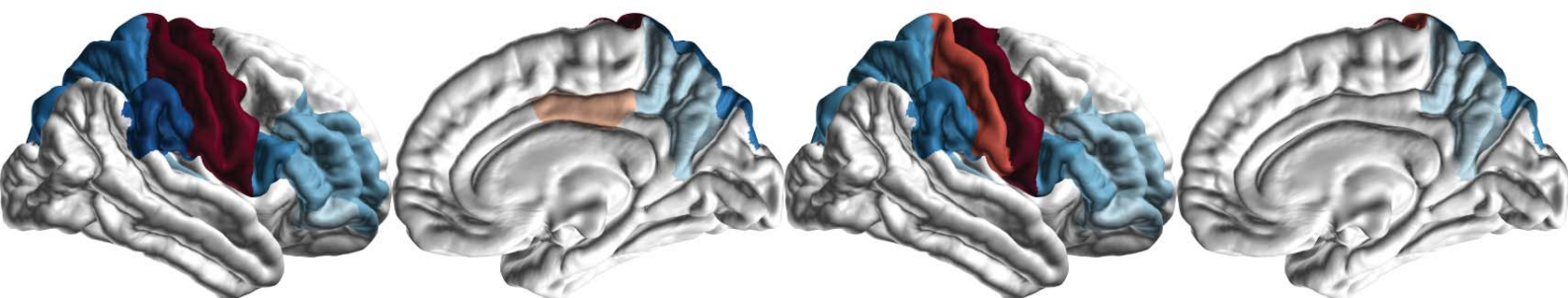


CP>GZ

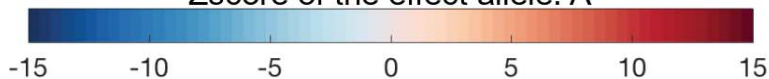


Corrected for total SA

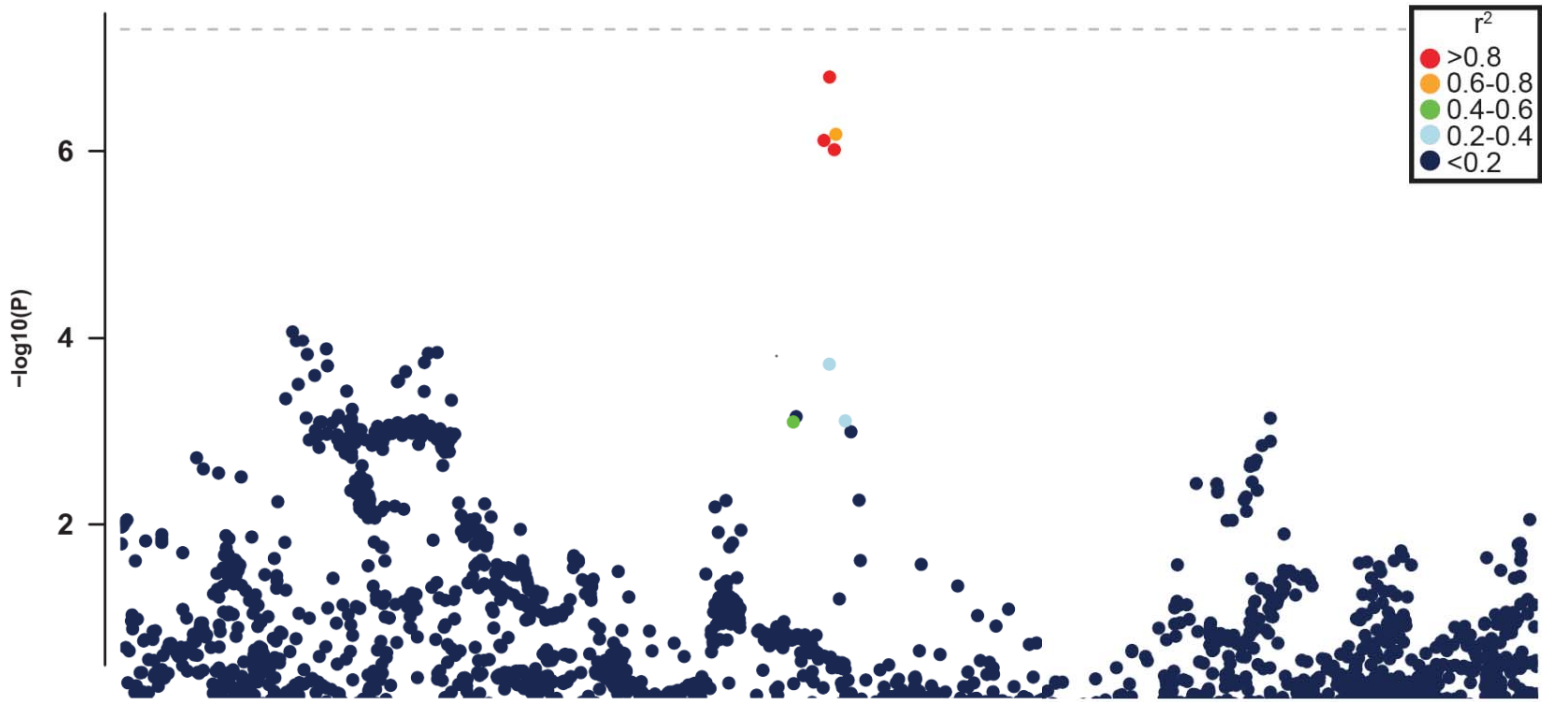
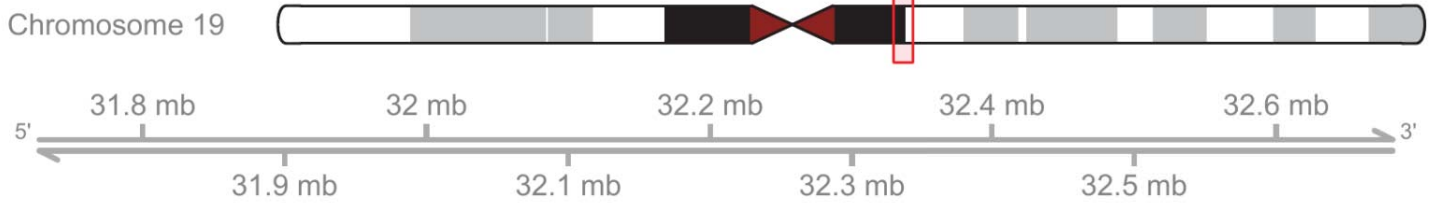
Uncorrected



Zscore of the effect allele: A



Postcentral SA: rs117623407



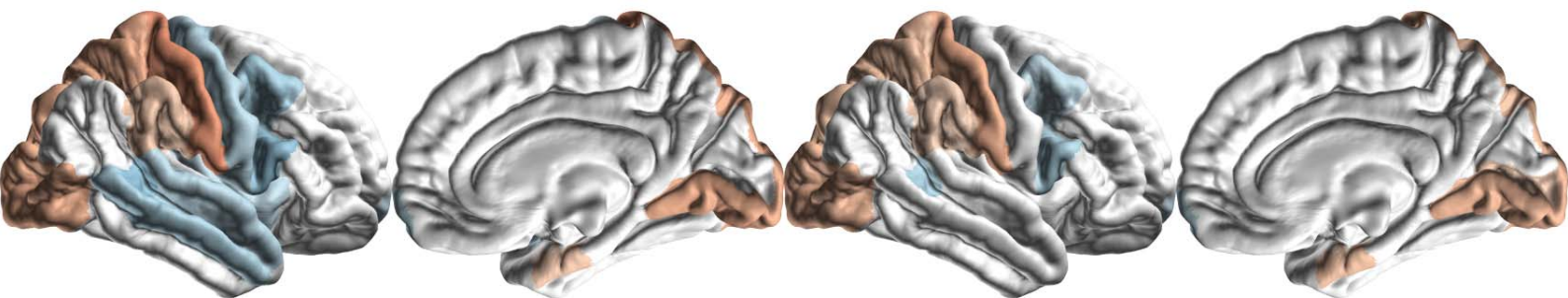
TSHZ3

GZ>CP

CP>GZ

Corrected for total SA

Uncorrected

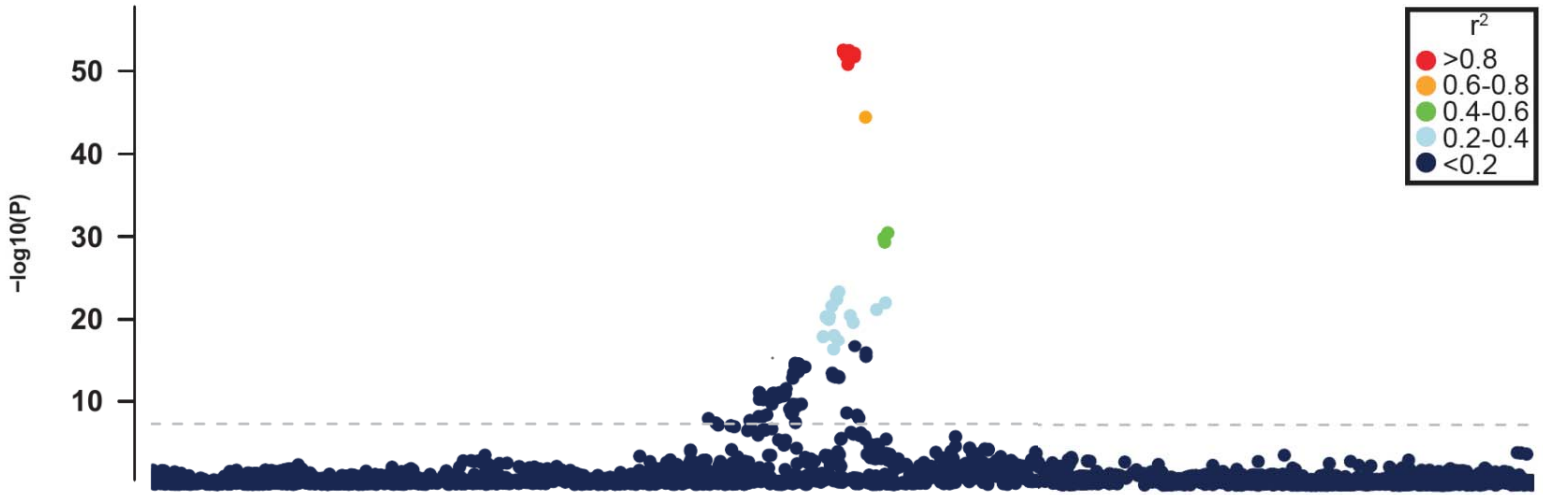
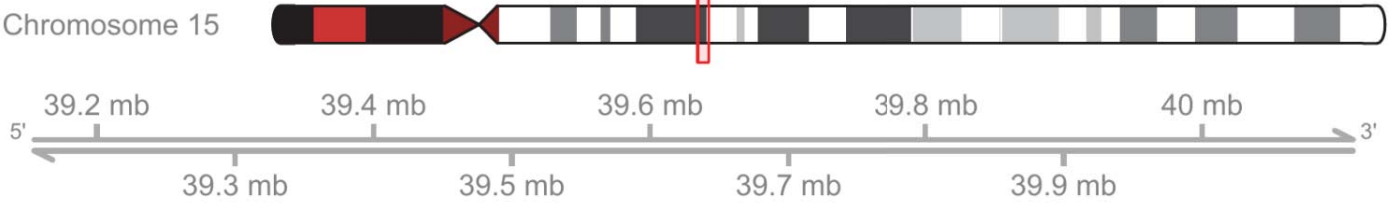


Zscore of the effect allele: A



Postcentral TH: rs11639391

Chromosome 15



C15orf54

THBS1

FSI

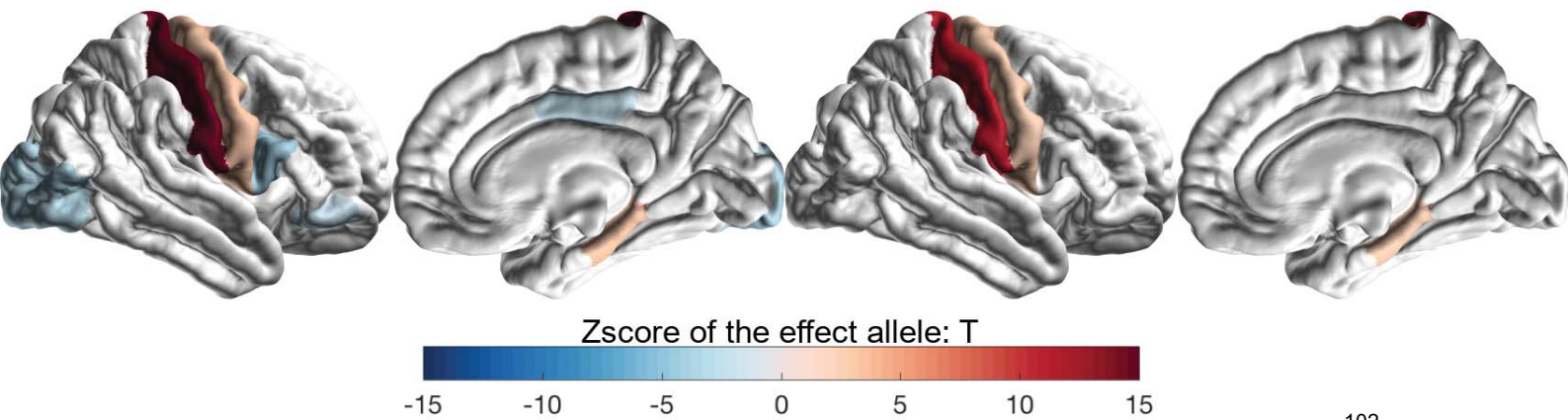
FSI

Angular
Temporal
dIPFC
Cingulate



Corrected for average TH

Uncorrected

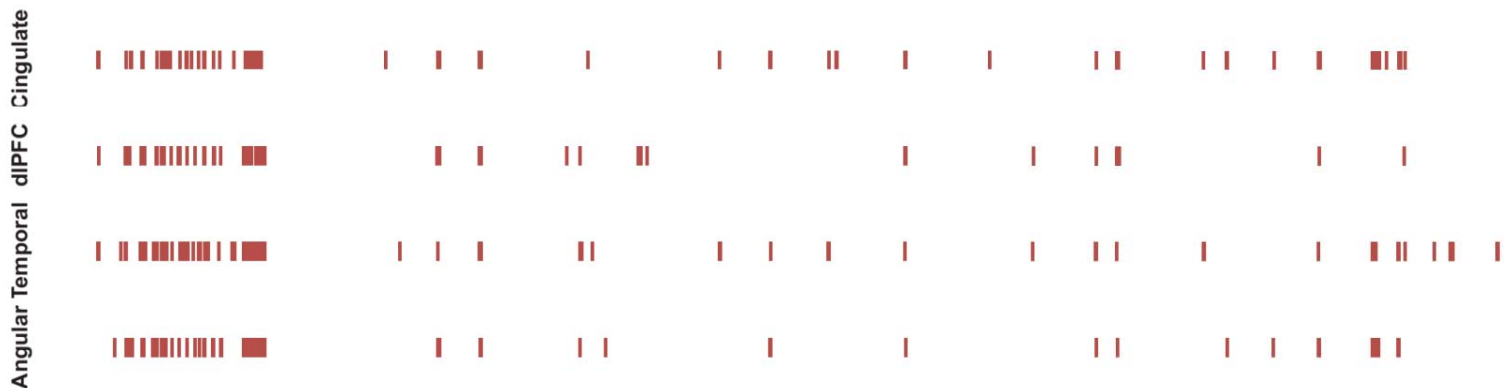
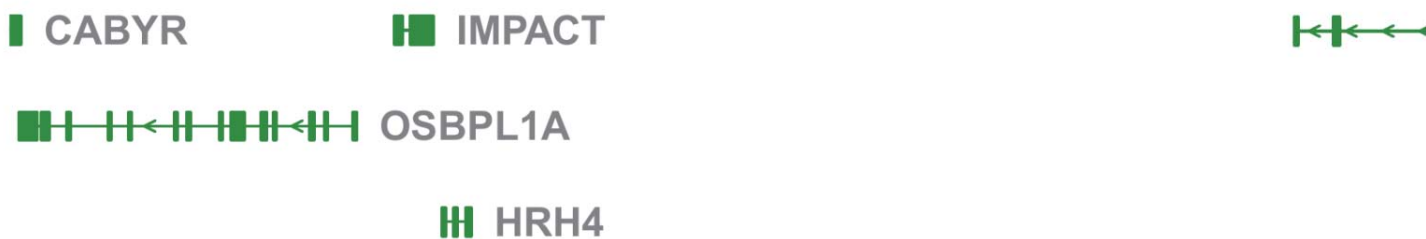
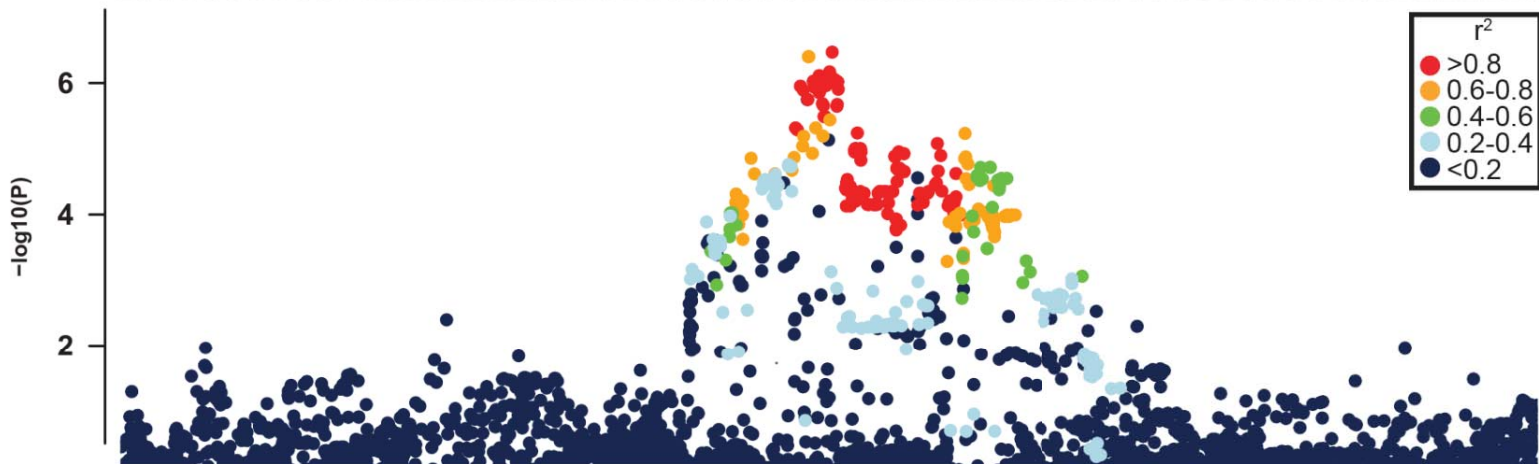
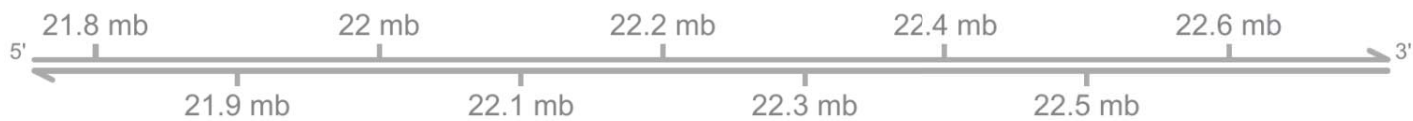


Zscore of the effect allele: T



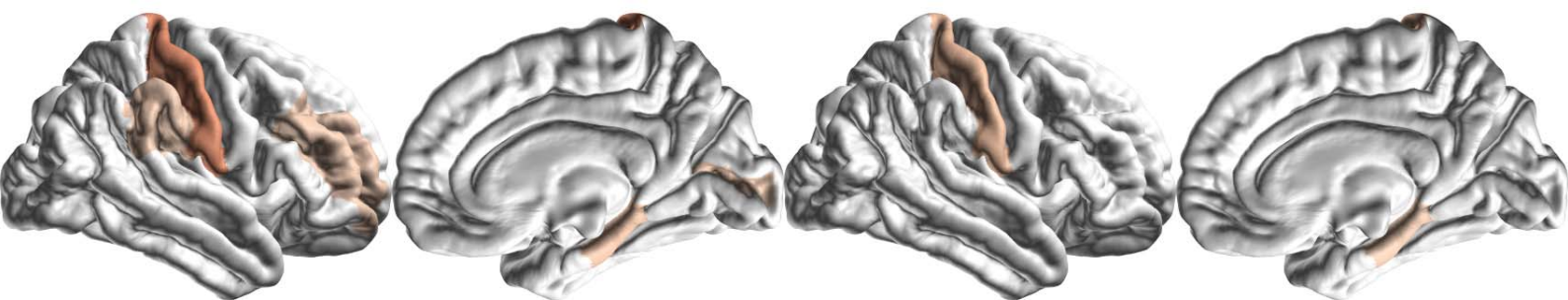
Postcentral TH: rs12958970

Chromosome 18

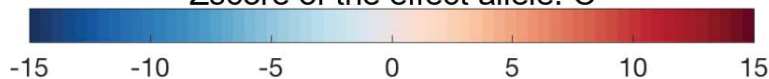


Corrected for average TH

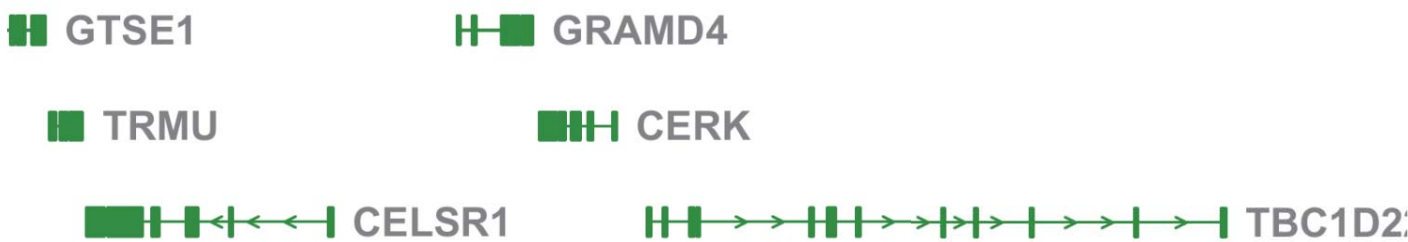
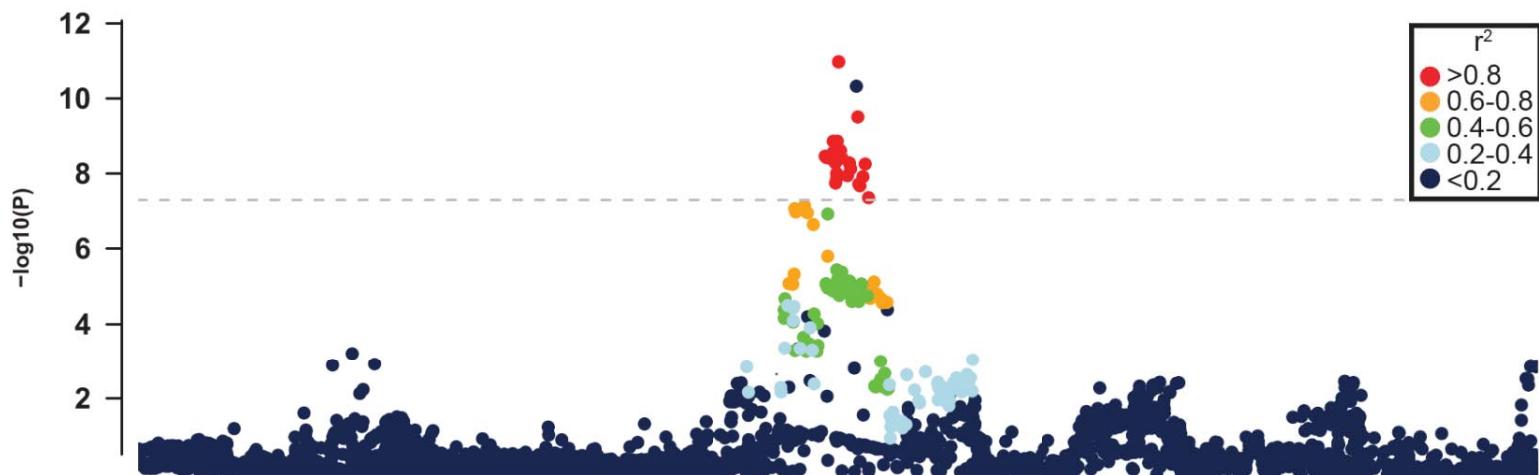
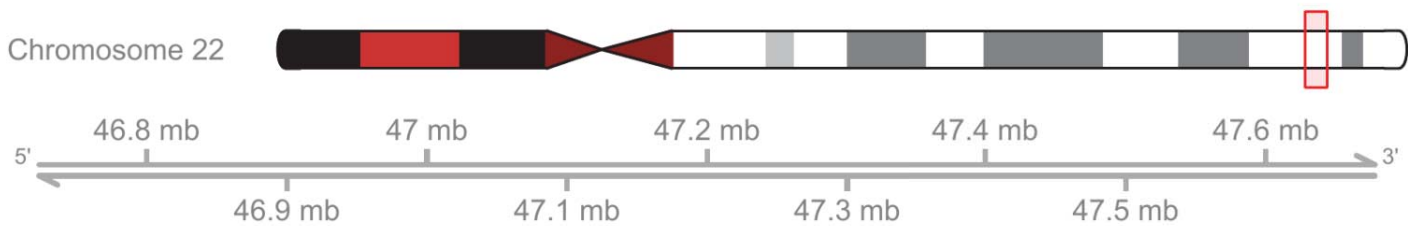
Uncorrected



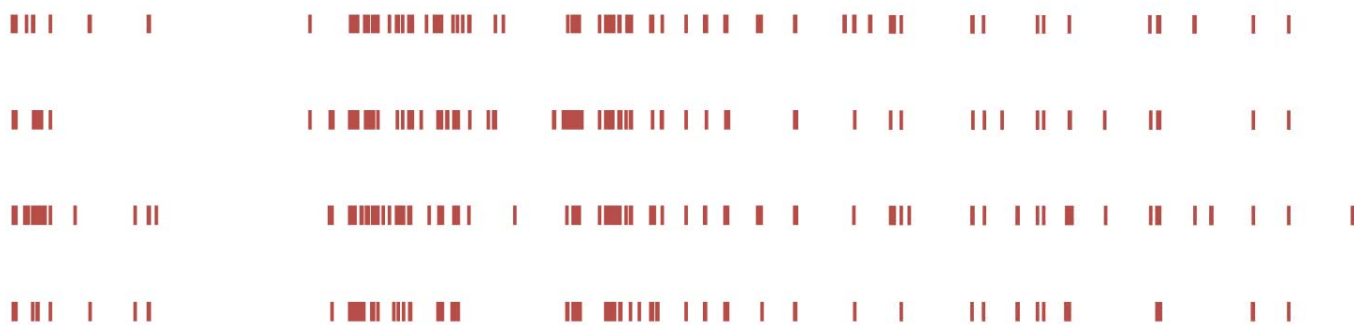
Zscore of the effect allele: C



Postcentral TH: rs4823877

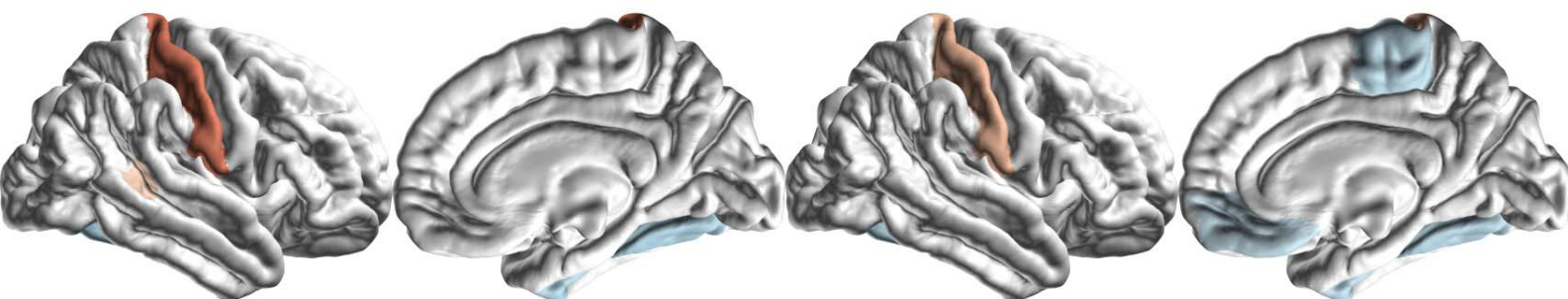


Angular Temporal dIPFC Cingulate

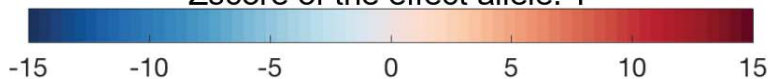


Corrected for average TH

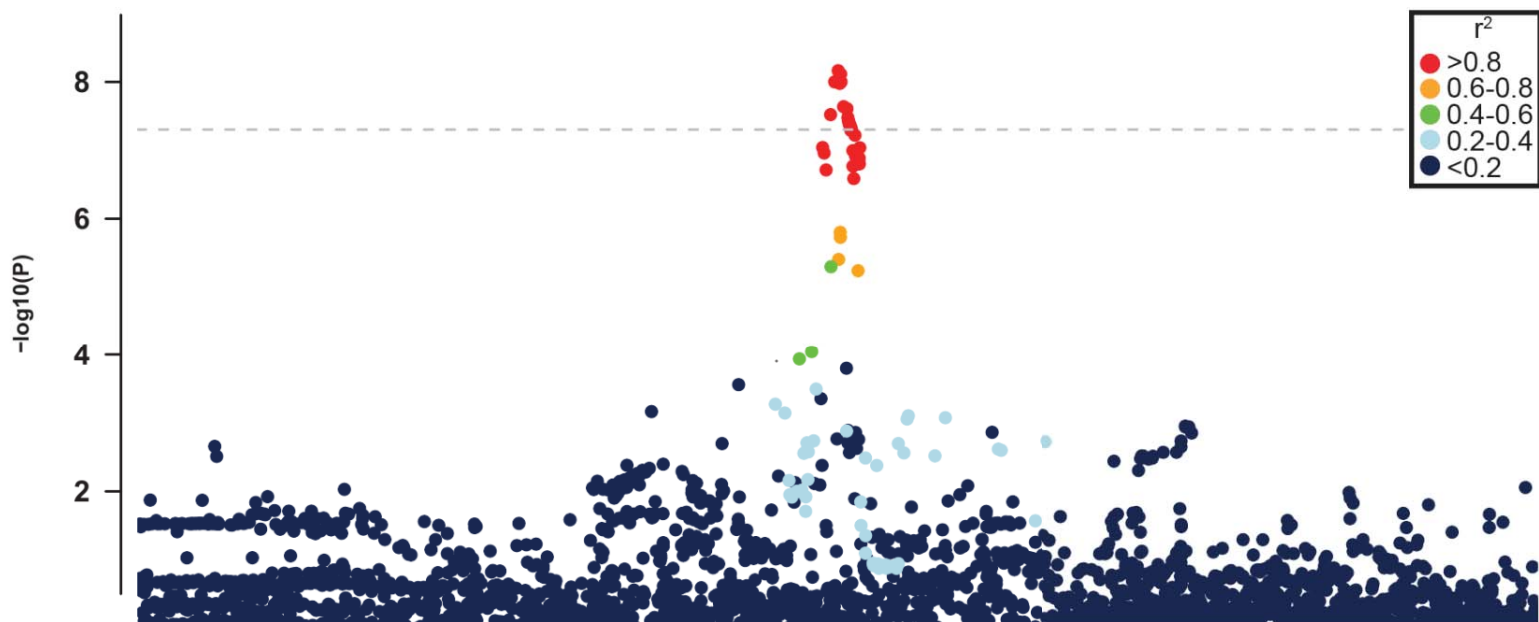
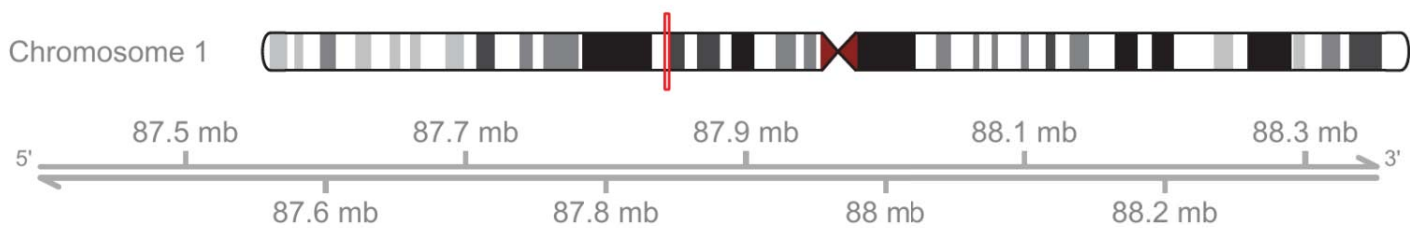
Uncorrected



Zscore of the effect allele: T



Precuneus SA: rs59373415



SEP15

LMO4

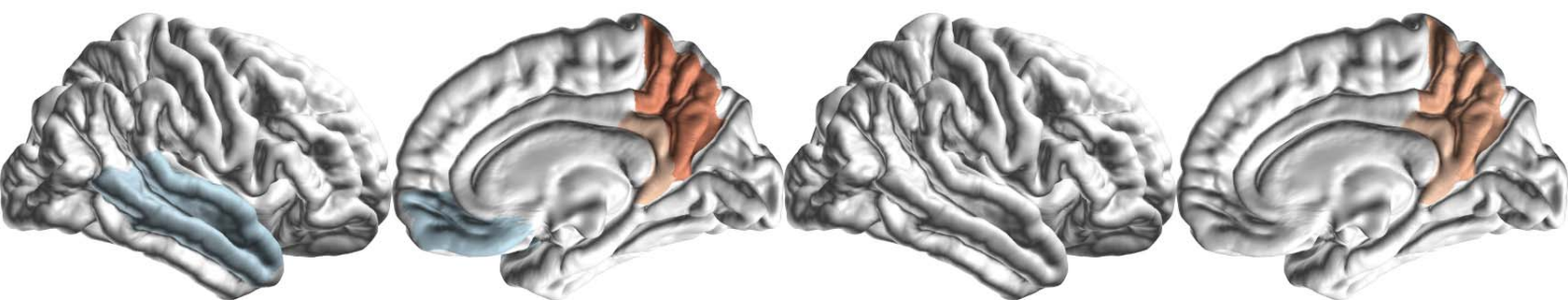
HS2ST1

GZ>CP

CP>GZ

Corrected for total SA

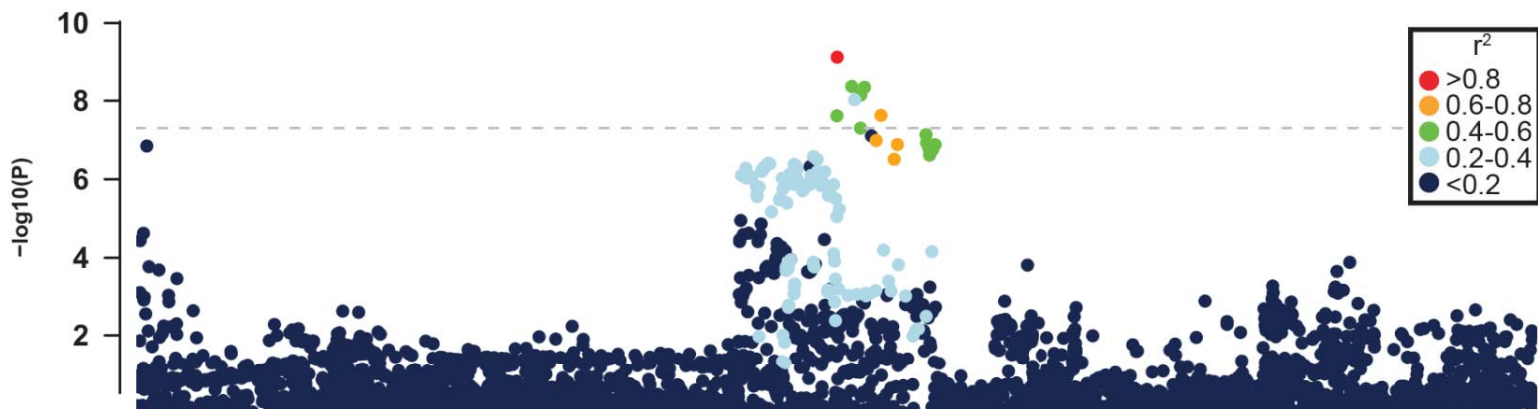
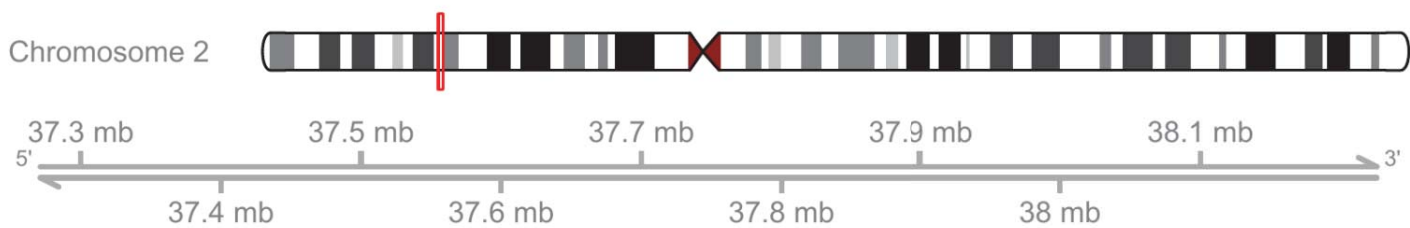
Uncorrected



Zscore of the effect allele: C

-15 -10 -5 0 5 10 15

Precuneus SA: rs7559976



HEATR5B QPCT CDC42EP3

GPATCH11

EIF2AK2

SULT6B1

CEBPZ

NDUFAF7

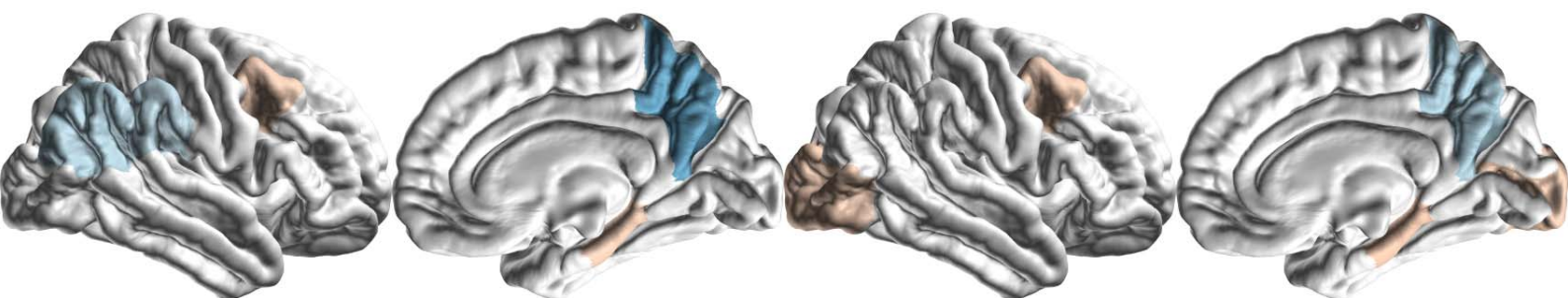
PRKD3

CP>GZ GZ>CP



Corrected for total SA

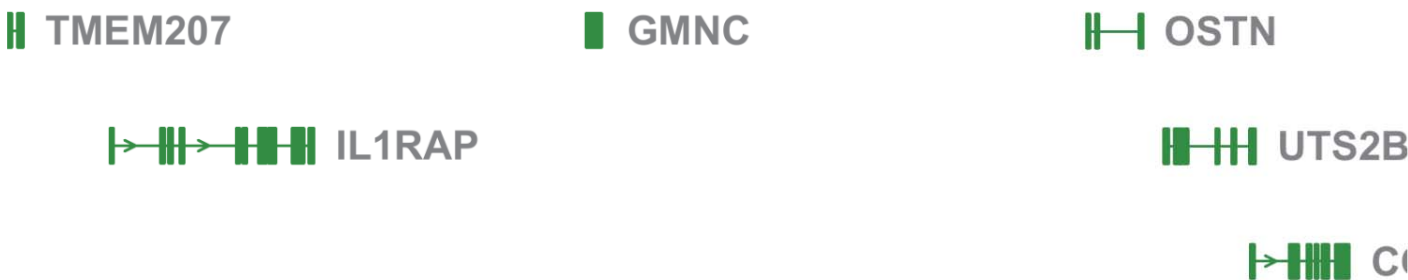
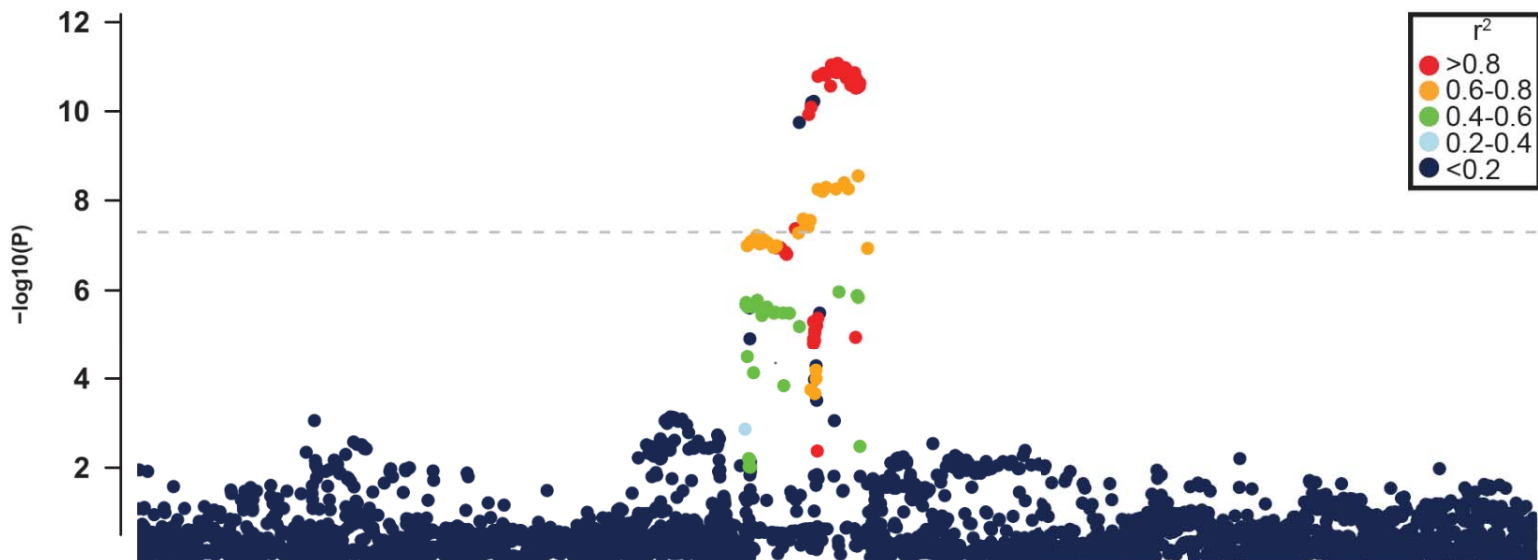
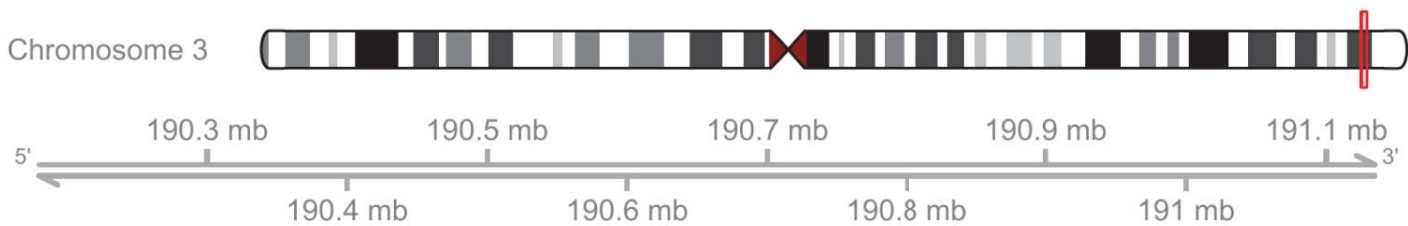
Uncorrected



Zscore of the effect allele: T

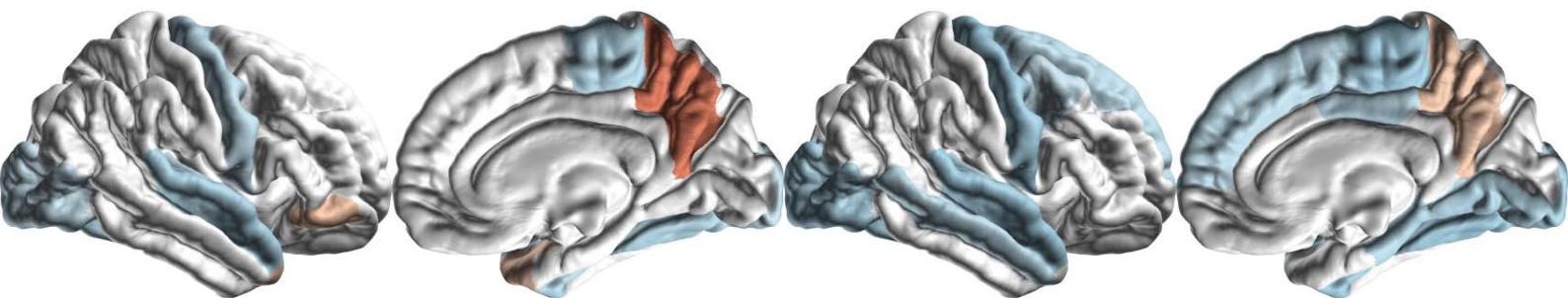


Precuneus SA: rs905124

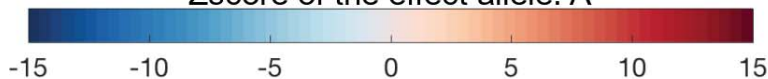


Corrected for total SA

Uncorrected

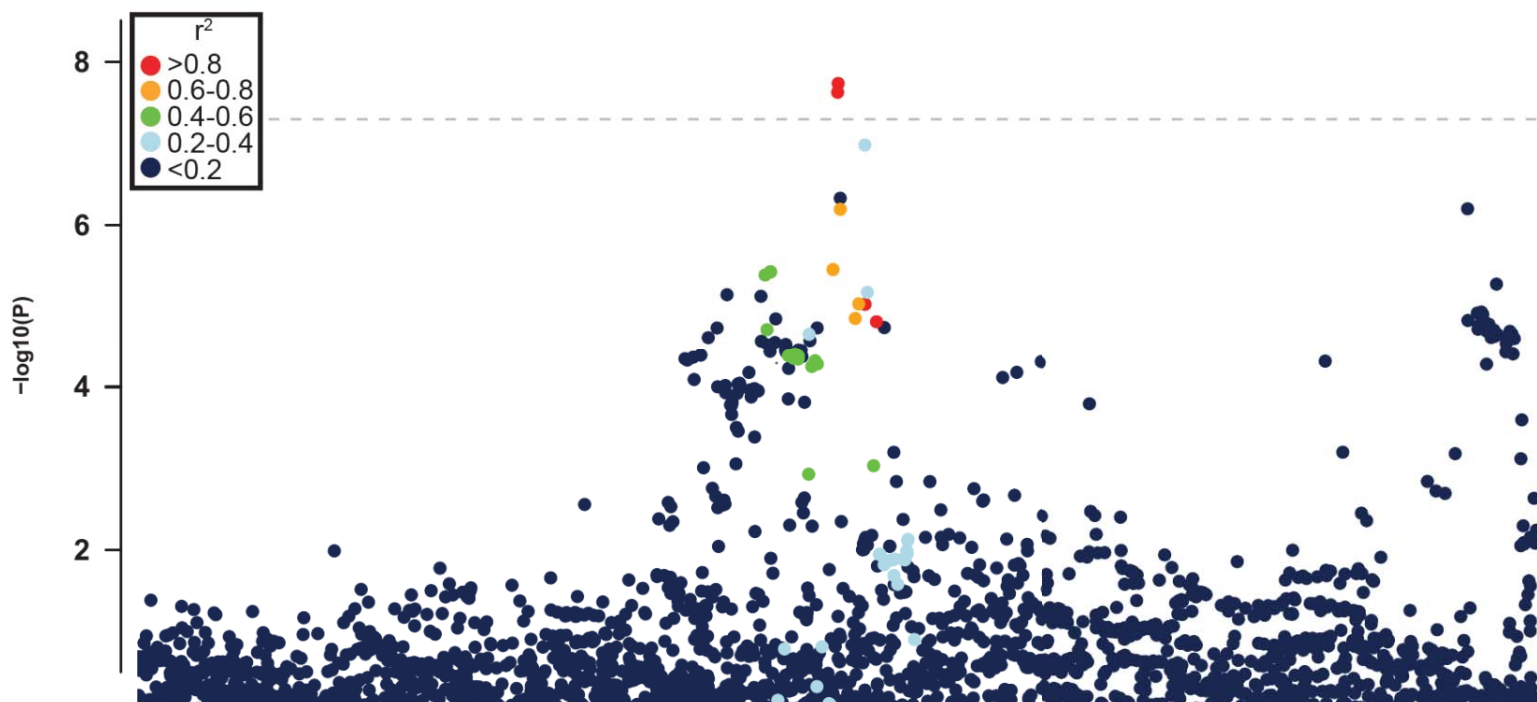
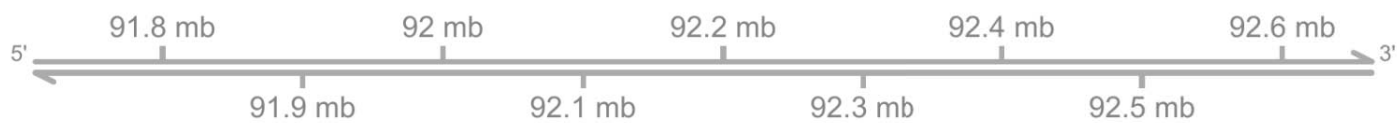


Zscore of the effect allele: A



Precuneus SA: rs888814

Chromosome 5



GZ>CP

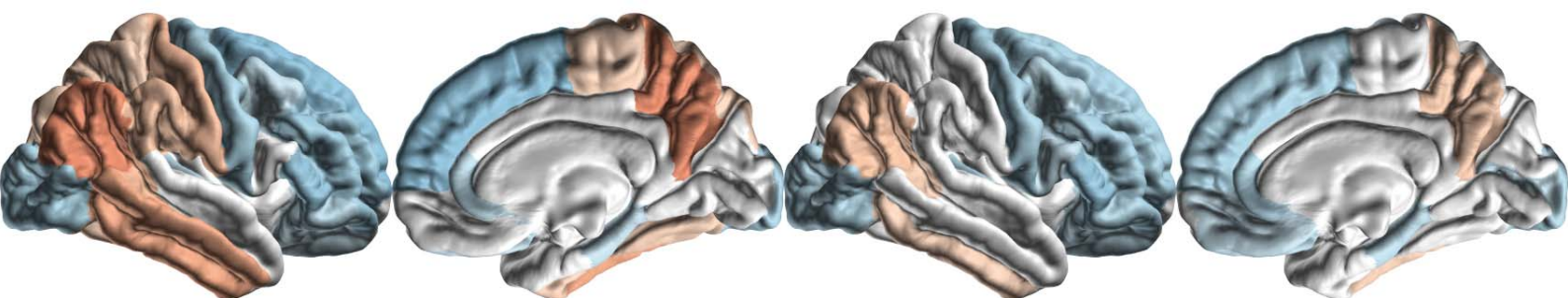


CP>GZ



Corrected for total SA

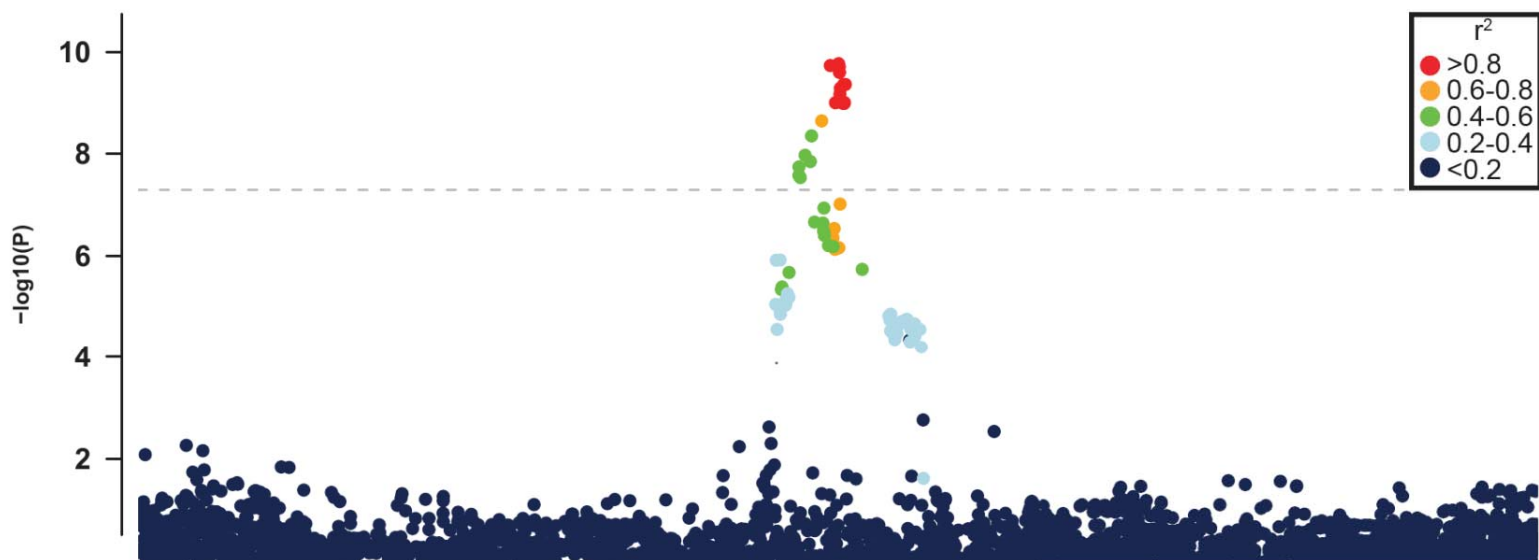
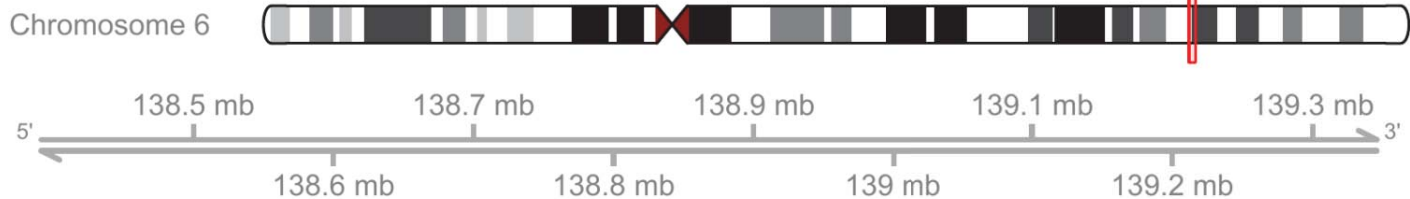
Uncorrected



Zscore of the effect allele: T

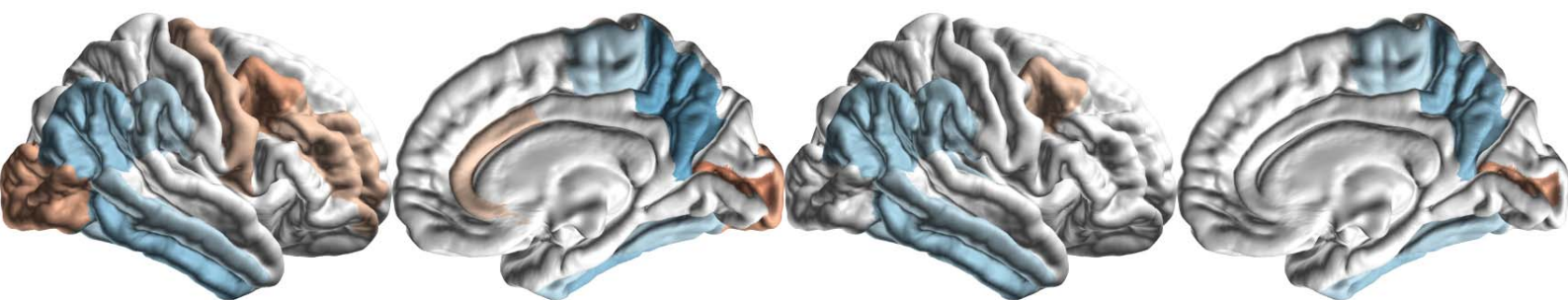


Precuneus SA: rs9399245



Corrected for total SA

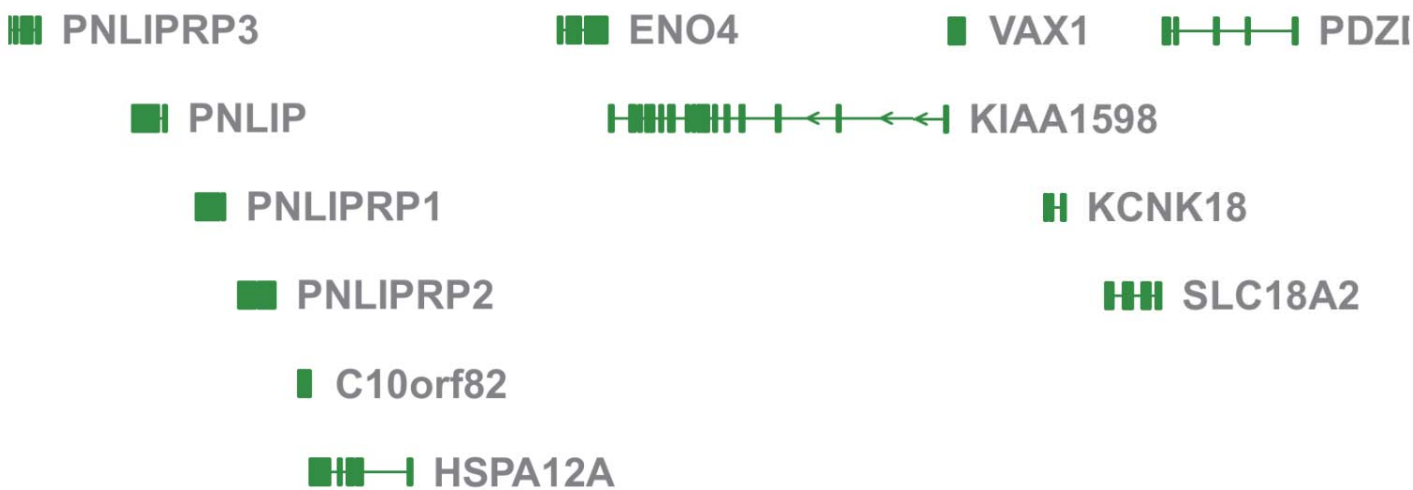
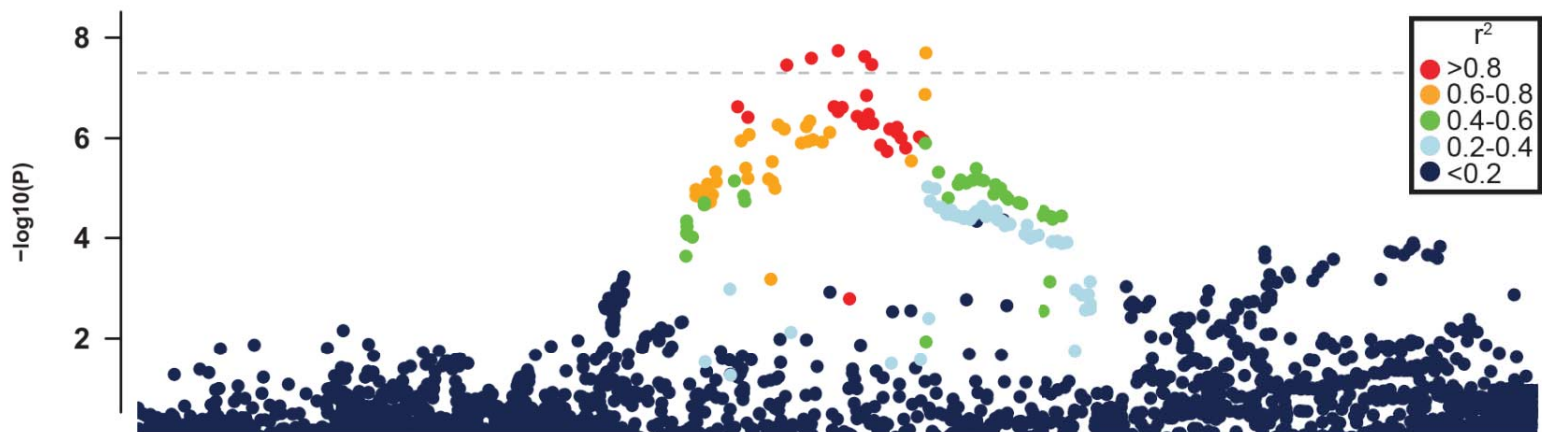
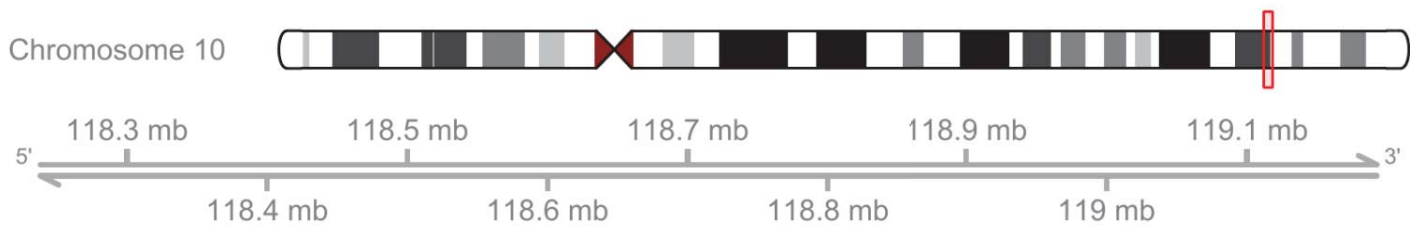
Uncorrected



Zscore of the effect allele: T



Precuneus SA: rs7902527

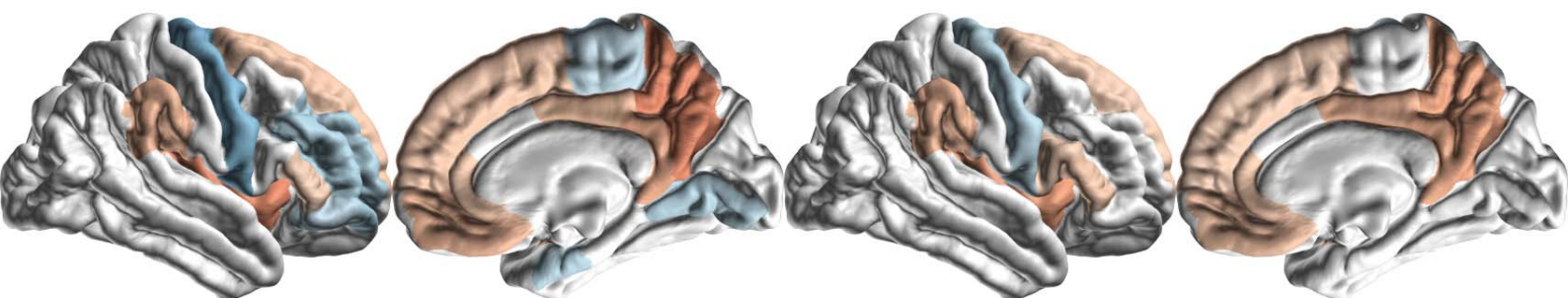


CP>GZ GZ>CP

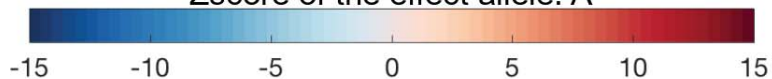


Corrected for total SA

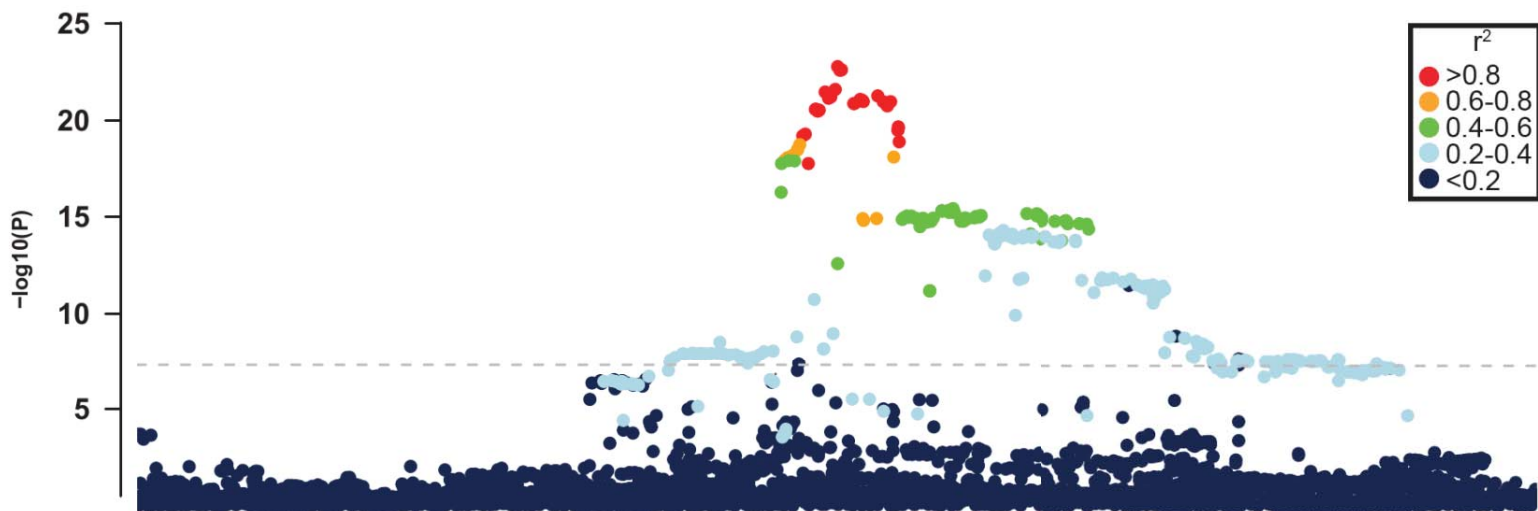
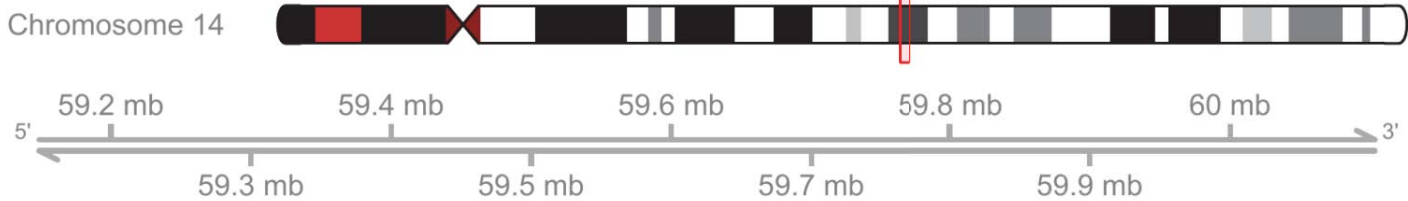
Uncorrected



Zscore of the effect allele: A



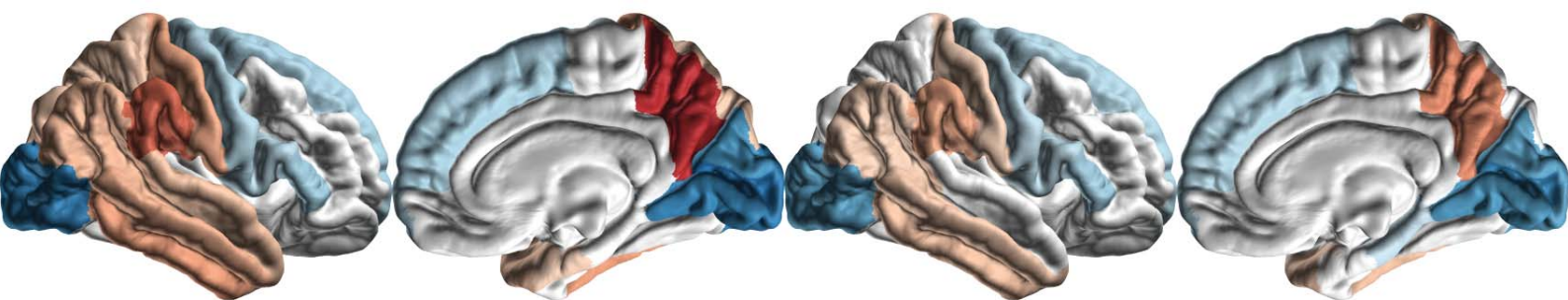
Precuneus SA: rs73313052



CP>GZ
GZ>CP

Corrected for total SA

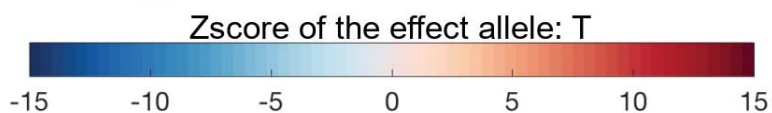
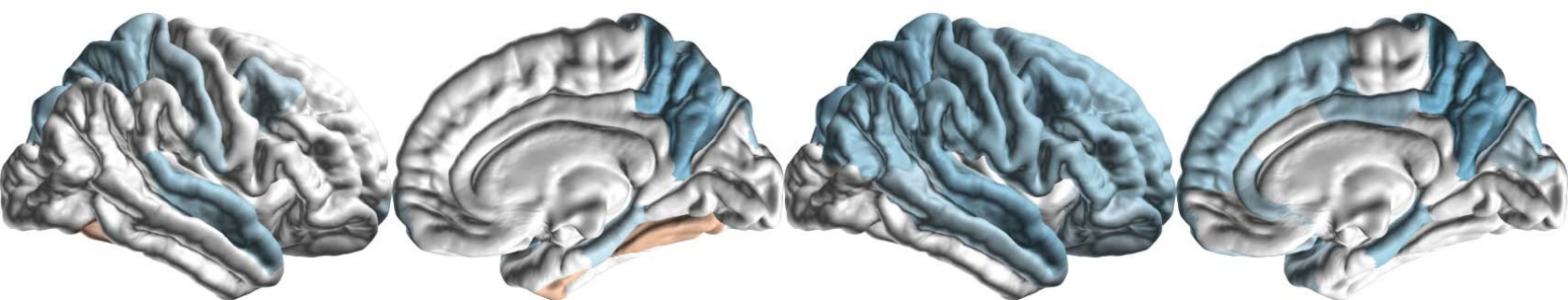
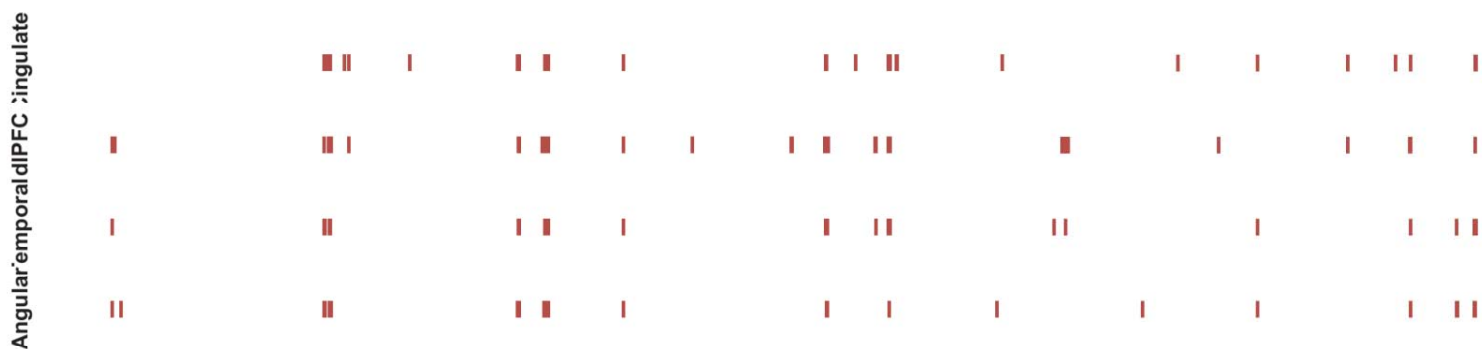
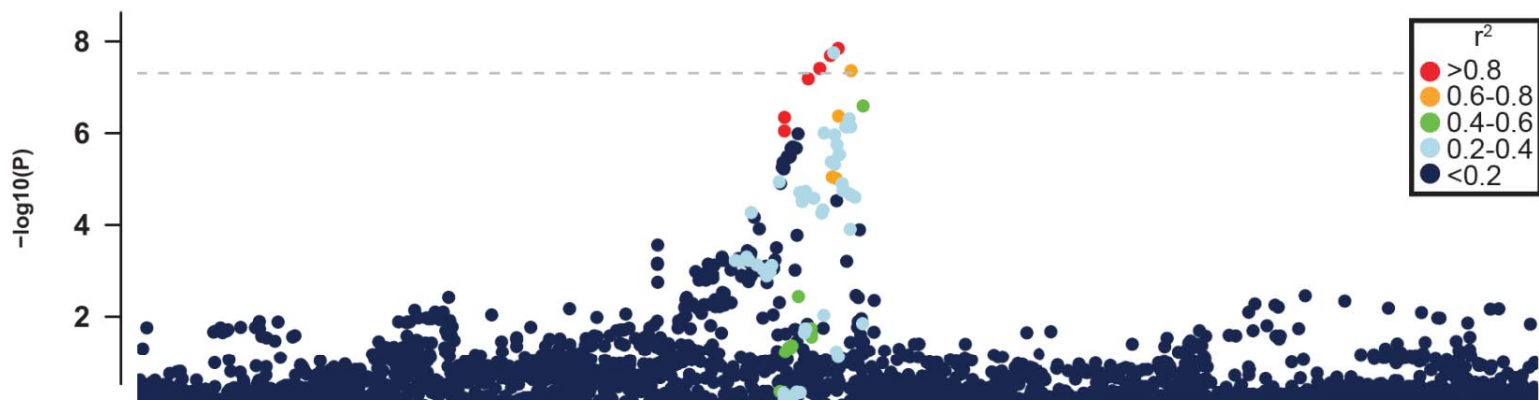
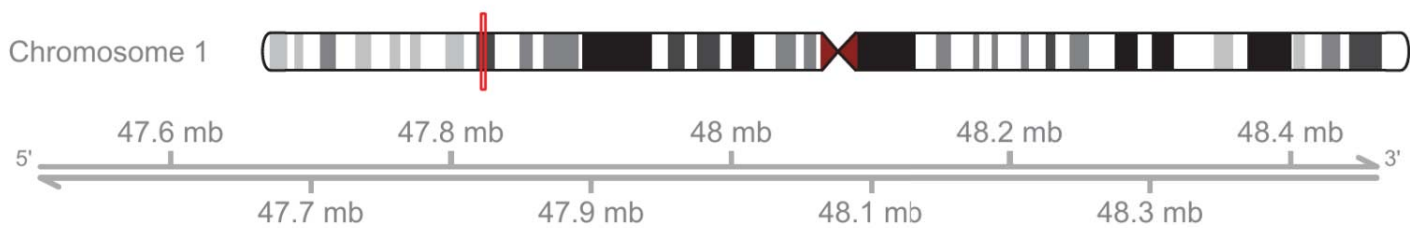
Uncorrected



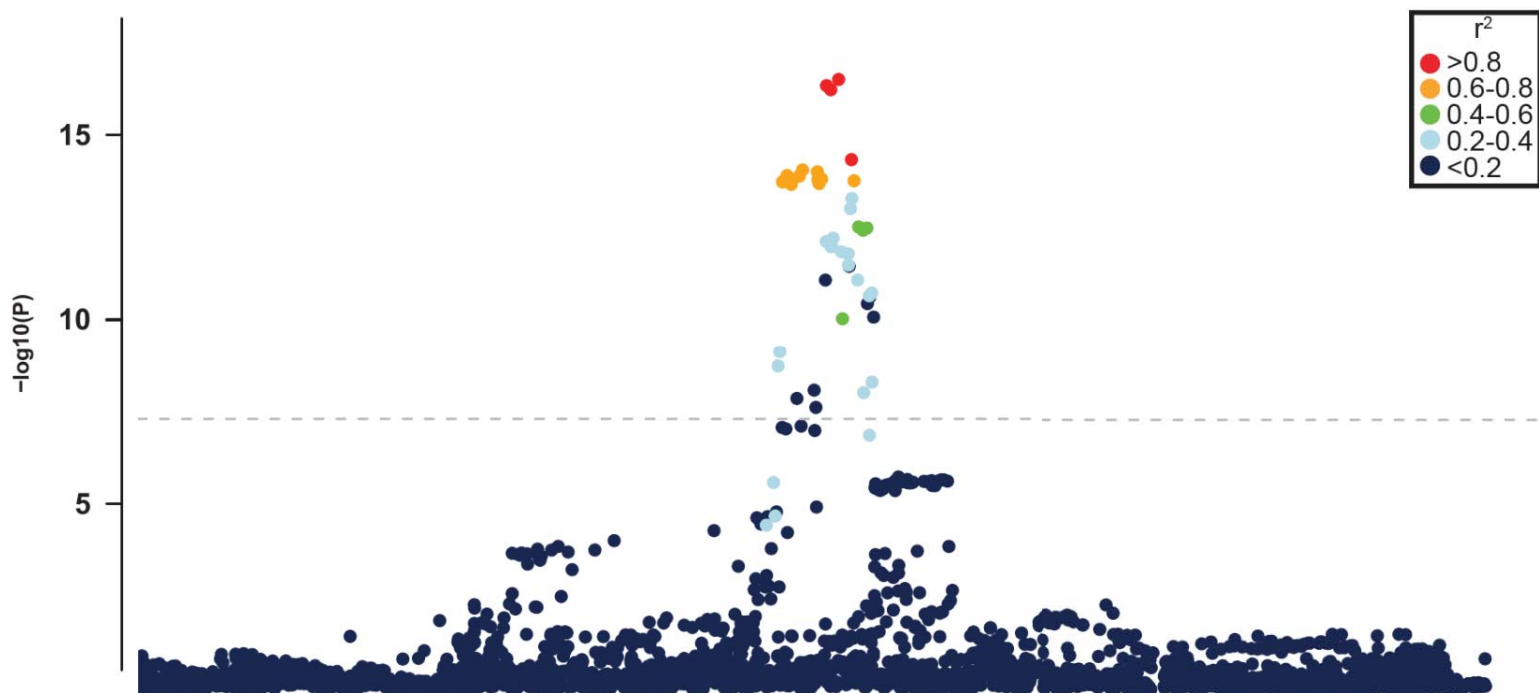
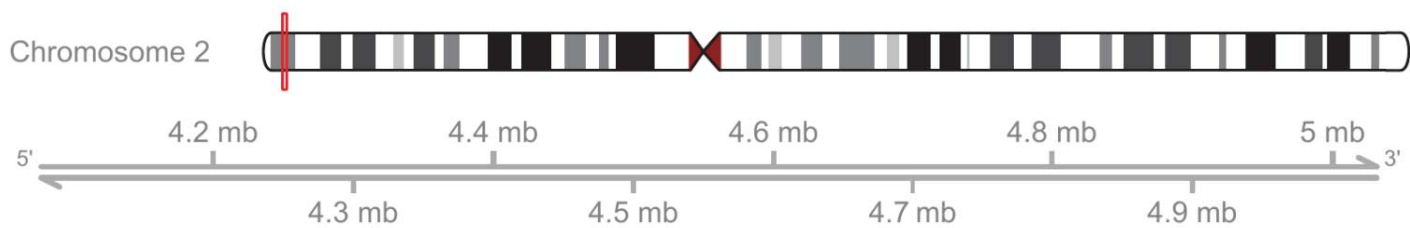
Zscore of the effect allele: A



Precuneus TH: rs12118339



Superior Parietal SA: rs639016

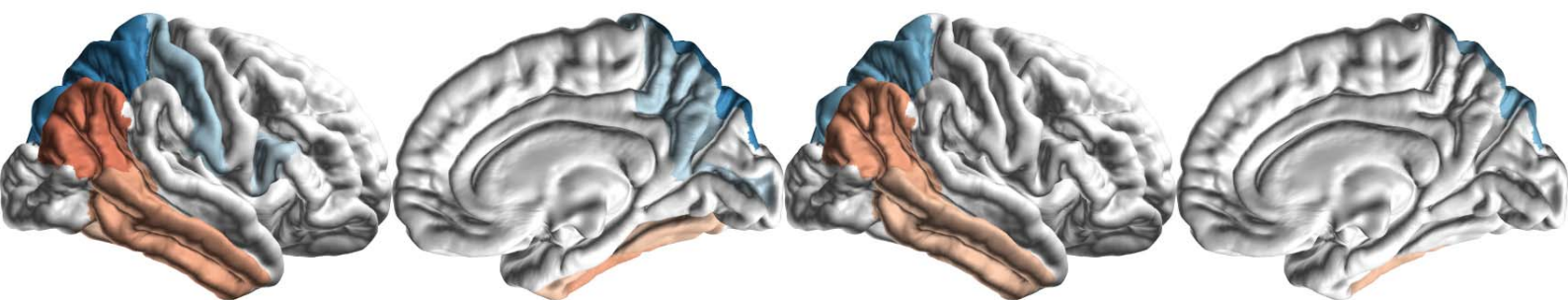


GZ>CP

CP>GZ

Corrected for total SA

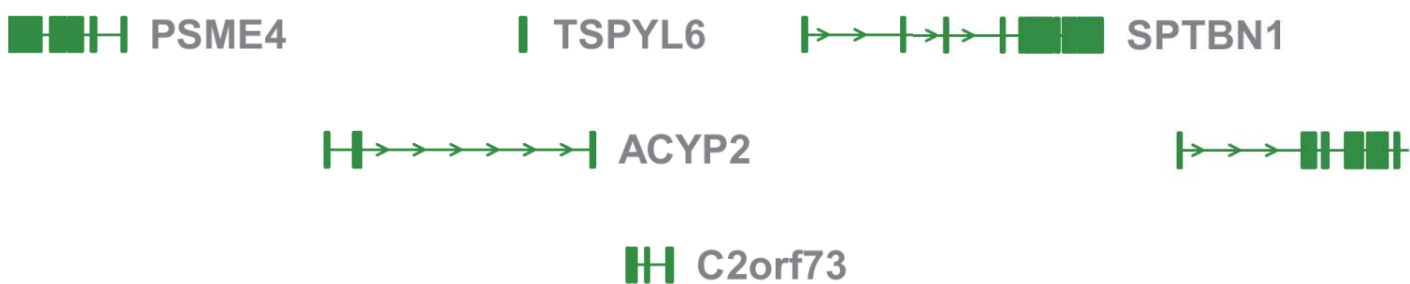
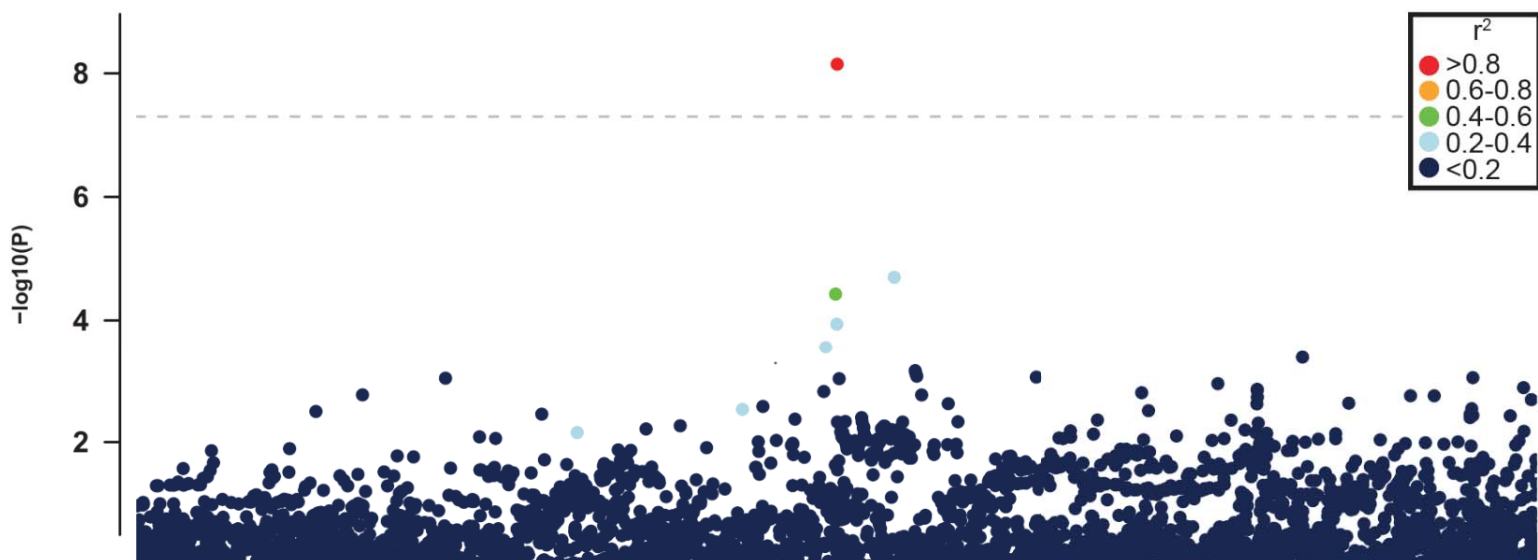
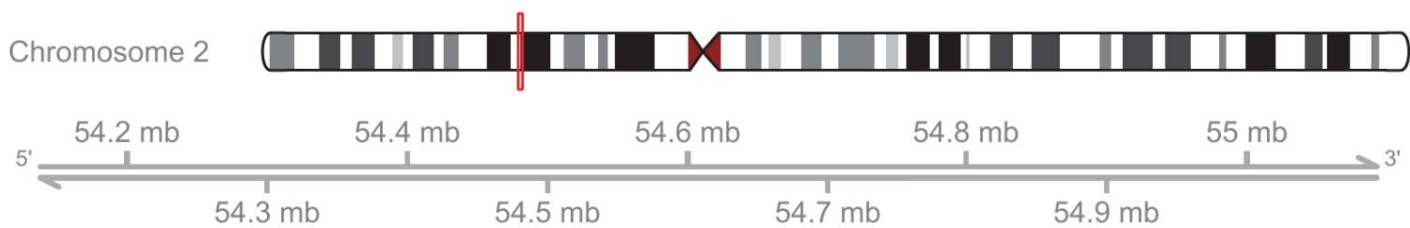
Uncorrected



Zscore of the effect allele: C

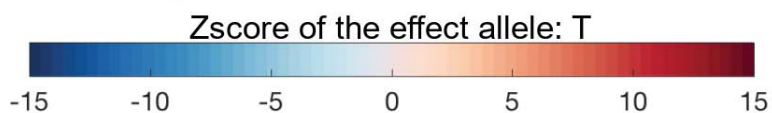
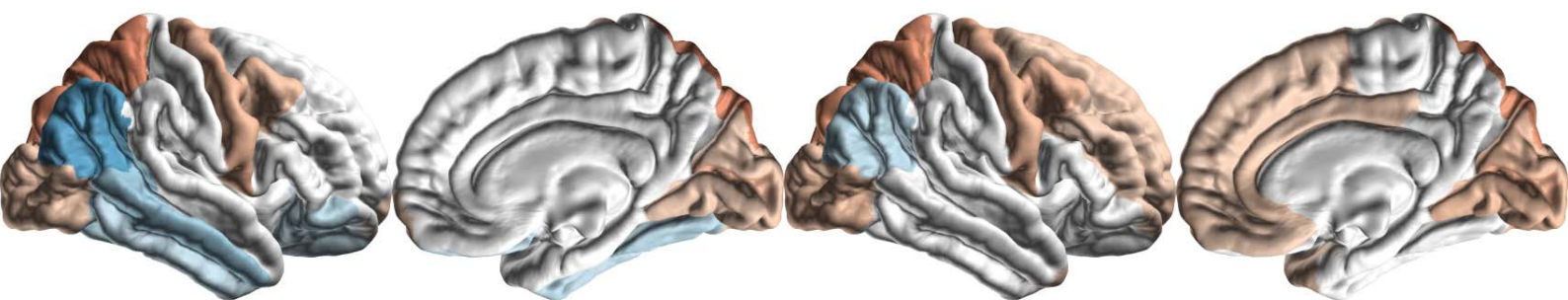
-15 -10 -5 0 5 10 15

Superior Parietal SA: rs79272390

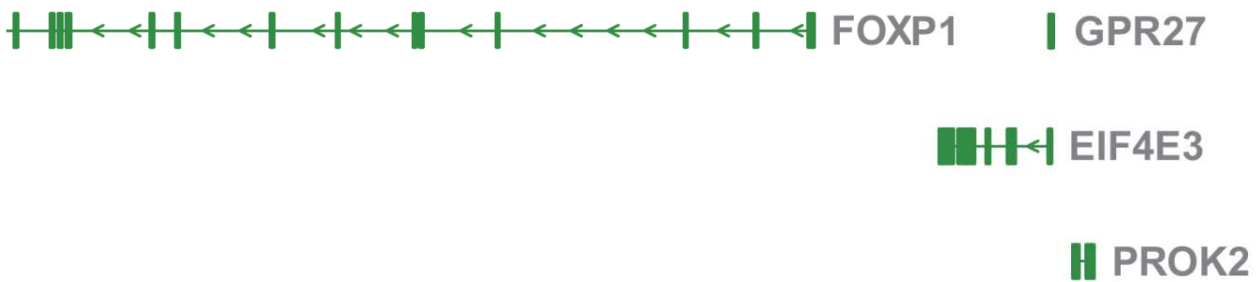
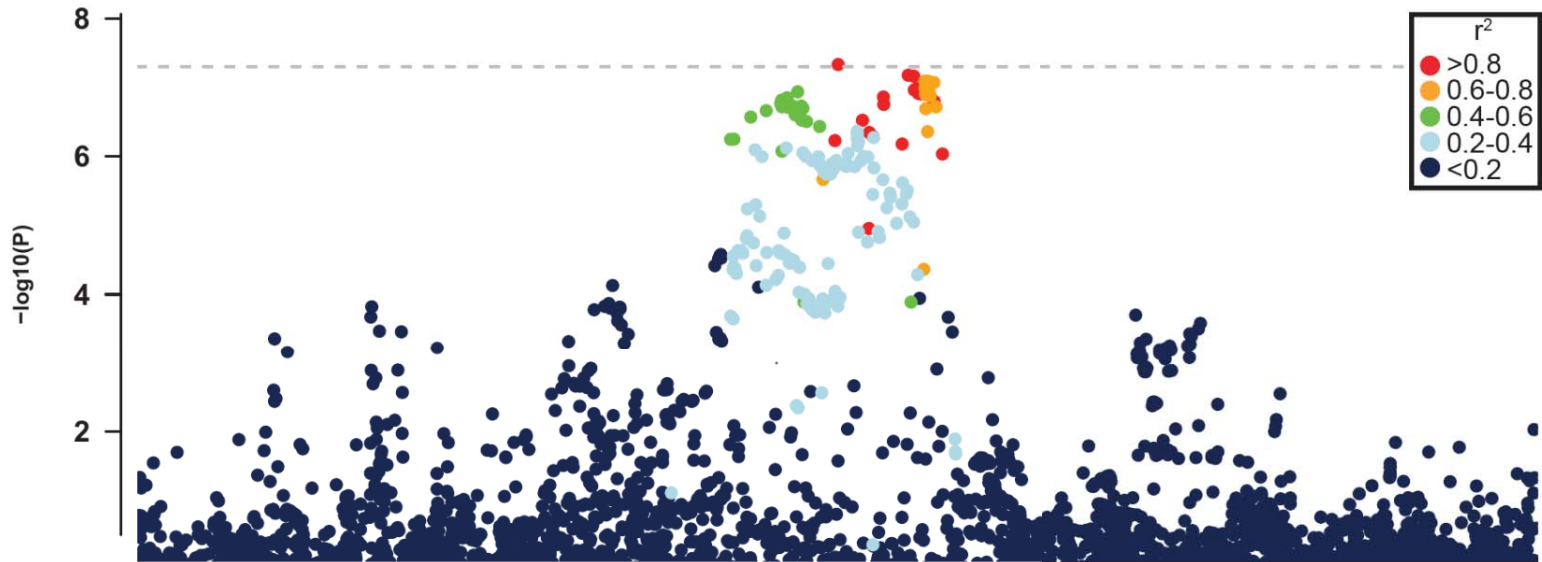
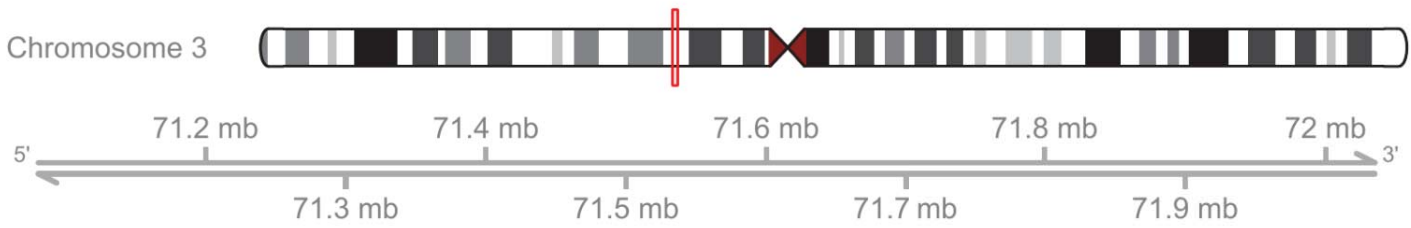


Corrected for total SA

Uncorrected

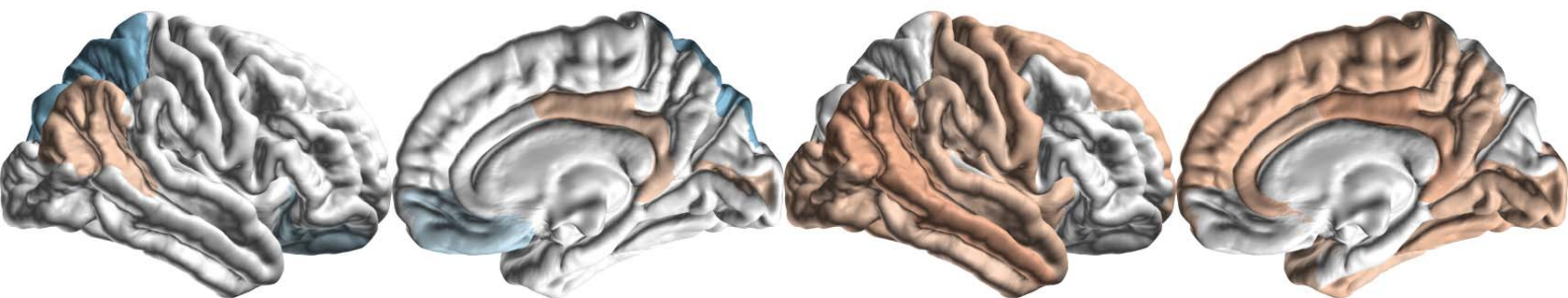


Superior Parietal SA: rs17718831



Corrected for total SA

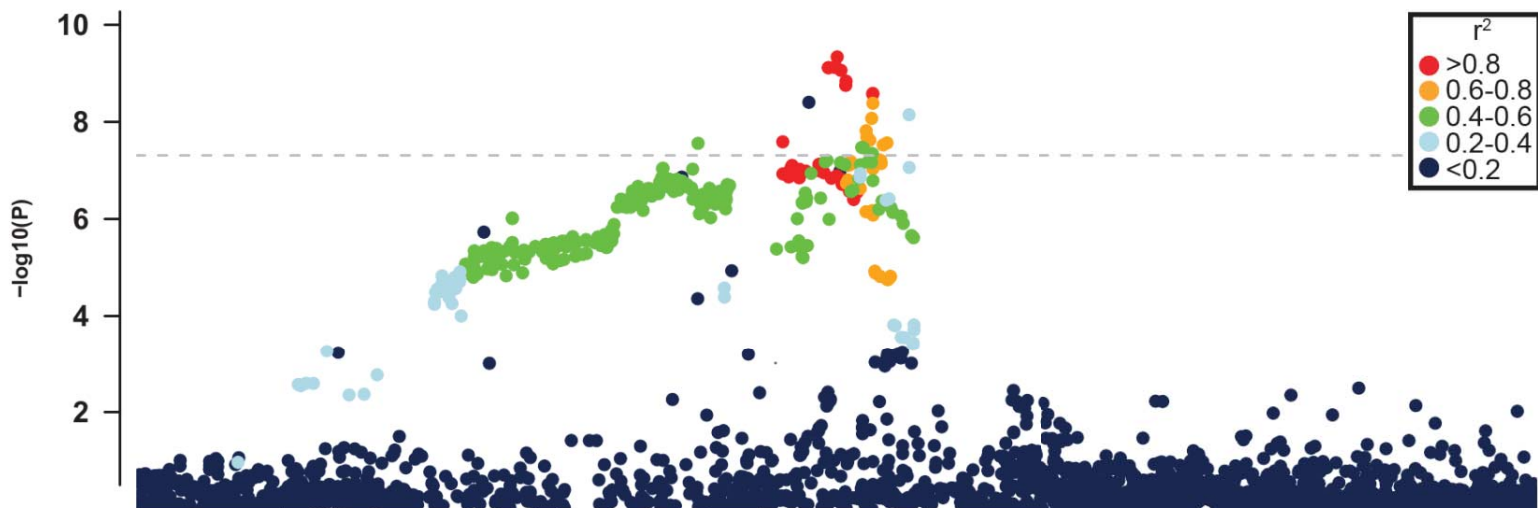
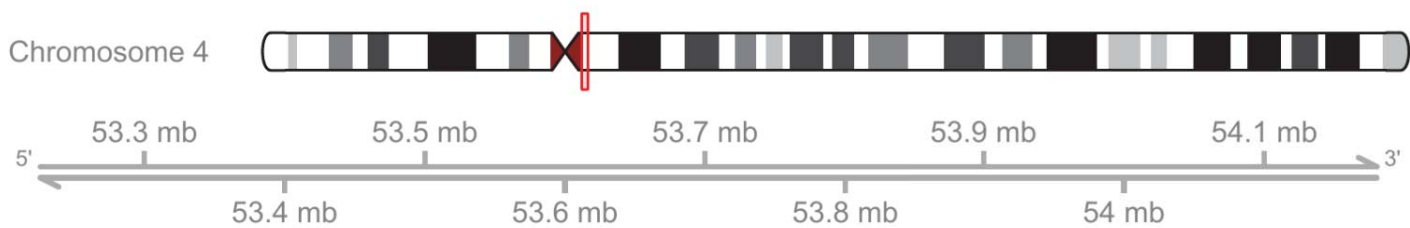
Uncorrected



Zscore of the effect allele: A



Superior Parietal SA: rs6554054



USP46 RASL11B

SNORA26

ERVMER34-1

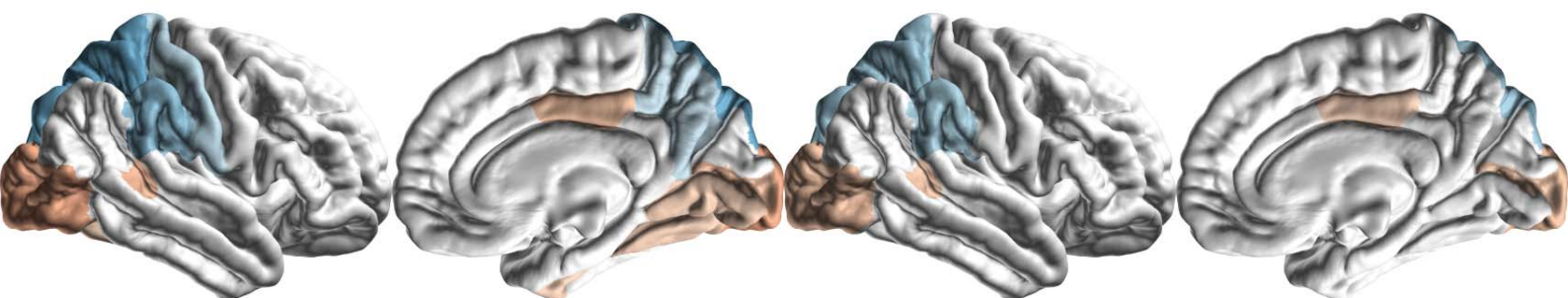


CP>GZ GZ>CP

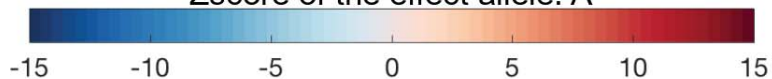


Corrected for total SA

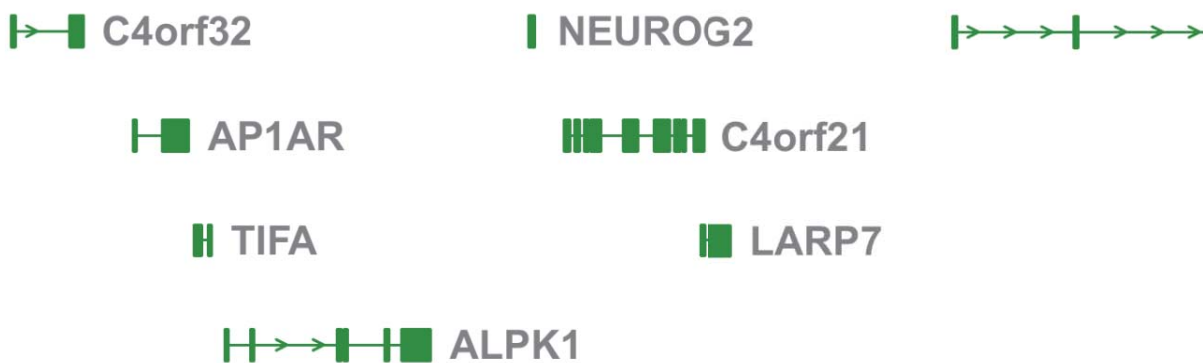
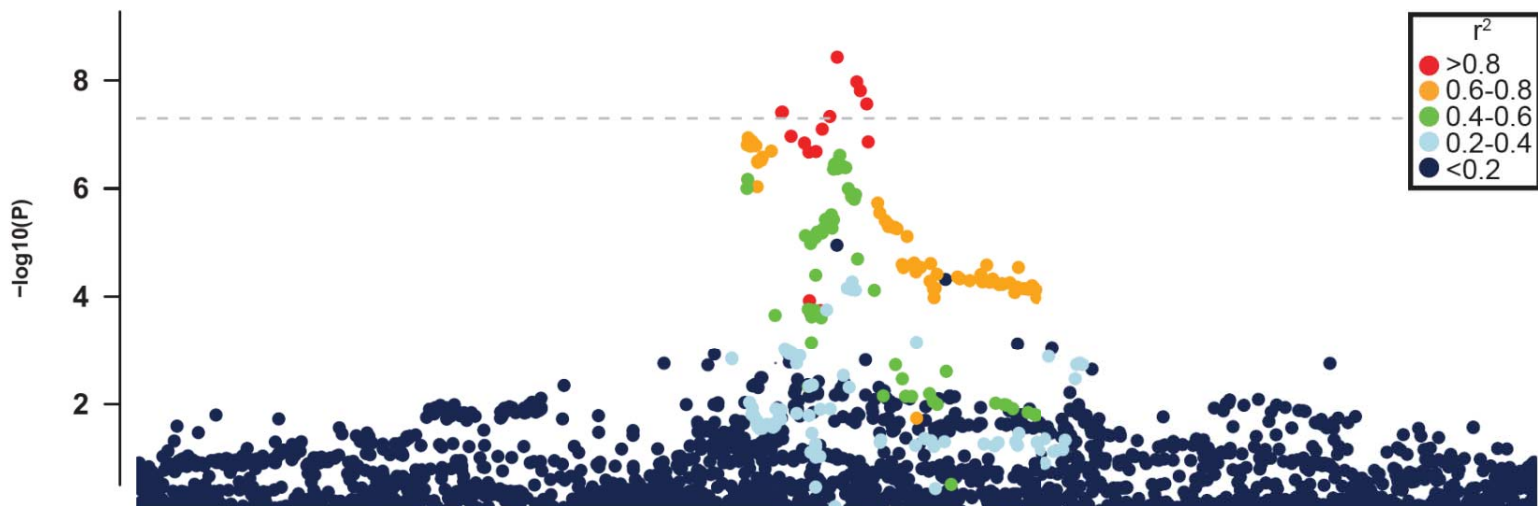
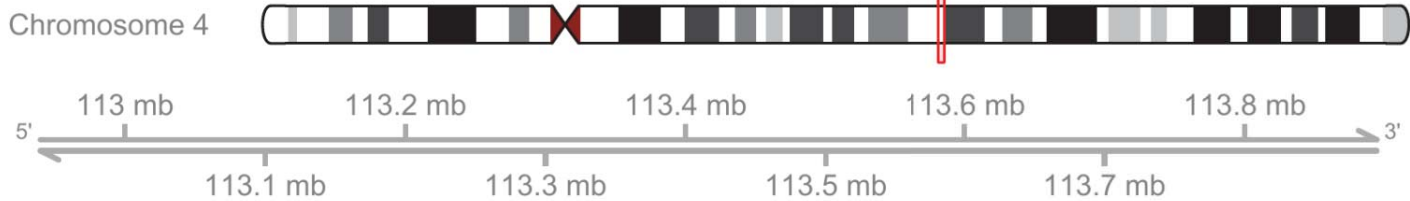
Uncorrected



Zscore of the effect allele: A

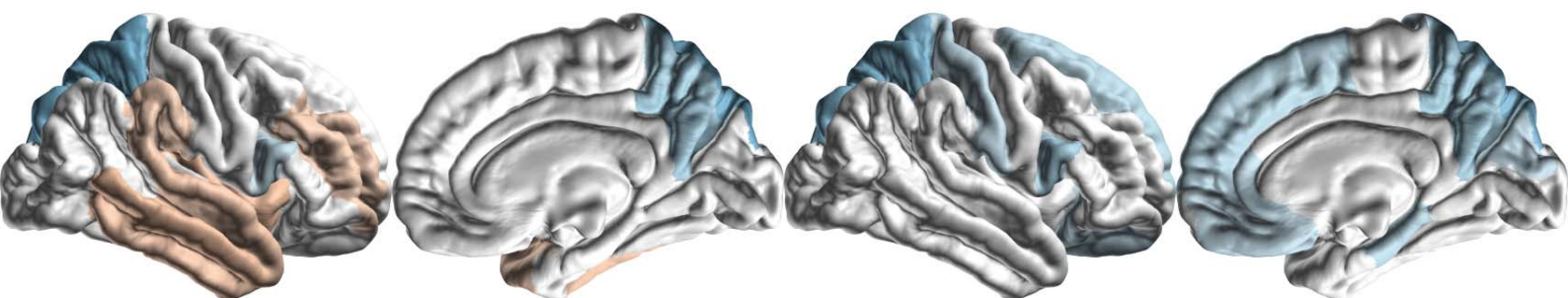


Superior Parietal SA: rs6840242



Corrected for total SA

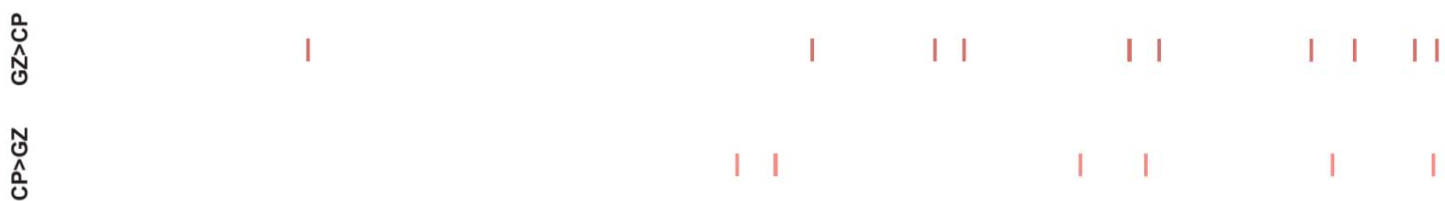
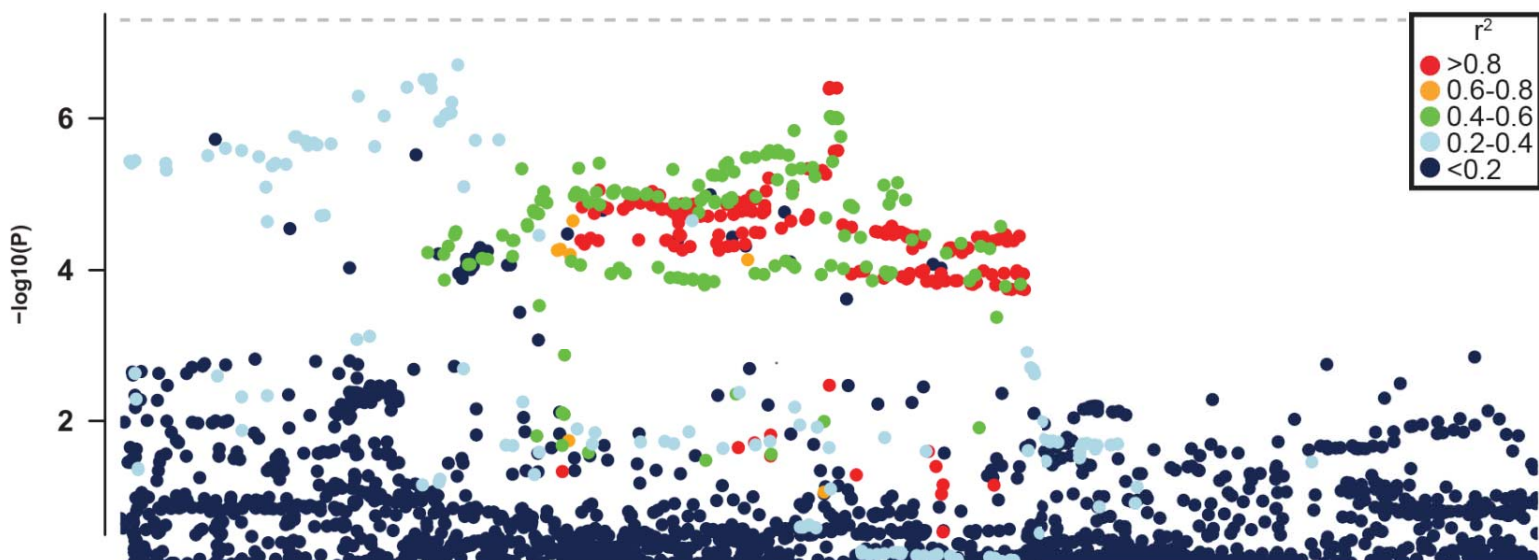
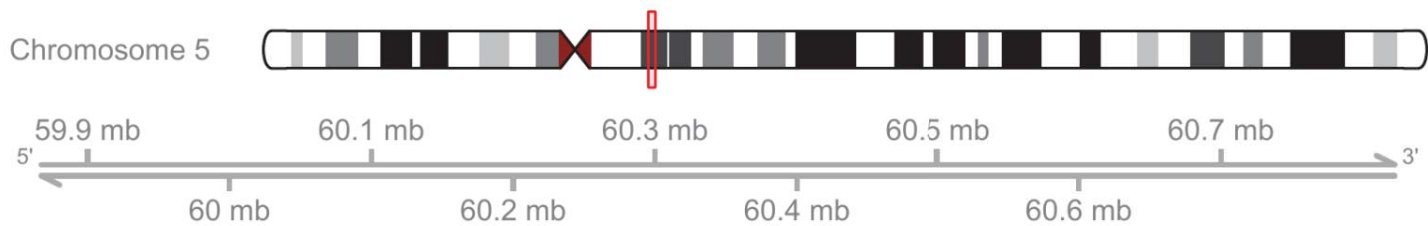
Uncorrected



Zscore of the effect allele: T

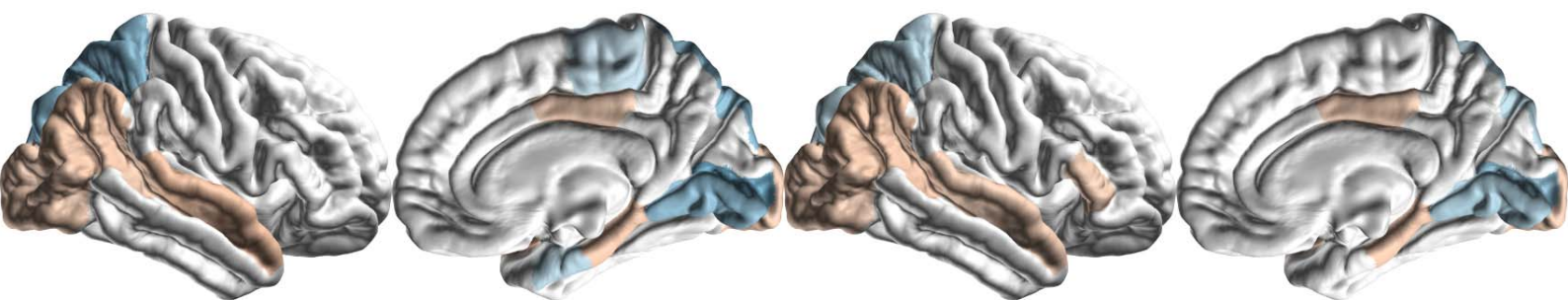


Superior Parietal SA: rs40084

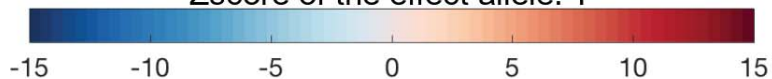


Corrected for total SA

Uncorrected

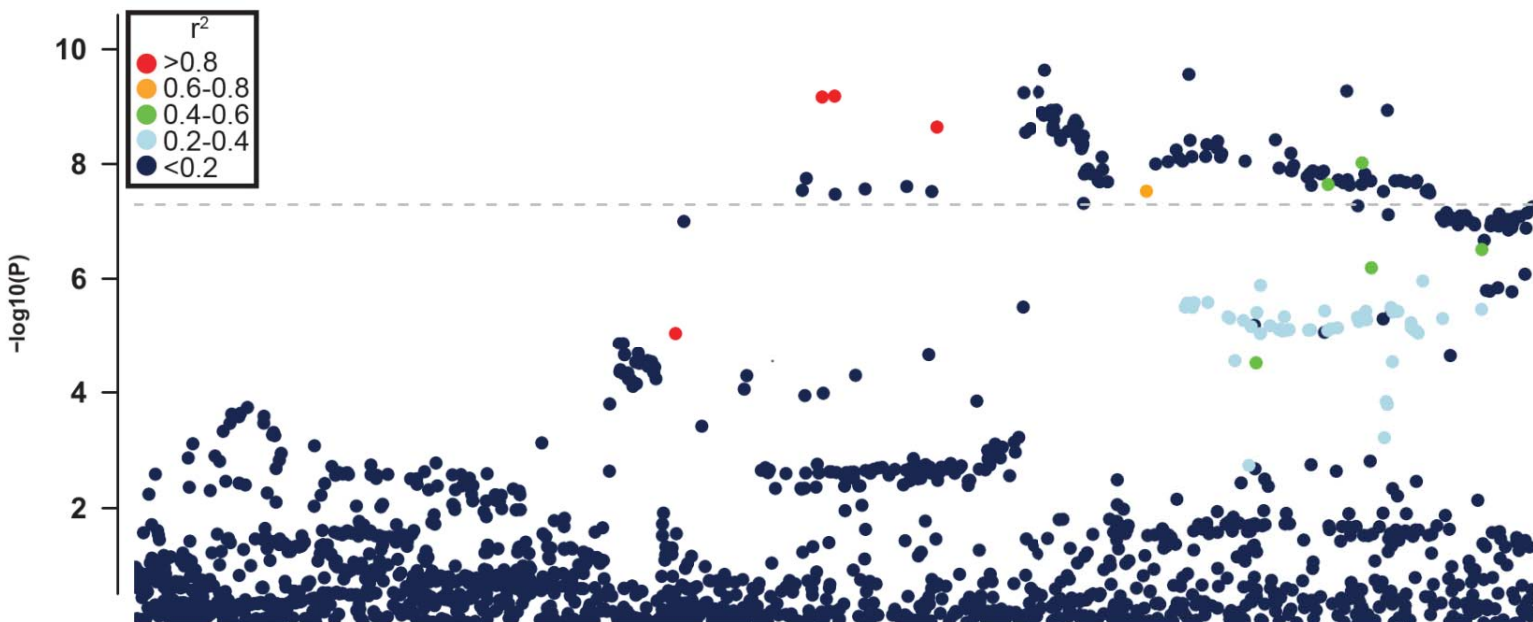


Zscore of the effect allele: T



Superior Parietal SA: rs115877304

Chromosome 5



NR2F1 POU5F2



GZ>CP

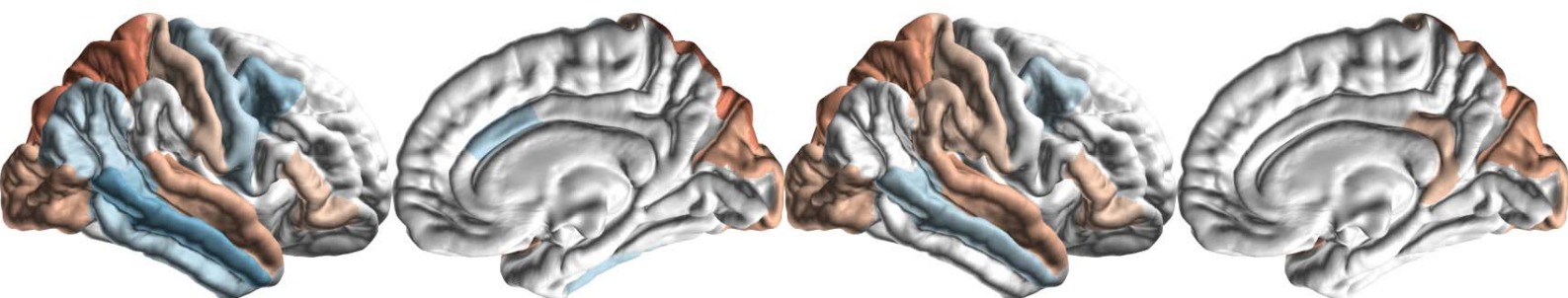


CP>GZ

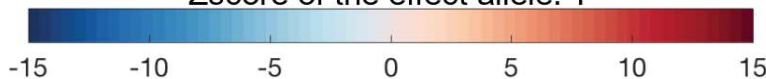


Corrected for total SA

Uncorrected

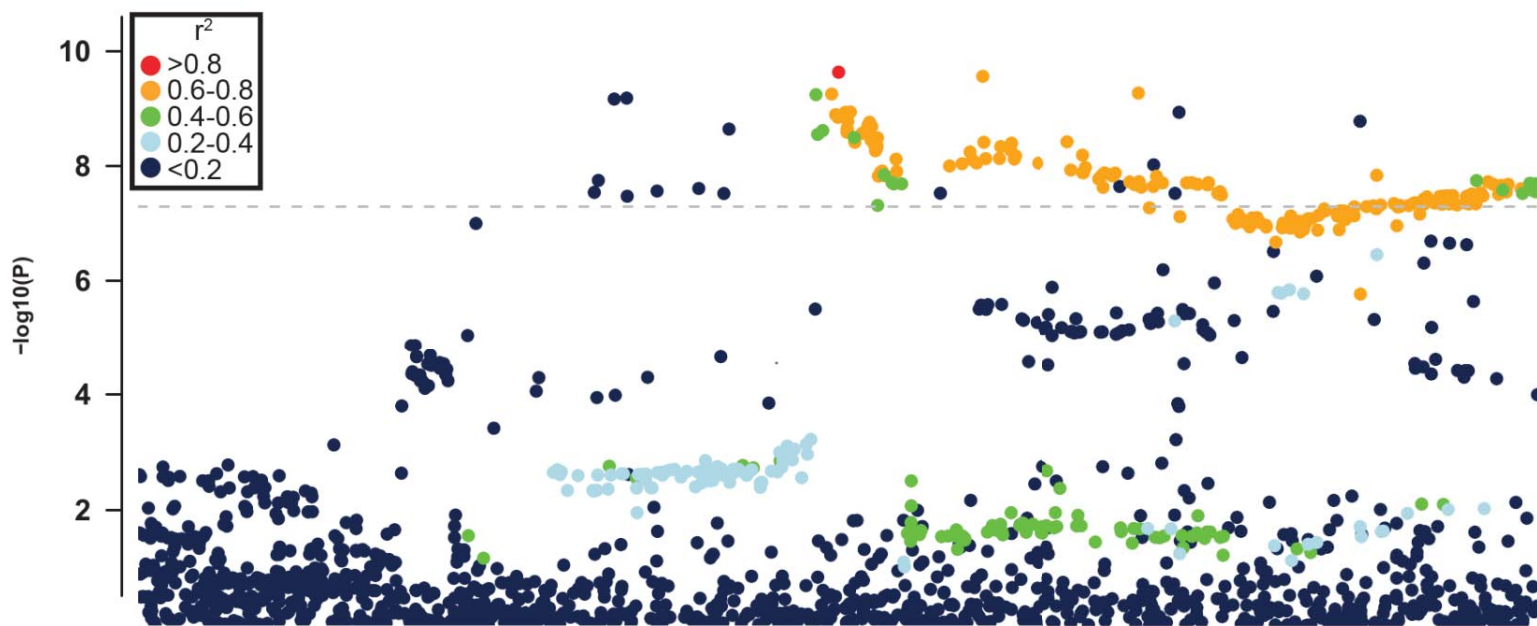


Zscore of the effect allele: T



Superior Parietal SA: rs114489117

Chromosome 5



NR2F1 POU5F2

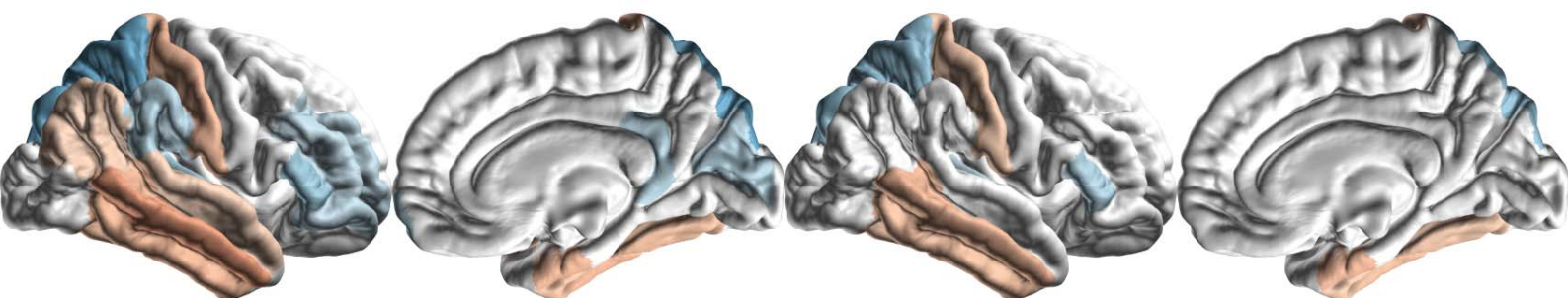


GZ>CP

CP>GZ

Corrected for total SA

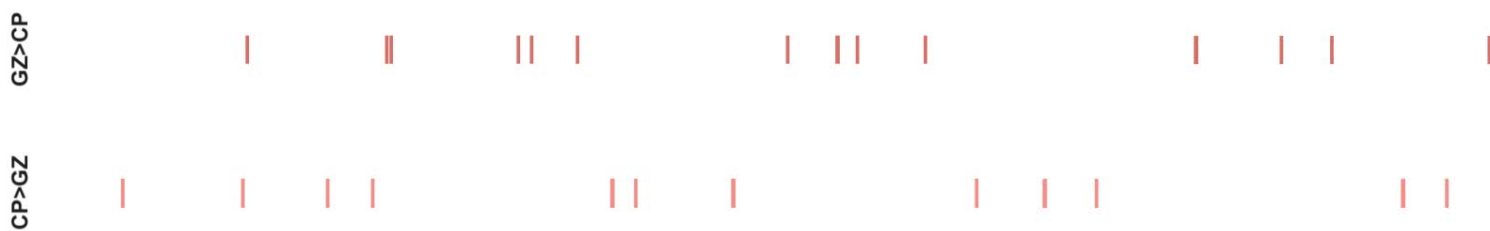
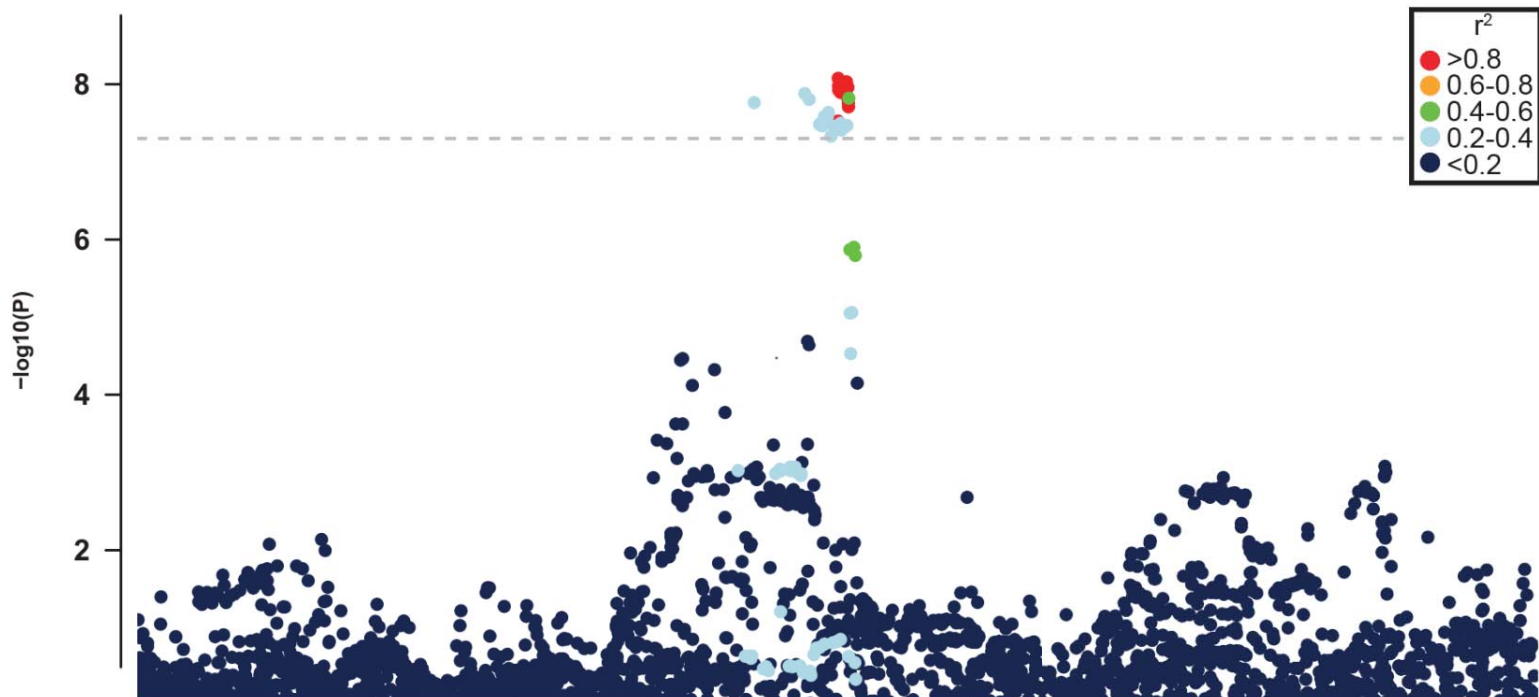
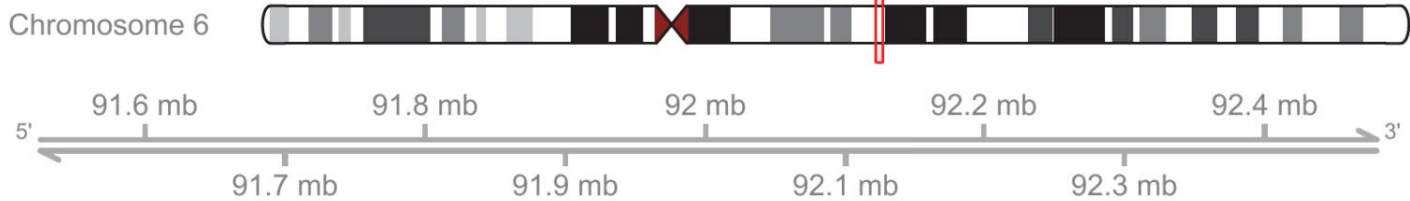
Uncorrected



Zscore of the effect allele: A

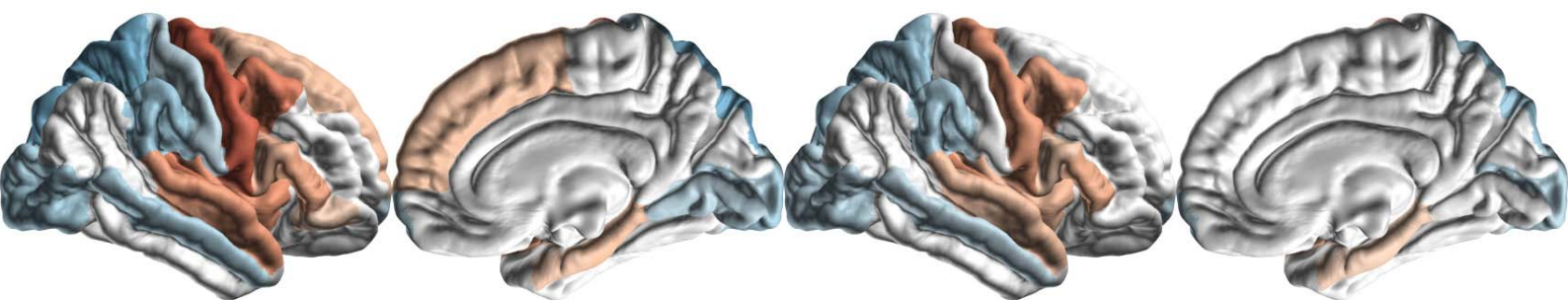


Superior Parietal SA: rs9345125



Corrected for total SA

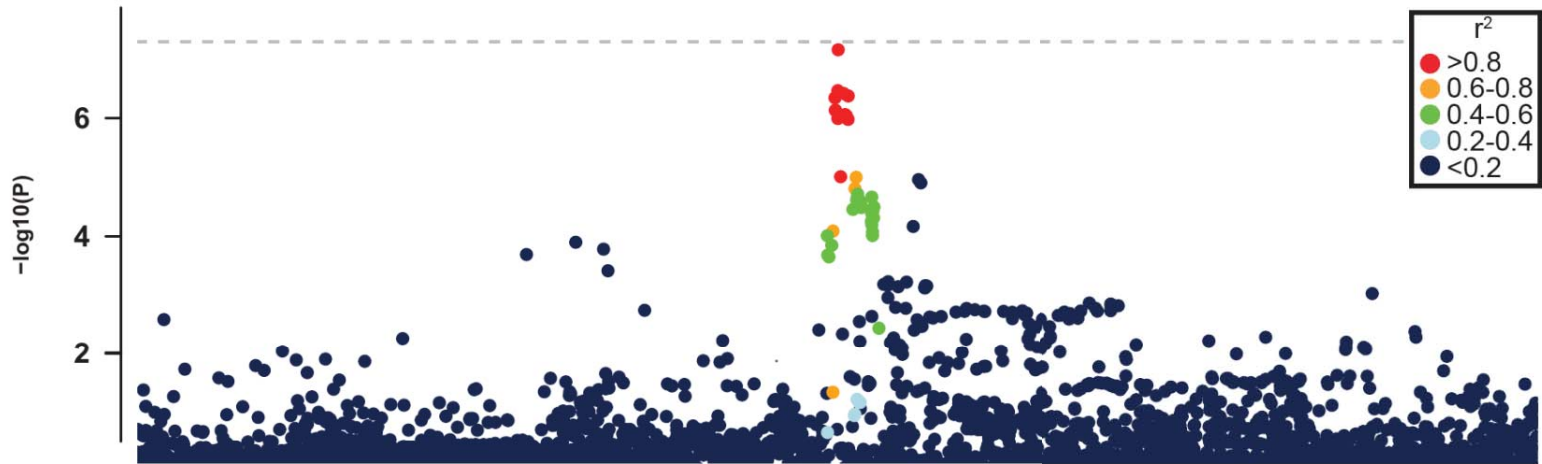
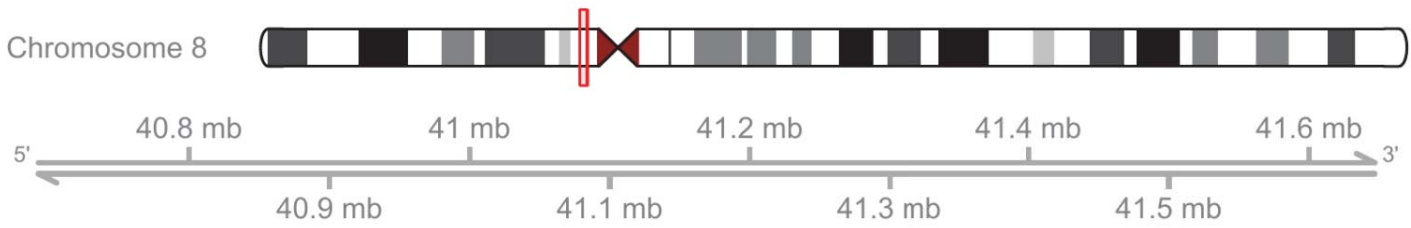
Uncorrected



Zscore of the effect allele: A

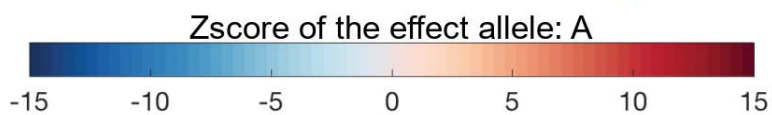
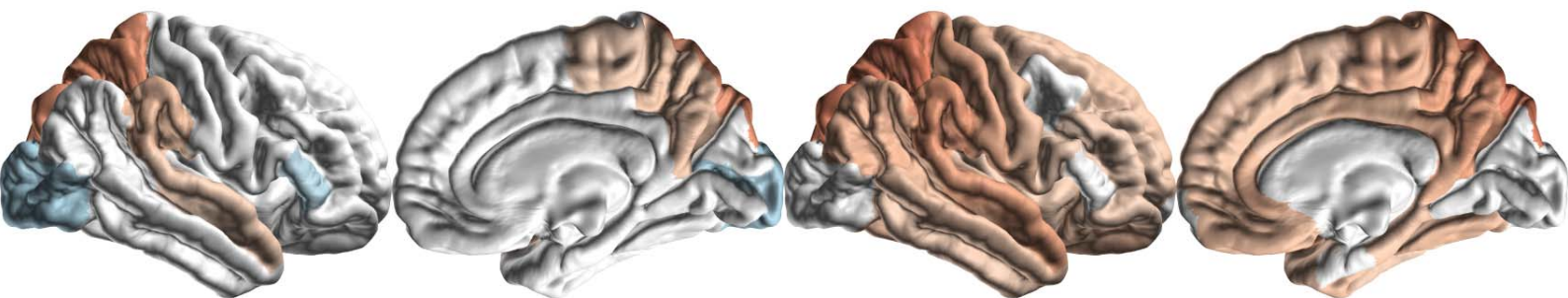


Superior Parietal SA: rs10109434

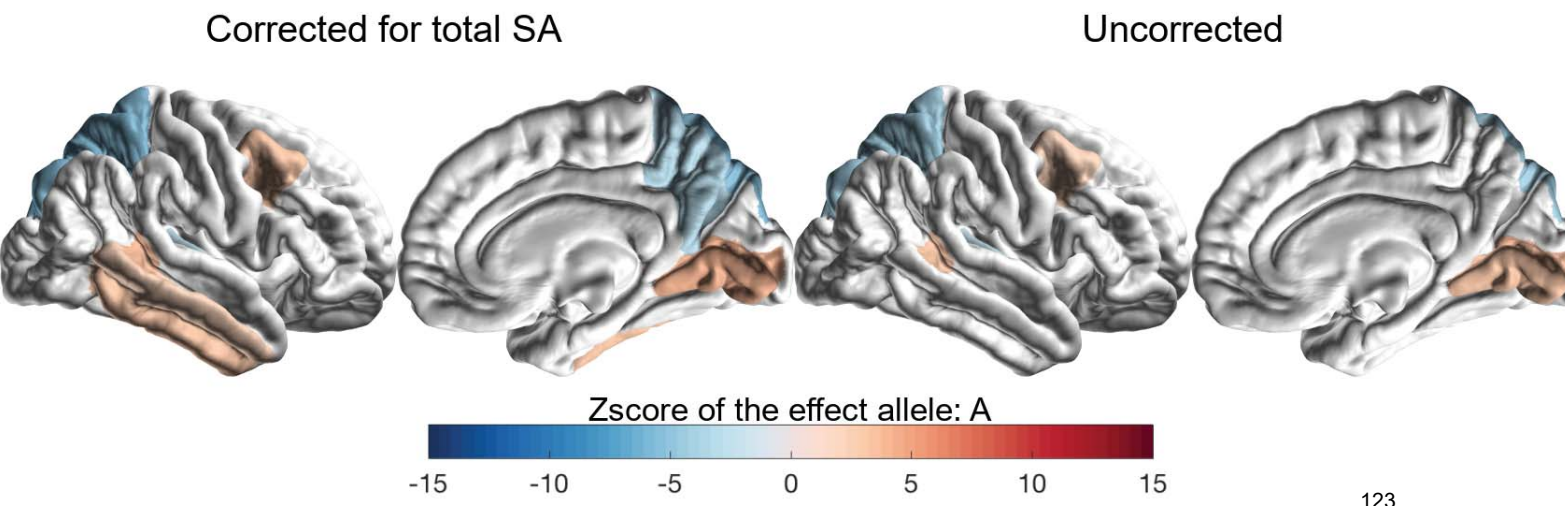
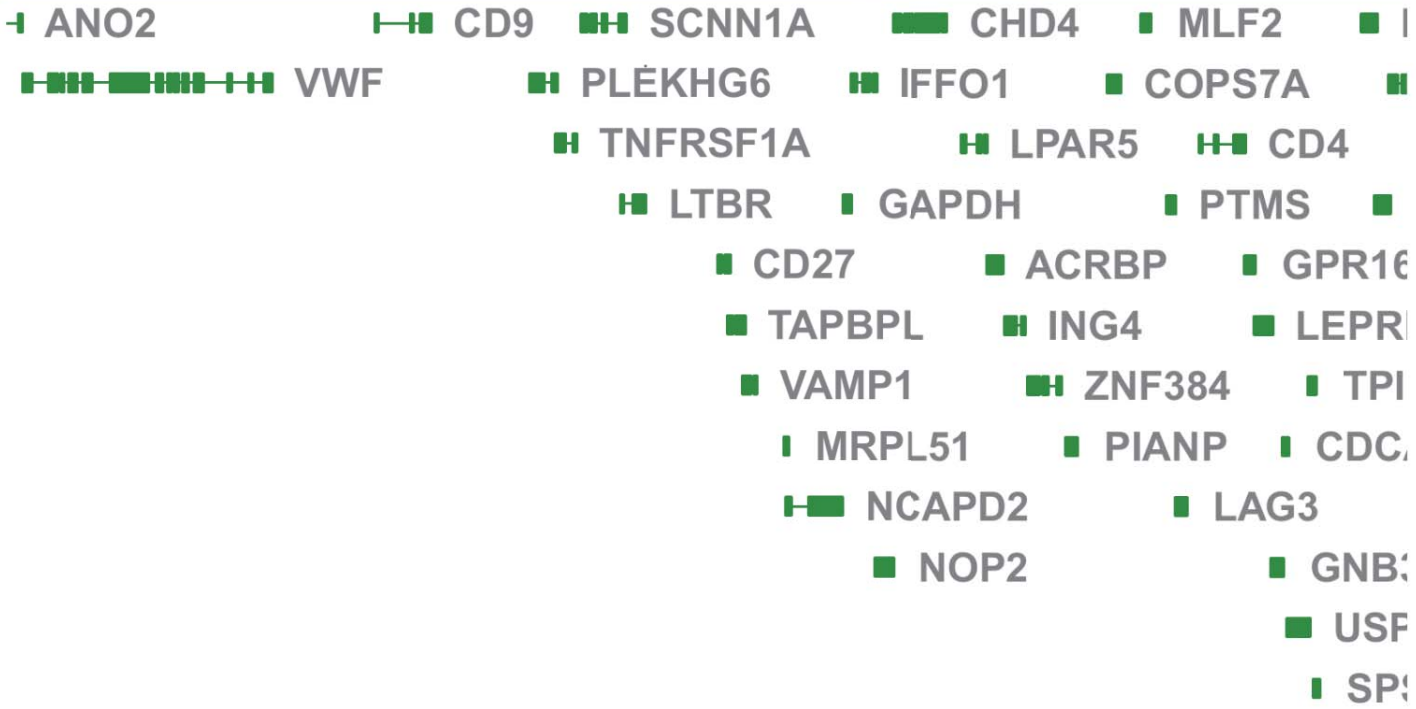
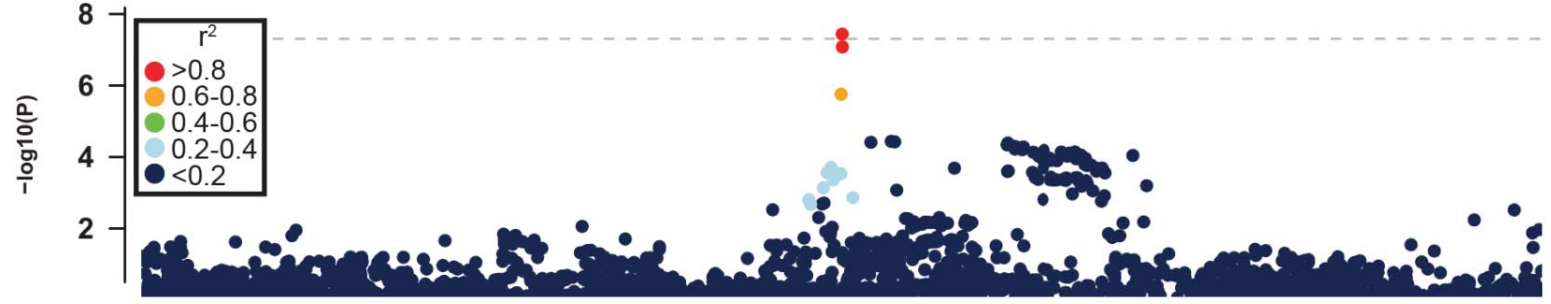
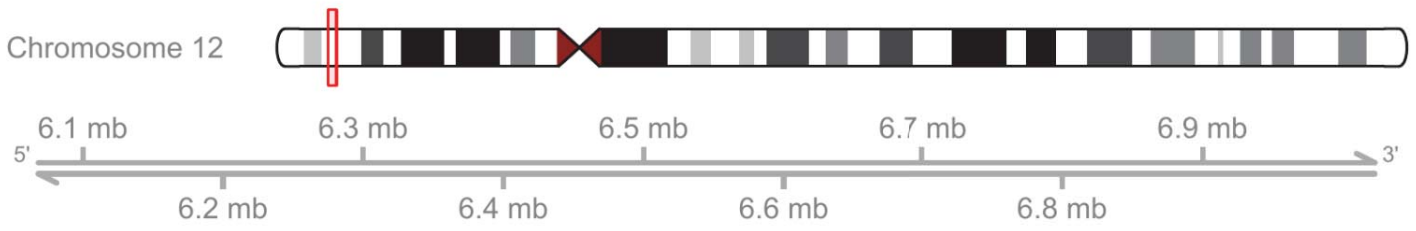


Corrected for total SA

Uncorrected

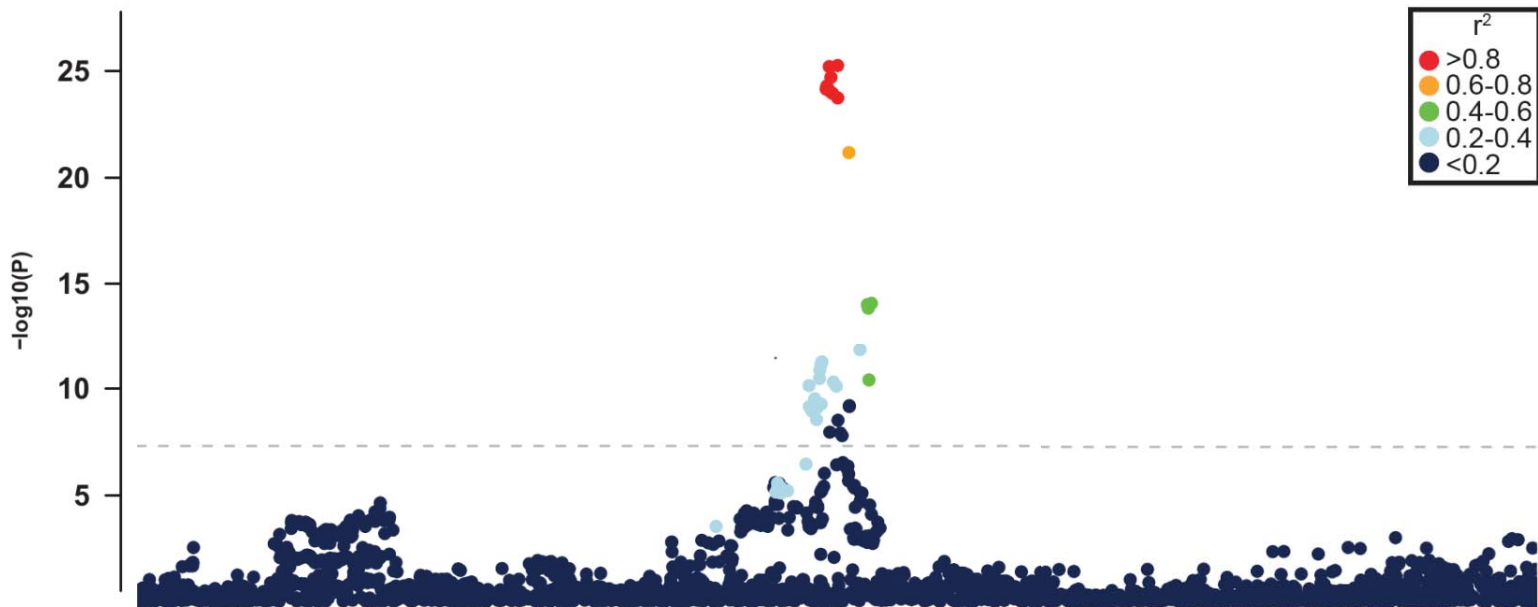
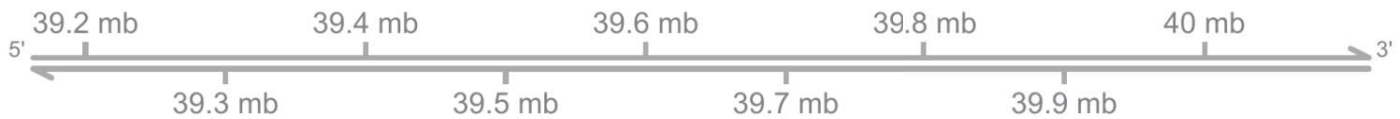


Superior Parietal SA: rs7980991



Superior Parietal SA: rs4924345

Chromosome 15



C15orf54

THBS1

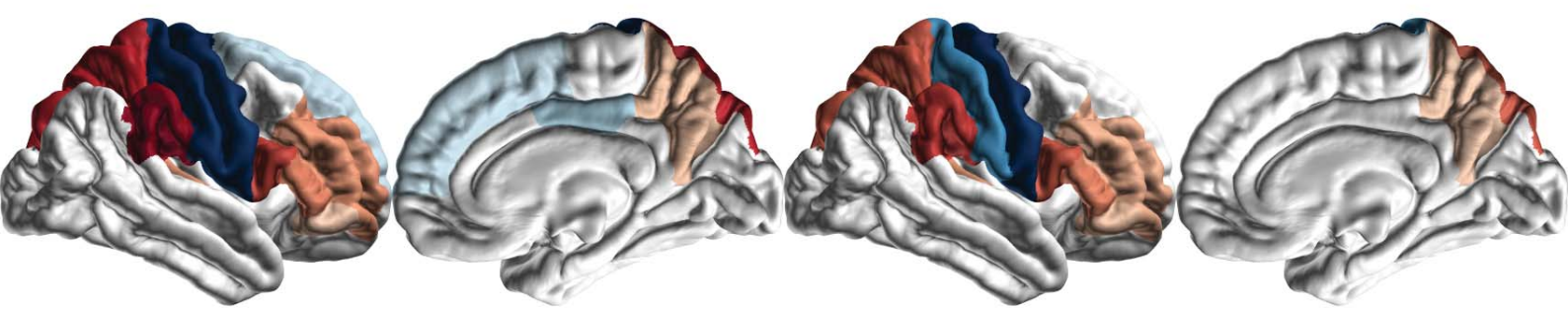


GZ>CP
CP>GZ

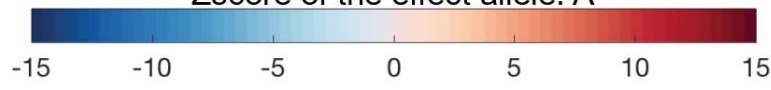


Corrected for total SA

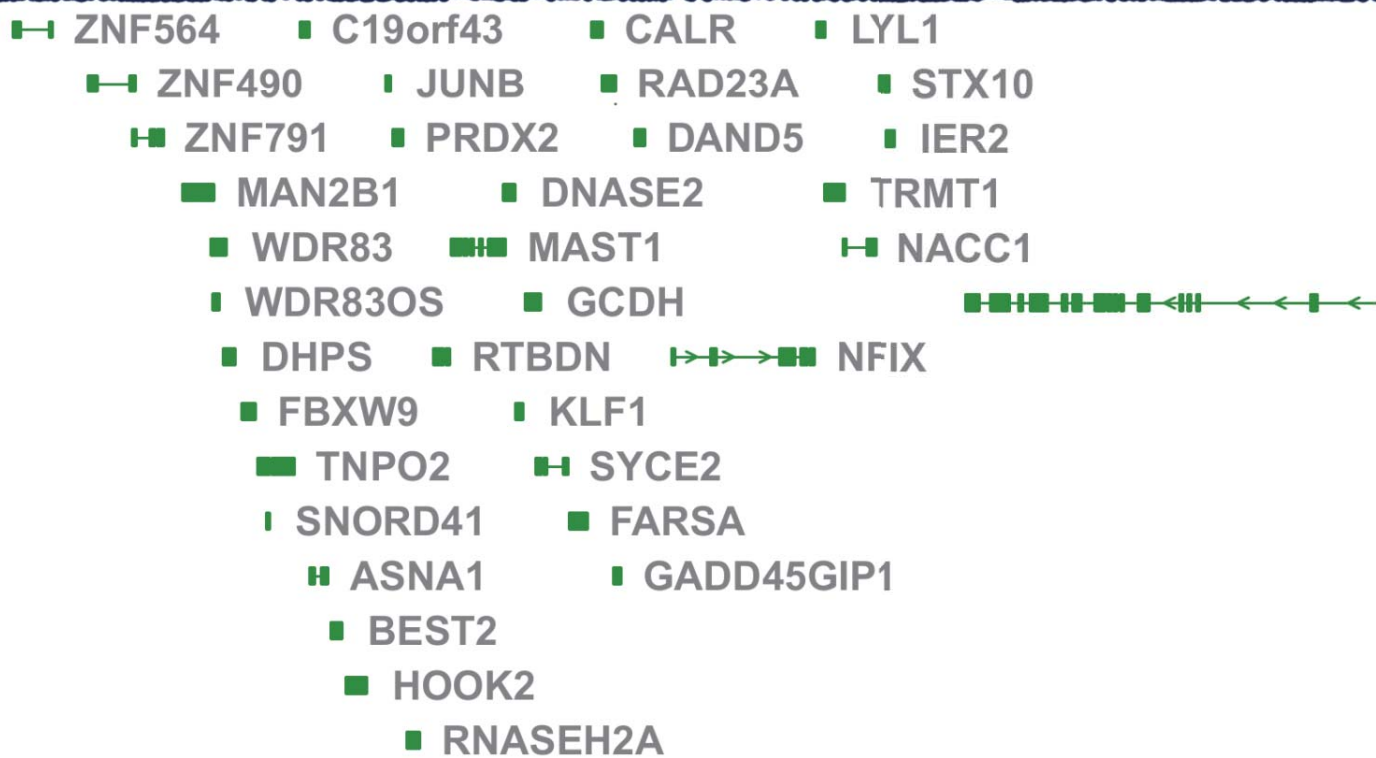
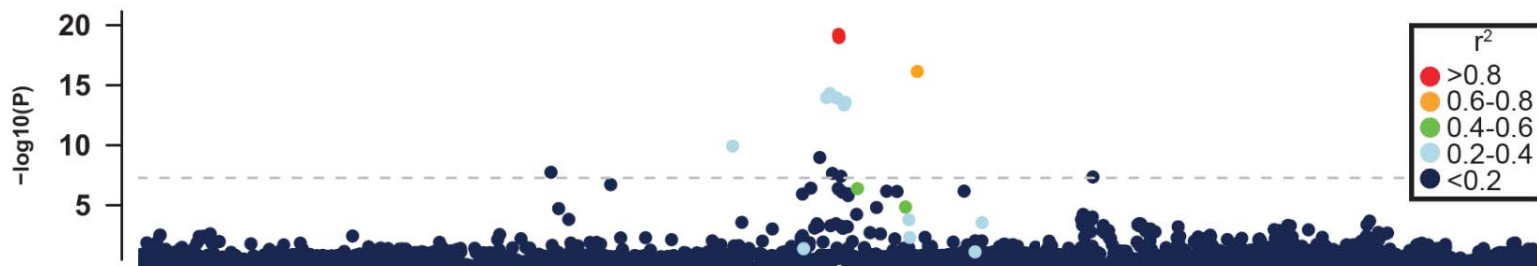
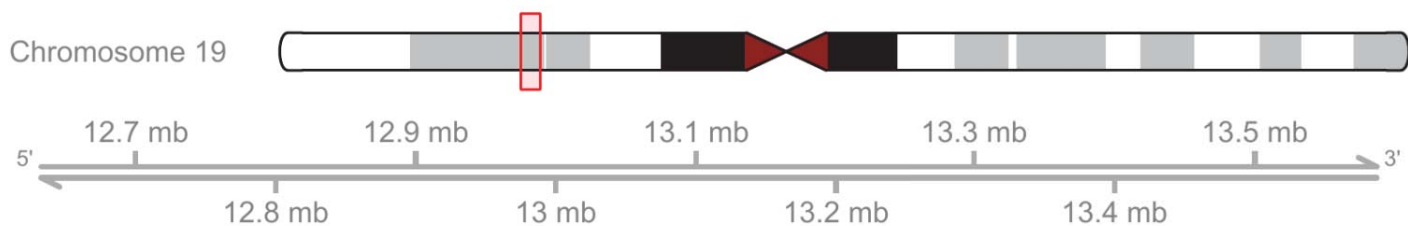
Uncorrected



Zscore of the effect allele: A



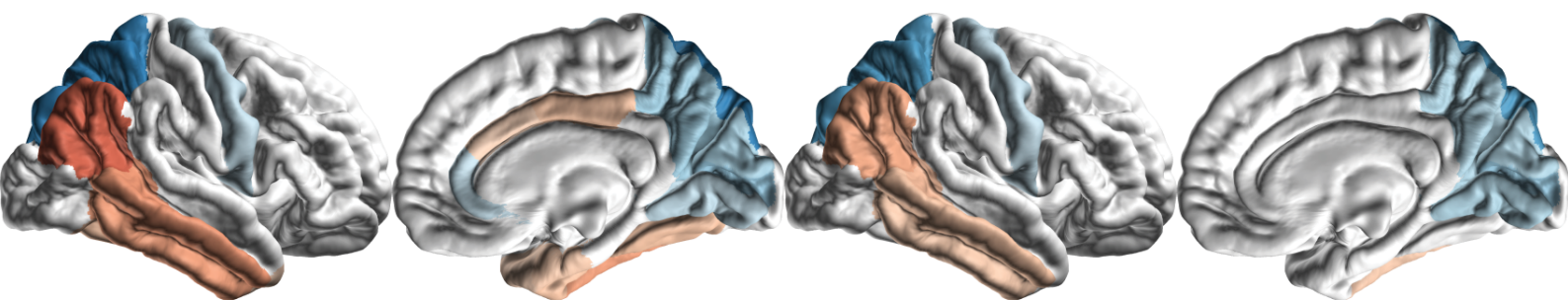
Superior Parietal SA: rs68175985



CP>GZZ>CP

Corrected for total SA

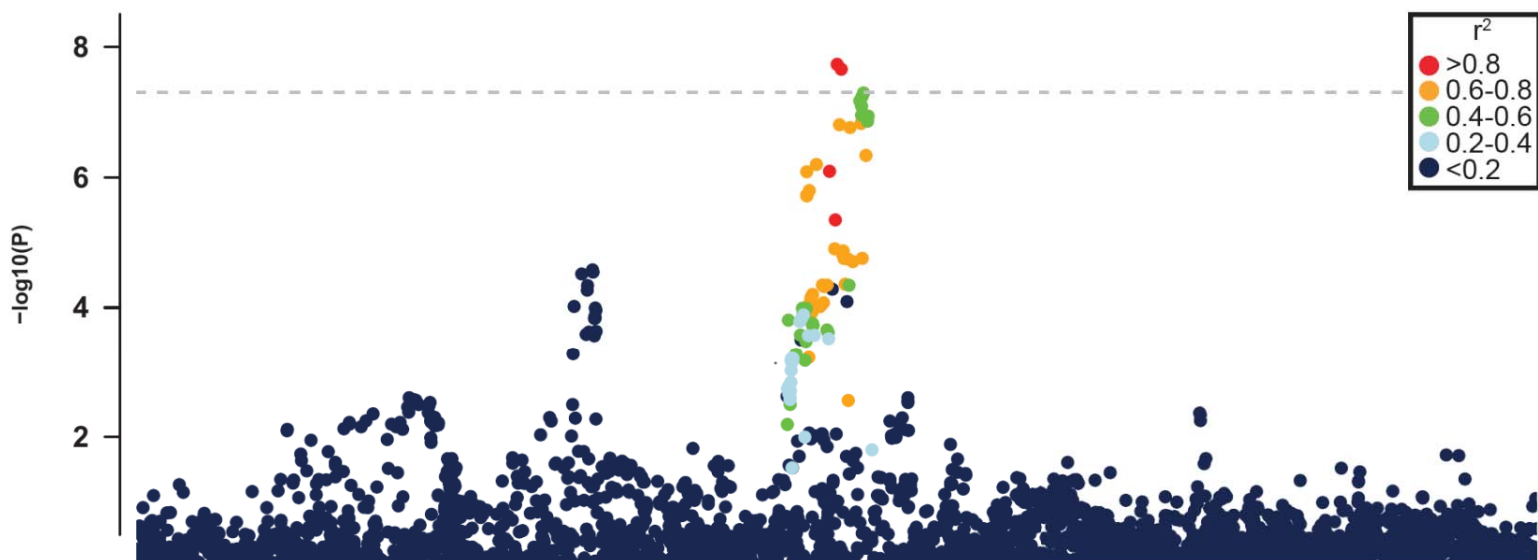
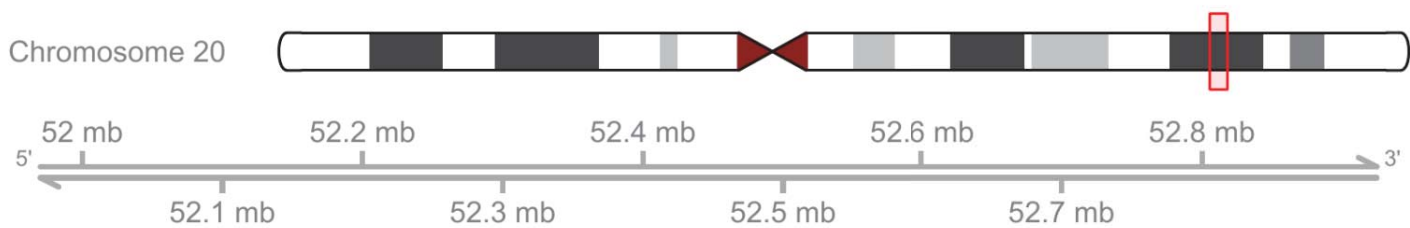
Uncorrected



Zscore of the effect allele: A



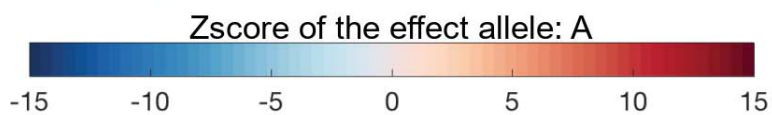
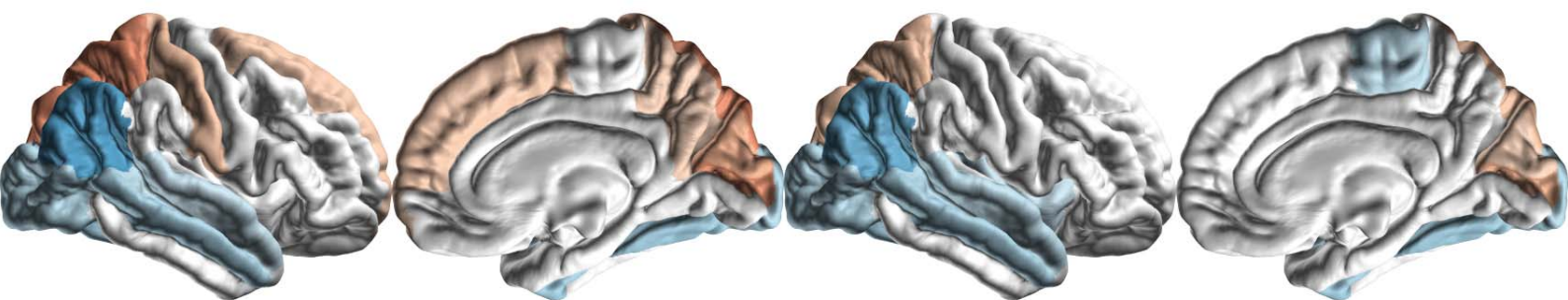
Superior Parietal SA: rs6022786



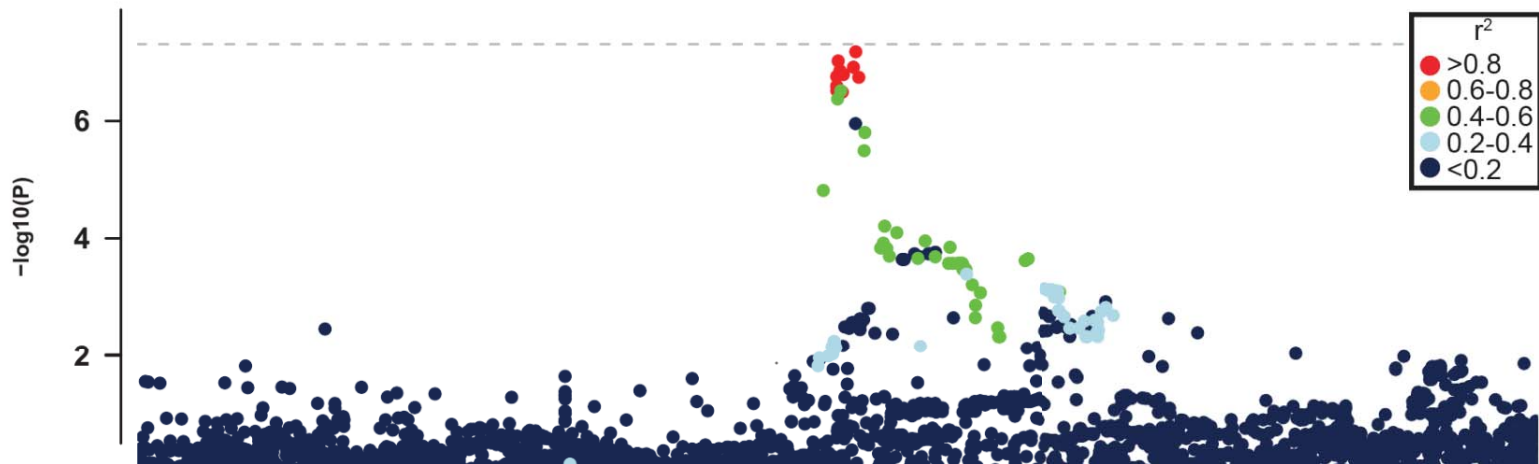
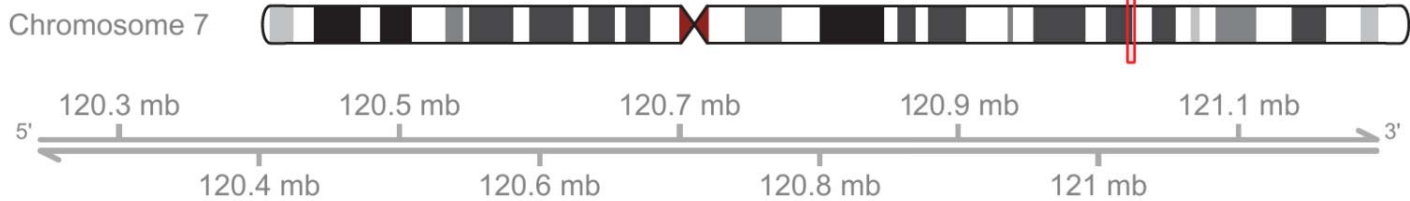
CP>GZ
GZ>CP

Corrected for total SA

Uncorrected



Superior Parietal TH: rs73215353

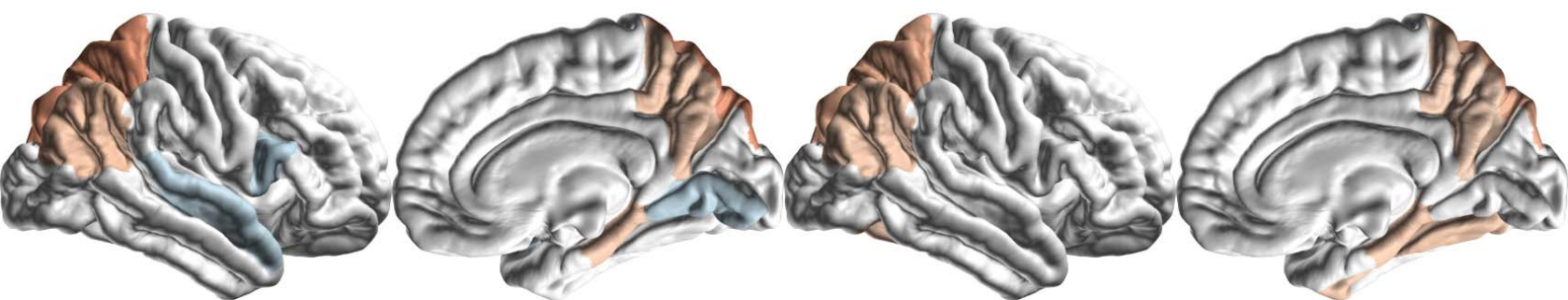


Angular Temporal dIPFC Cingulate

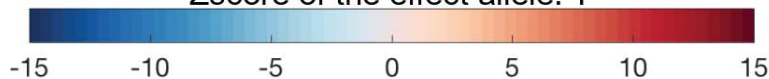


Corrected for average TH

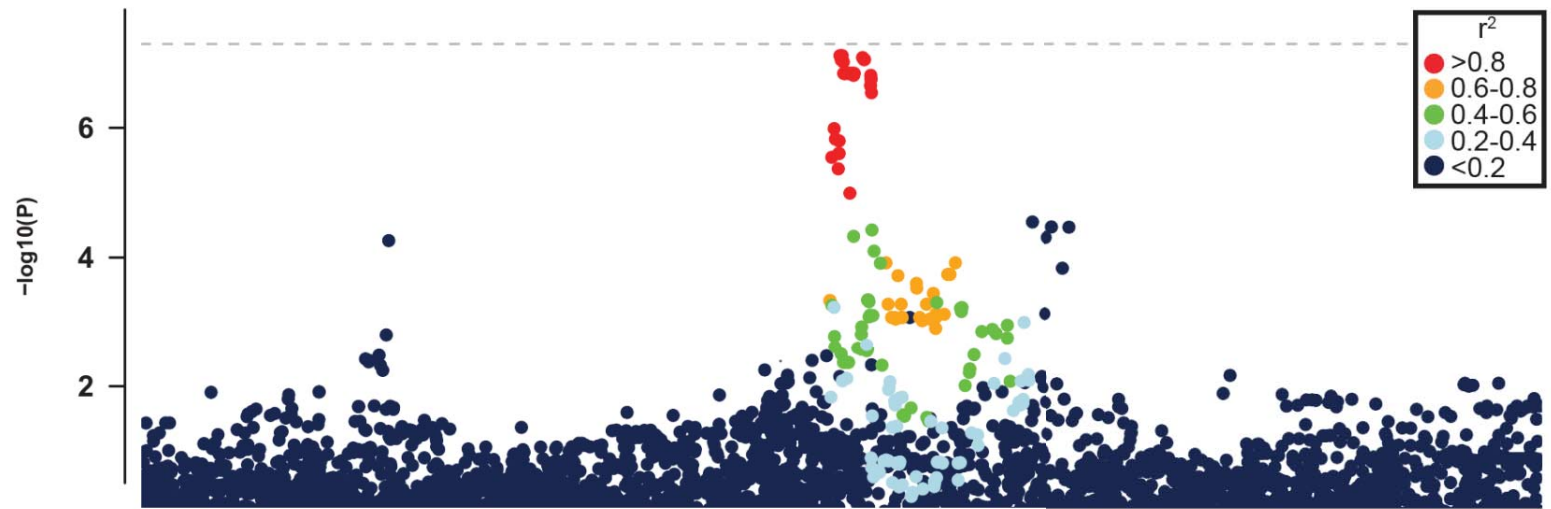
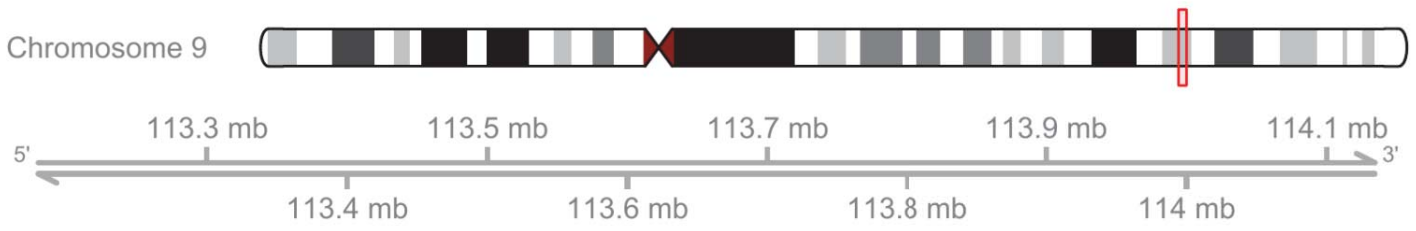
Uncorrected



Zscore of the effect allele: T

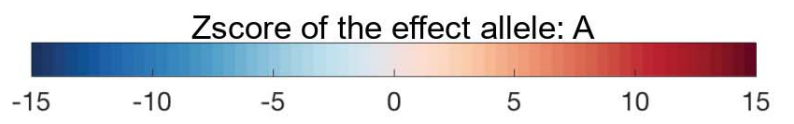
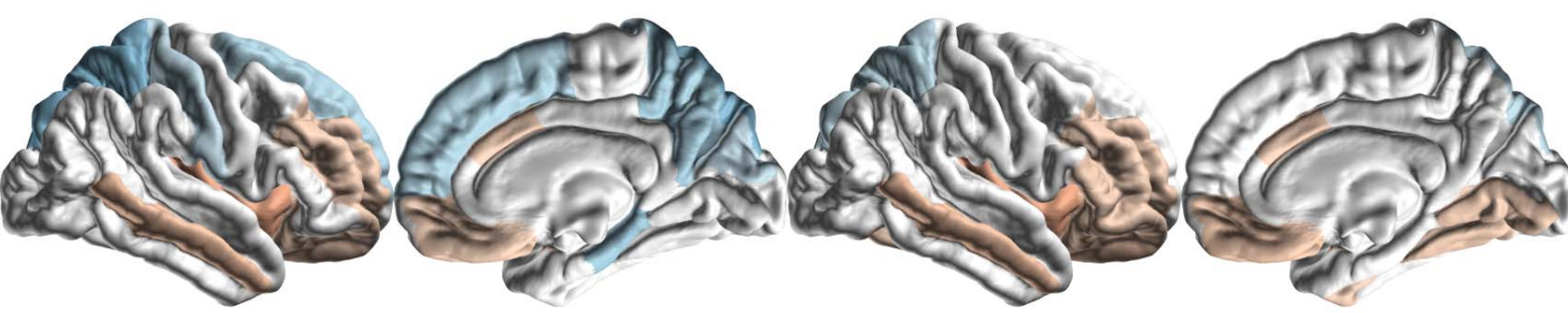


Superior Parietal TH: rs1326892

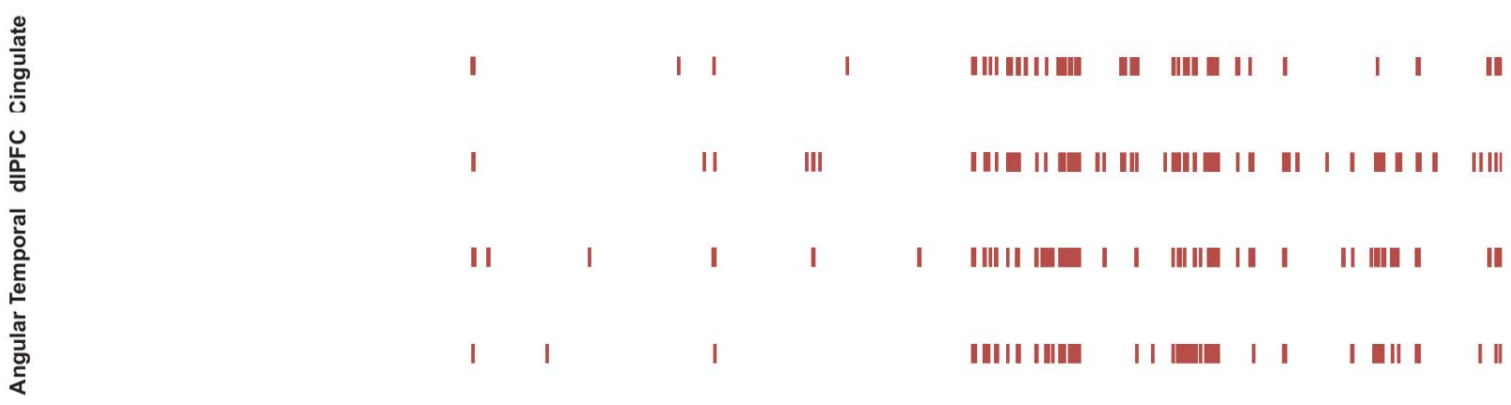
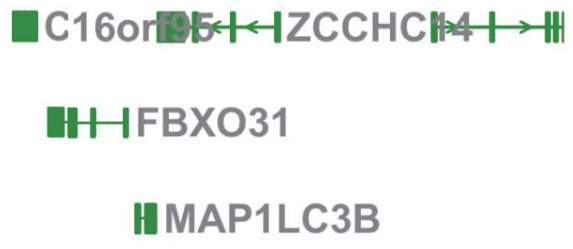
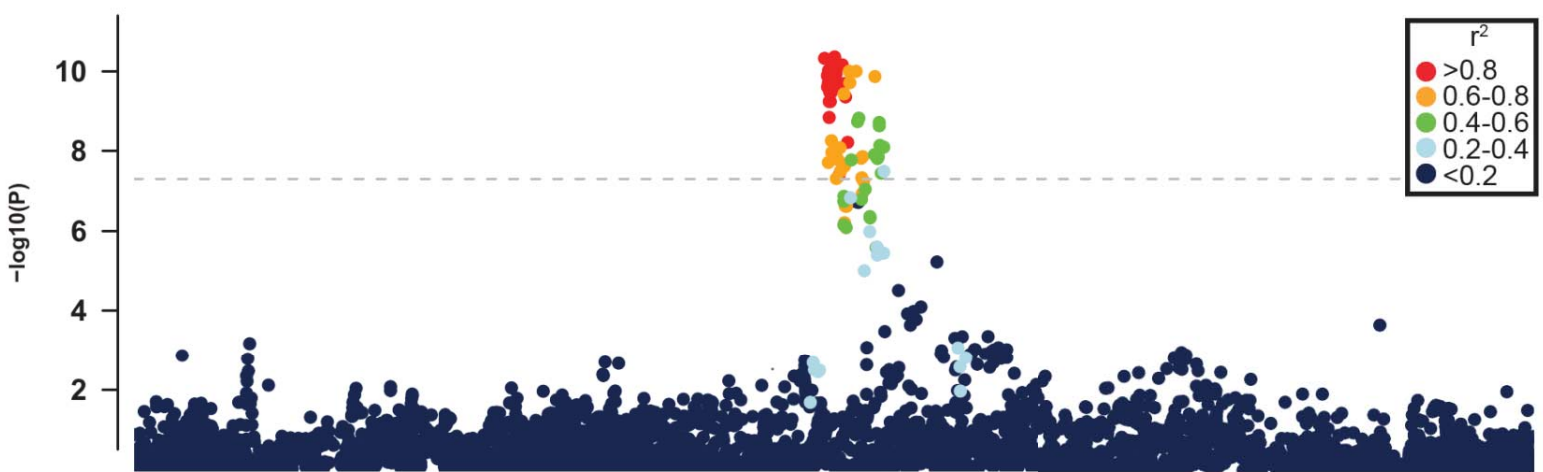
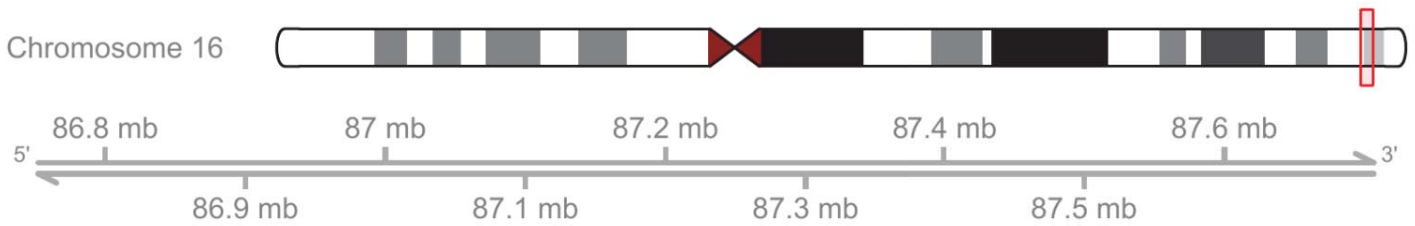


Corrected for average TH

Uncorrected

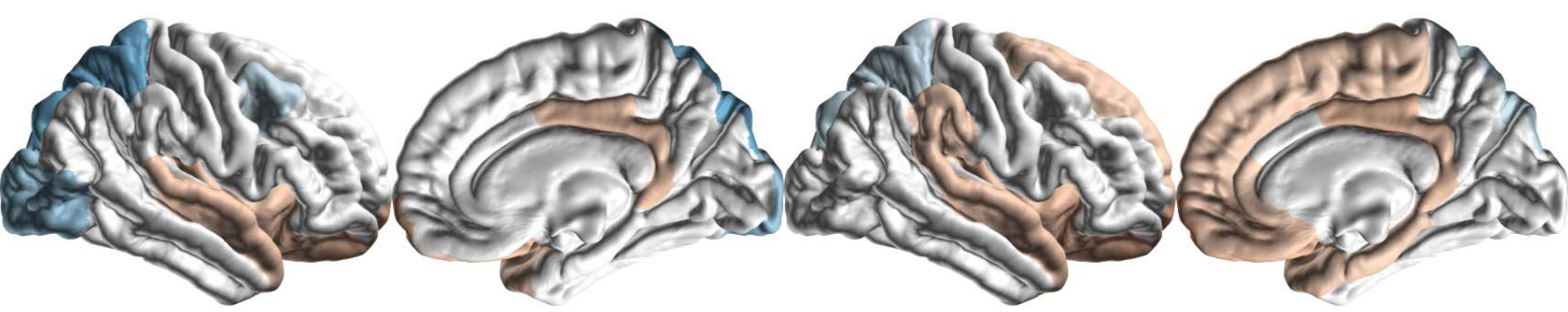


Superior Parietal TH: rs7203251

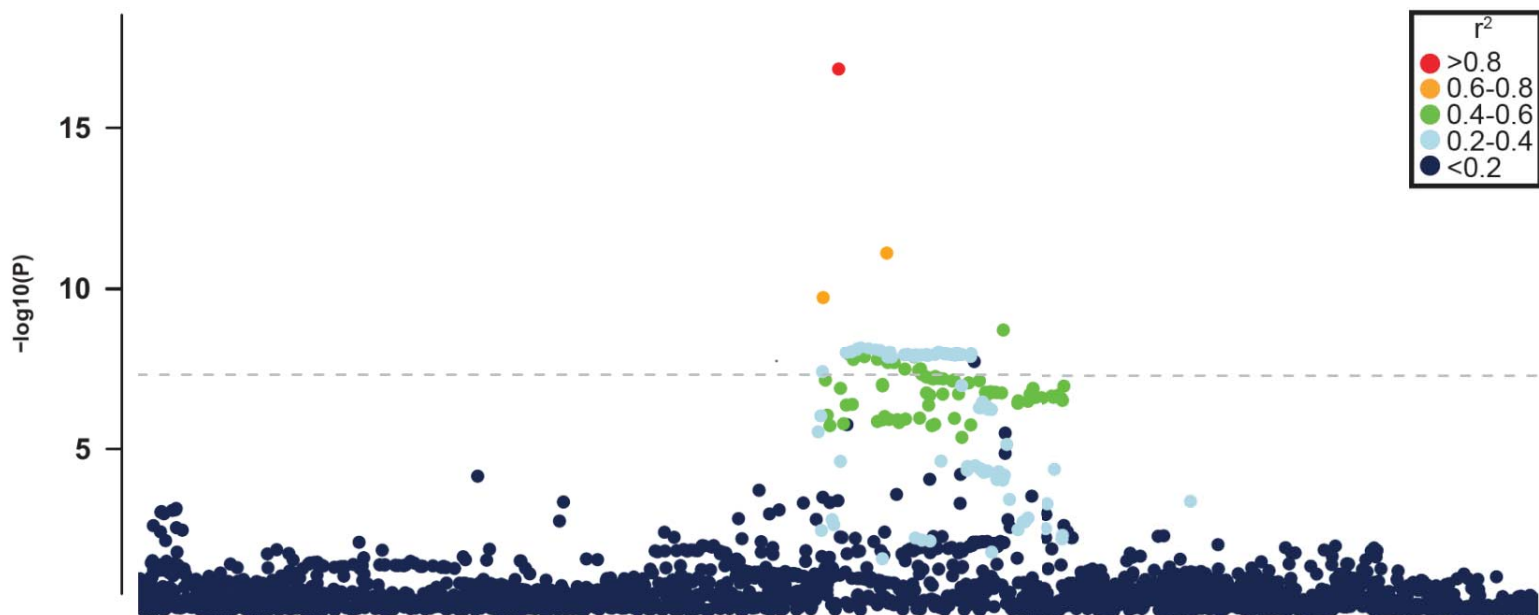
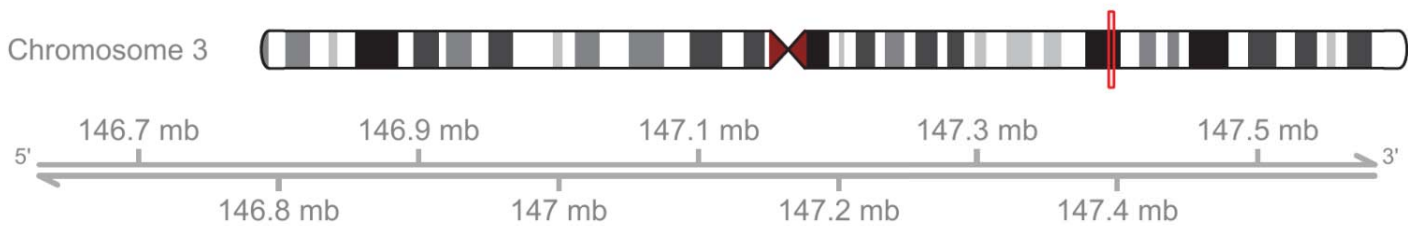


Corrected for average TH

Uncorrected



Supramarginal SA: rs2279829



ZIC4

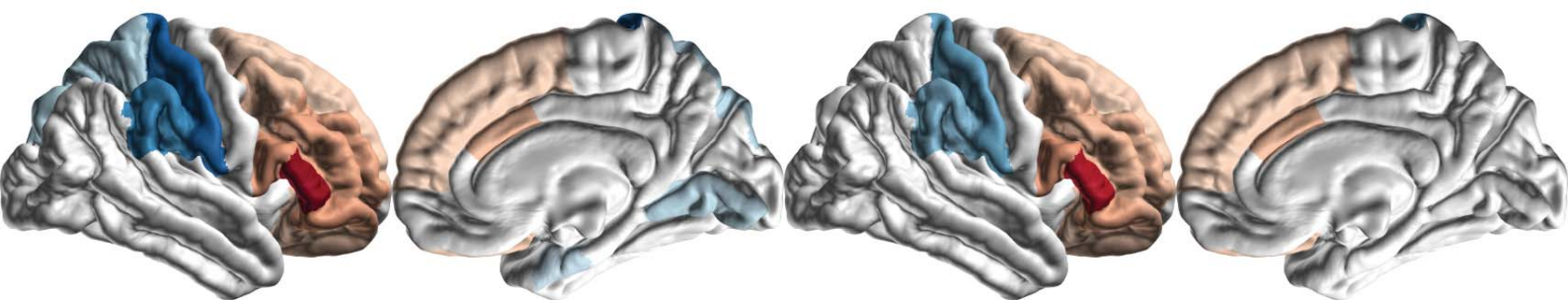
ZIC1

GZ>CP

CP>GZ

Corrected for total SA

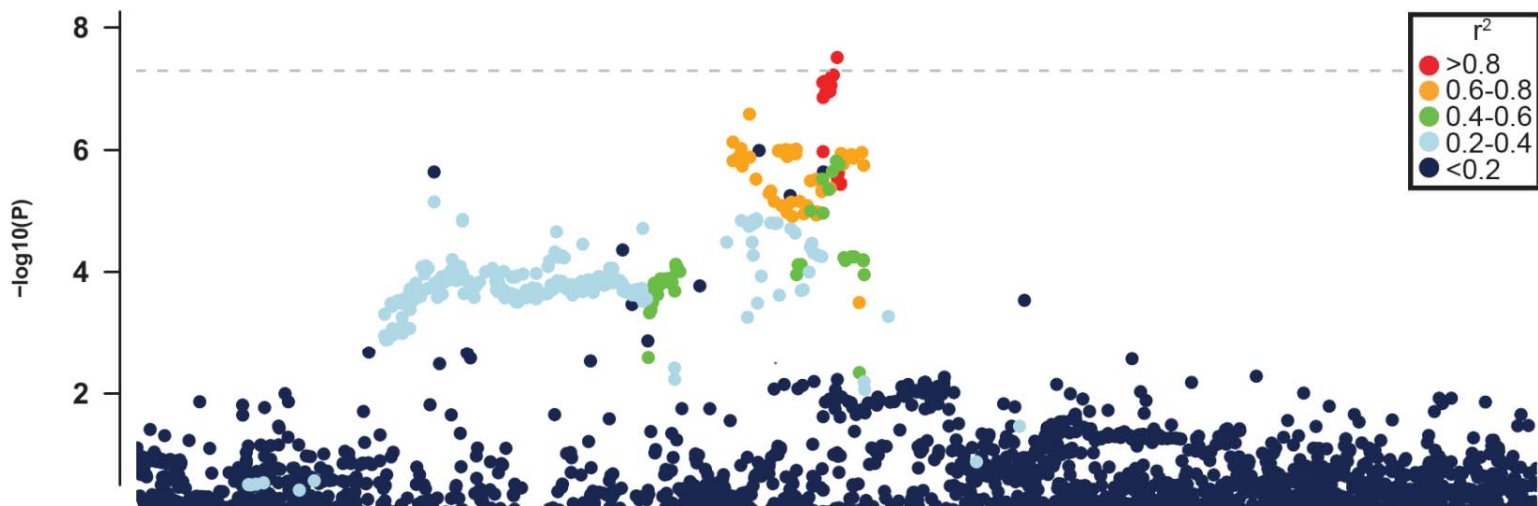
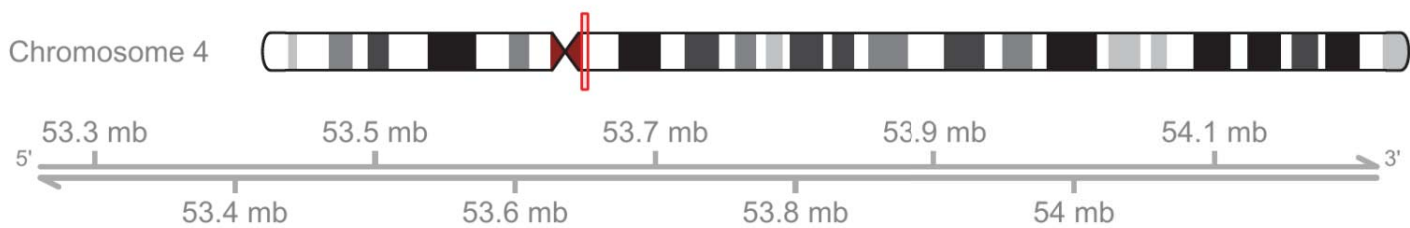
Uncorrected



Zscore of the effect allele: T



Supramarginal SA: rs2200225



USP46

RASL11B

SNORA26

ERVMER34-1

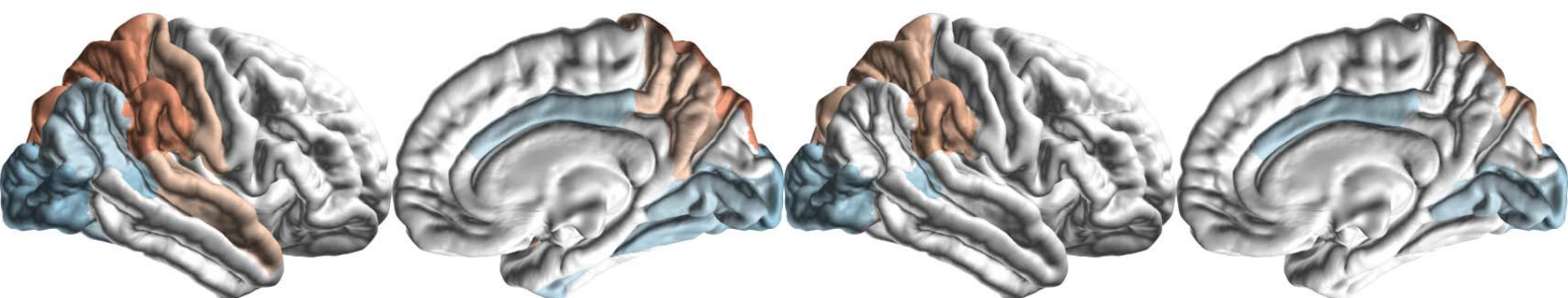


CP>GZ GZ>CP



Corrected for total SA

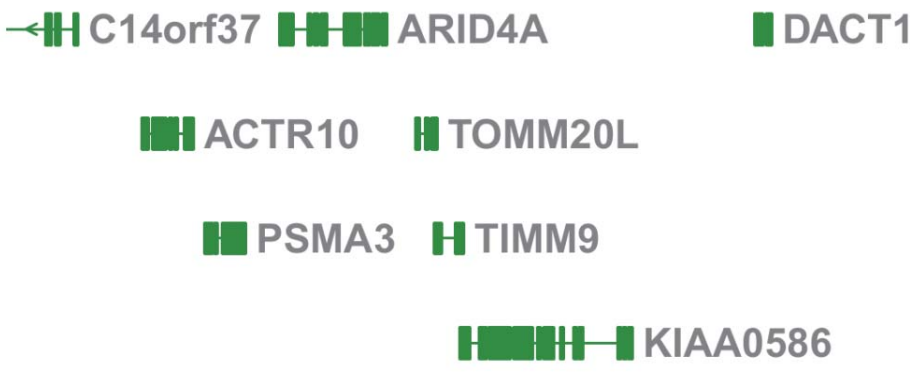
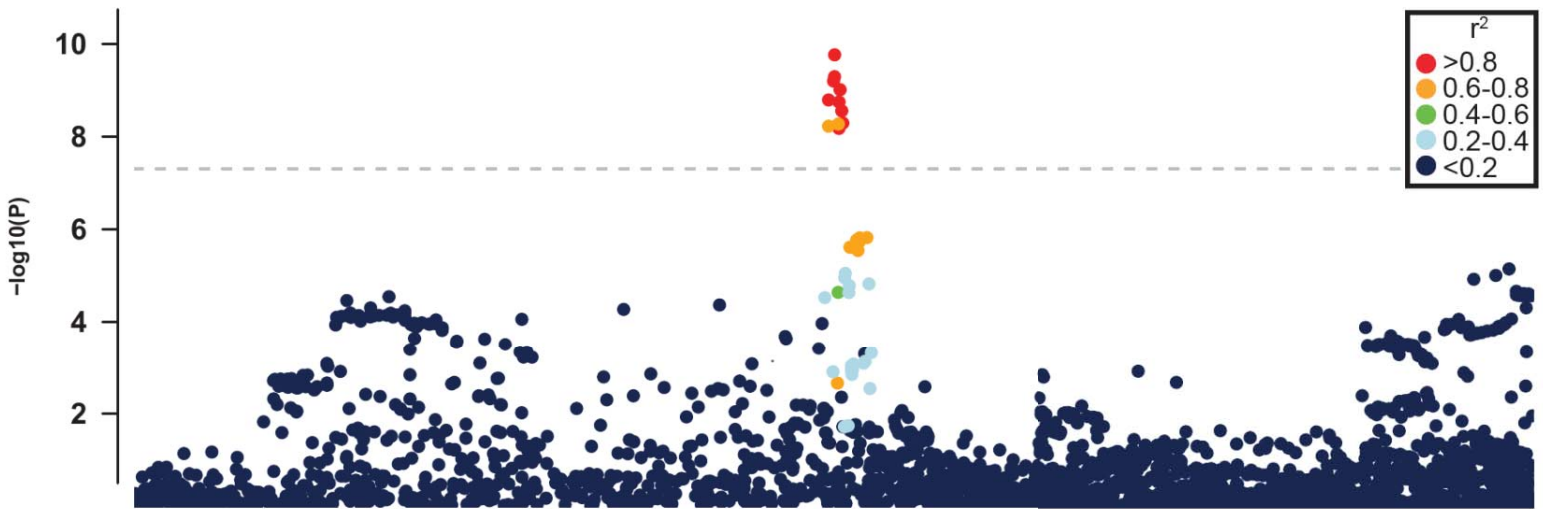
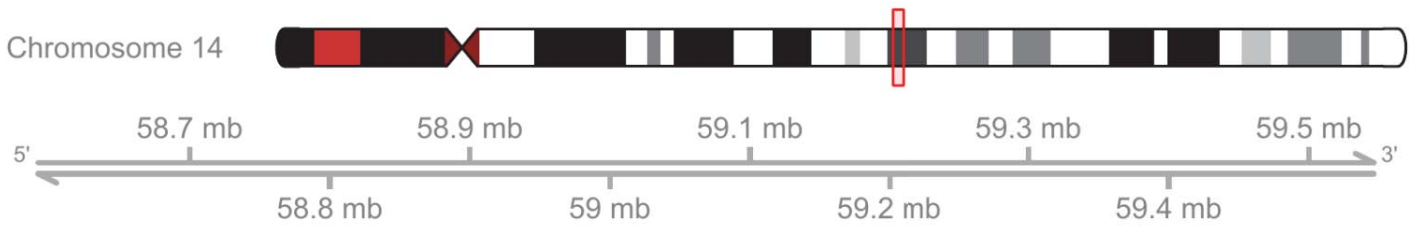
Uncorrected



Zscore of the effect allele: A

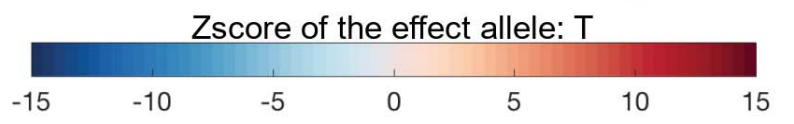
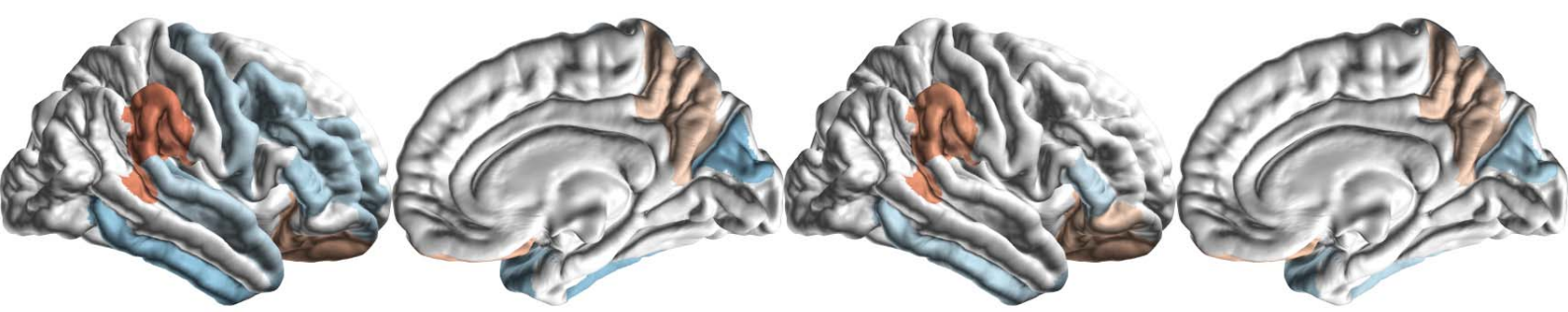


Supramarginal SA: rs221326



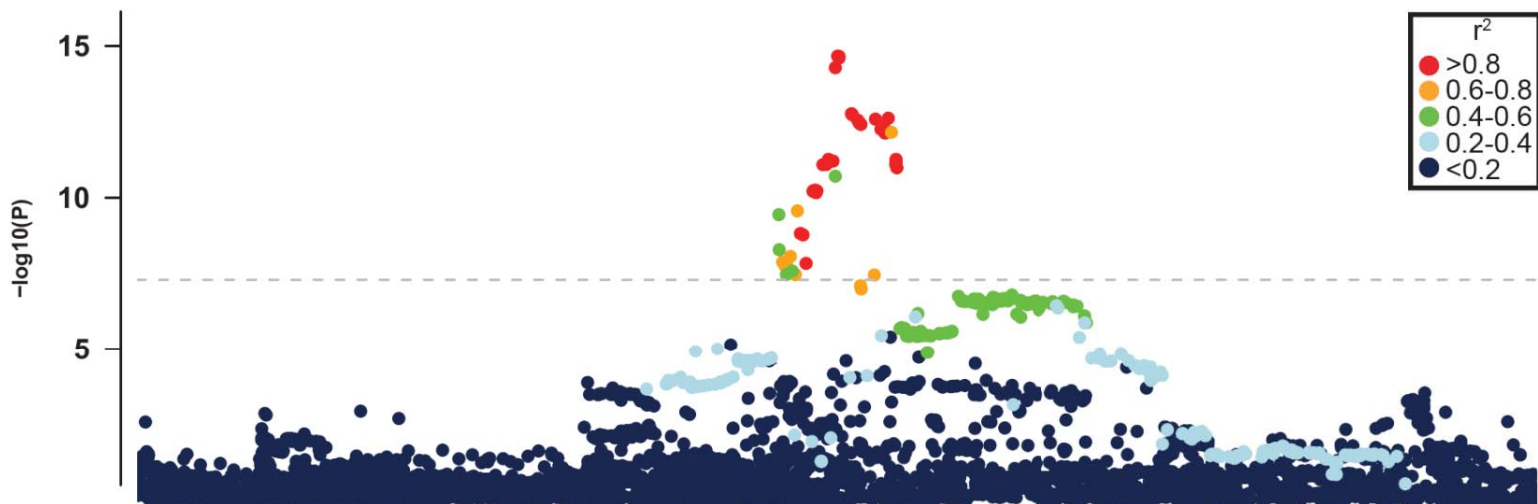
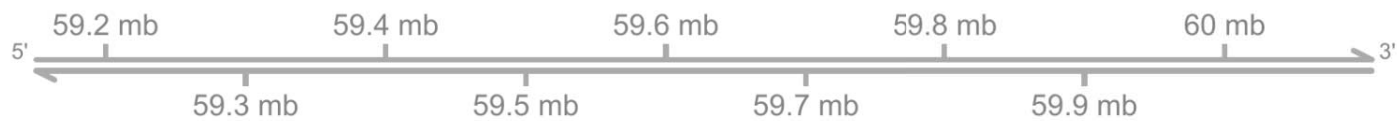
Corrected for total SA

Uncorrected



Supramarginal SA: rs2164950

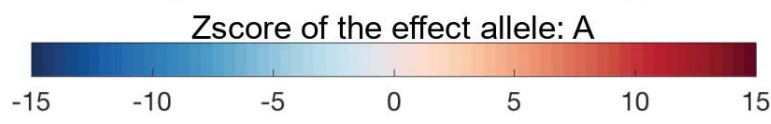
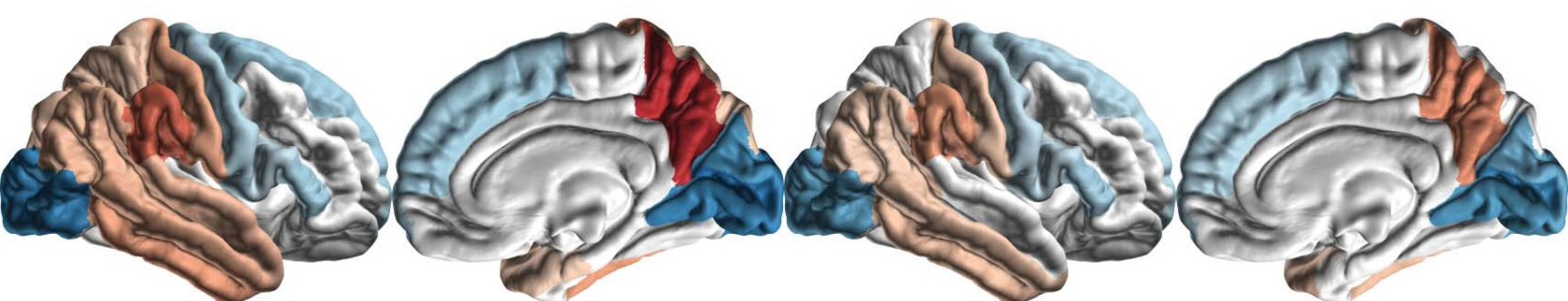
Chromosome 14



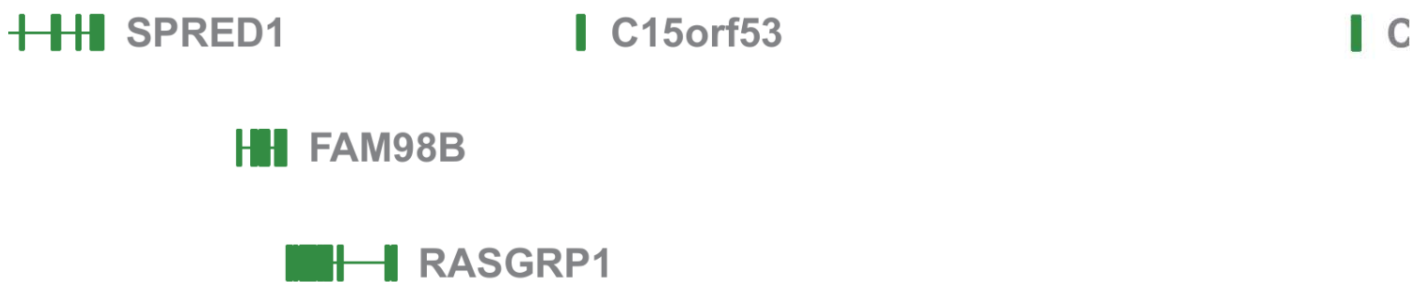
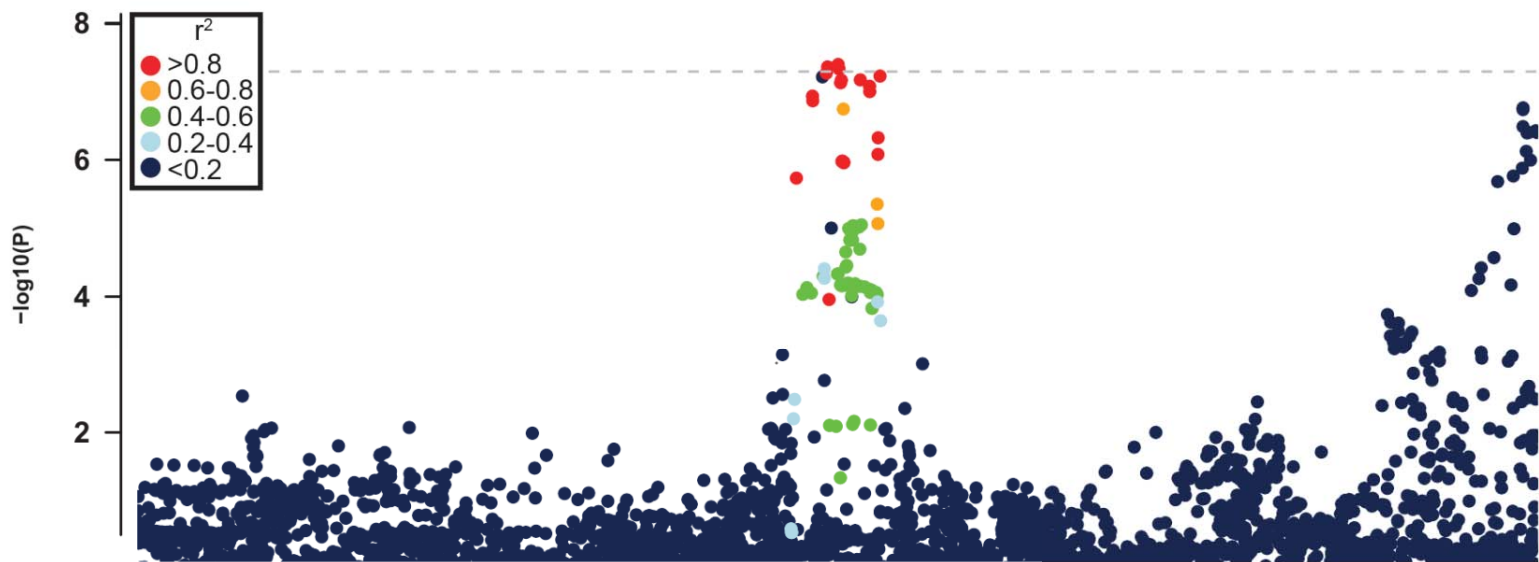
CP>GZ
GZ>CP

Corrected for total SA

Uncorrected

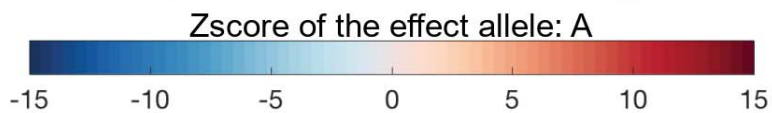
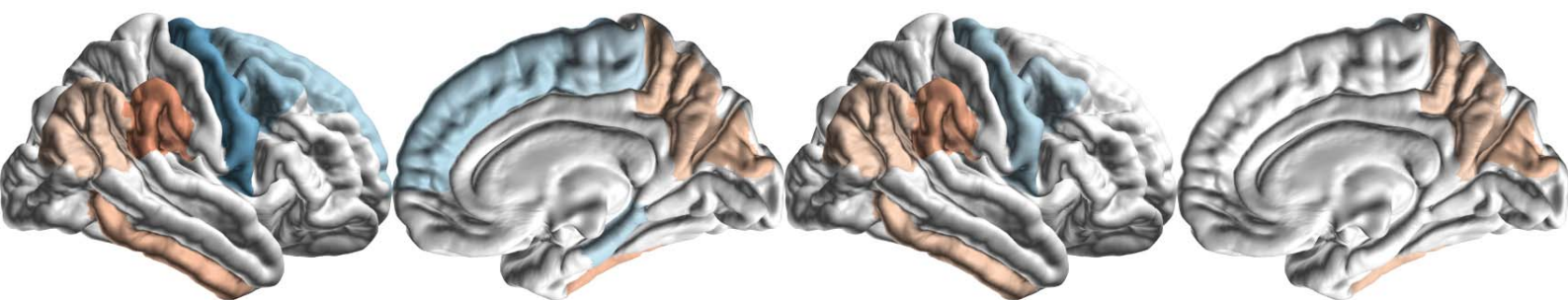


Supramarginal SA: rs62007727



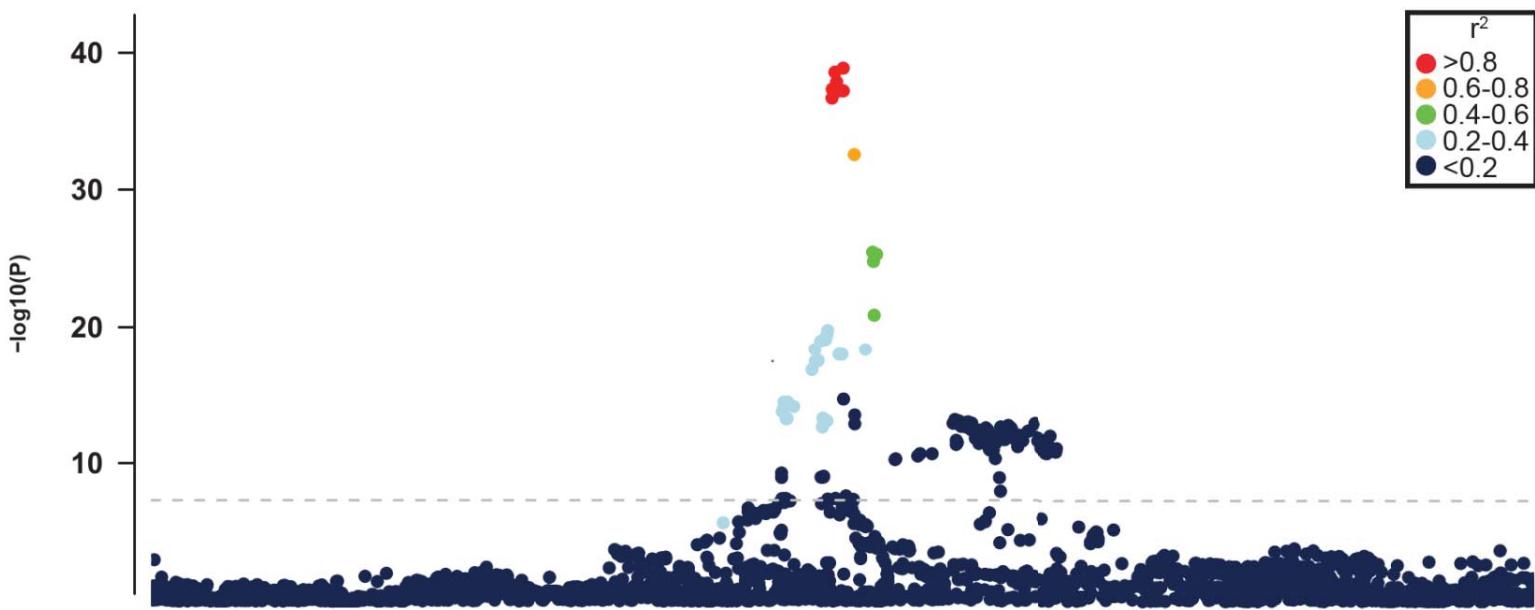
Corrected for total SA

Uncorrected



Supramarginal SA: rs4924345

Chromosome 15



C15orf54

THBS1



GZ>CP

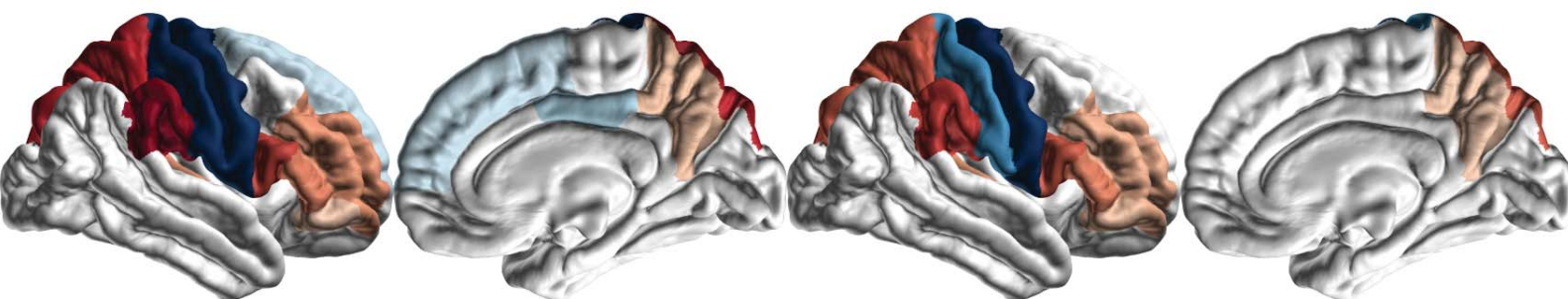


CP>GZ

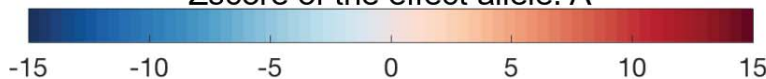


Corrected for total SA

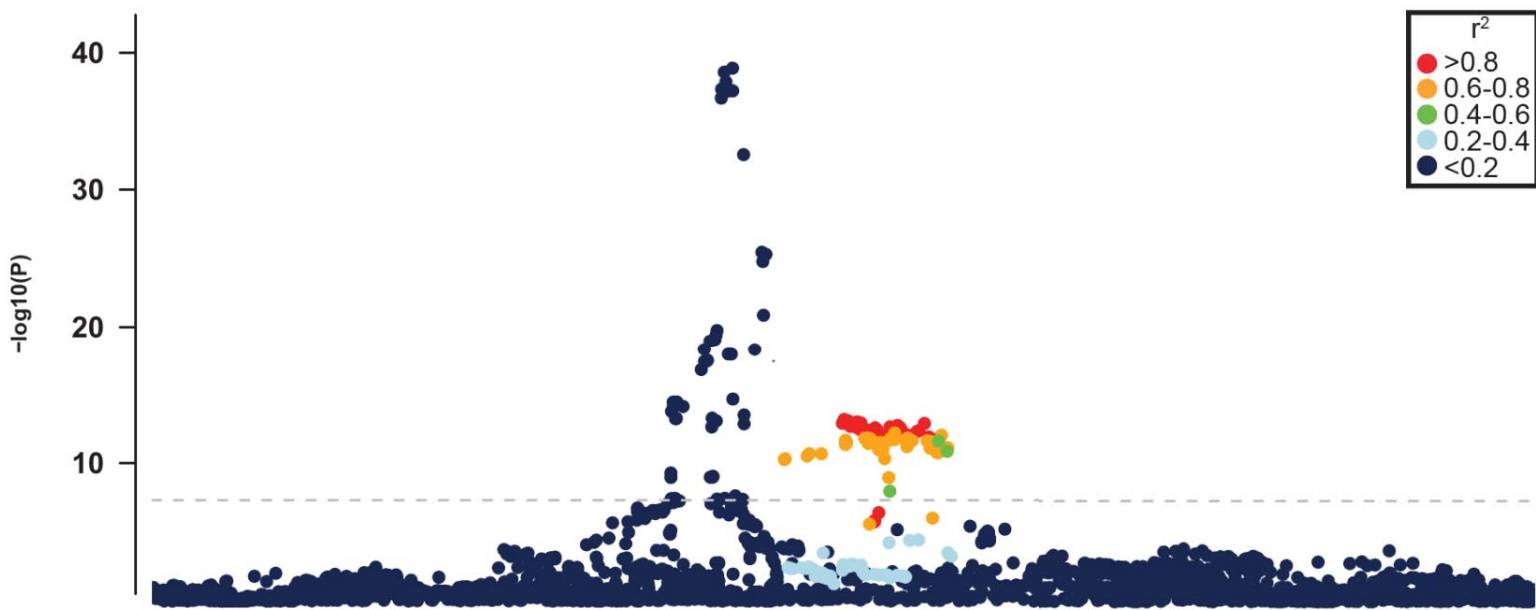
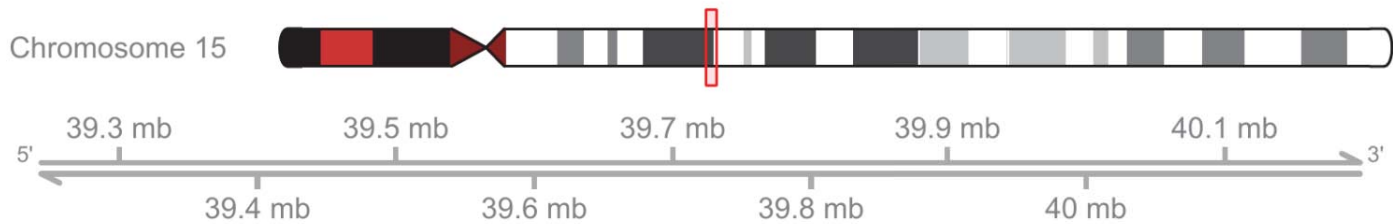
Uncorrected



Zscore of the effect allele: A



Supramarginal SA: rs78502100



C15orf54

THBS1



GZ>CP

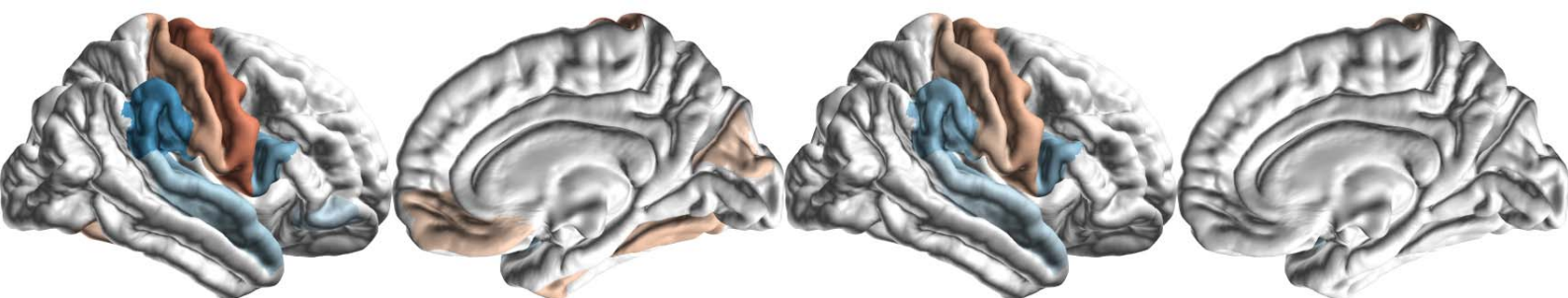


CP>GZ

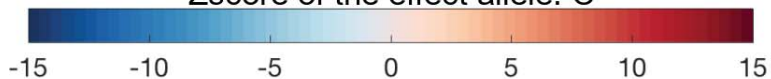


Corrected for total SA

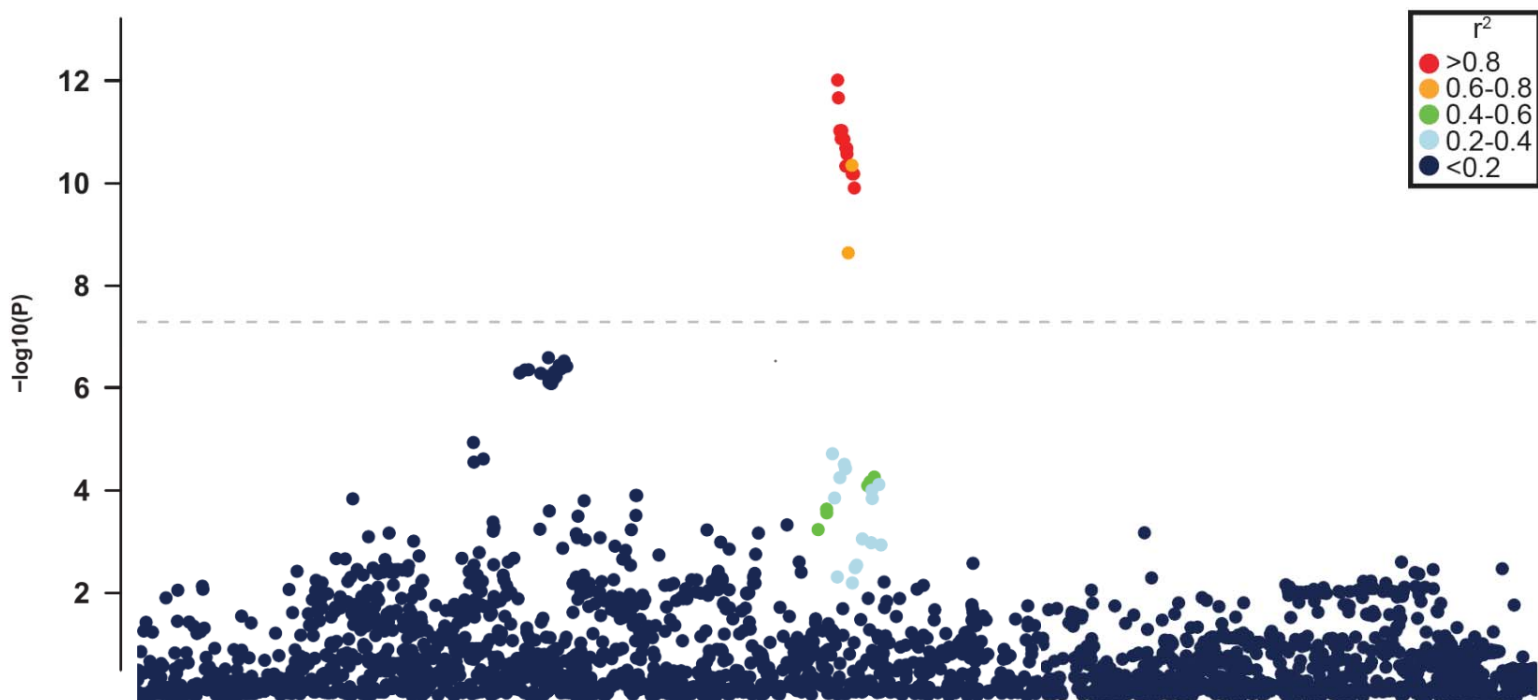
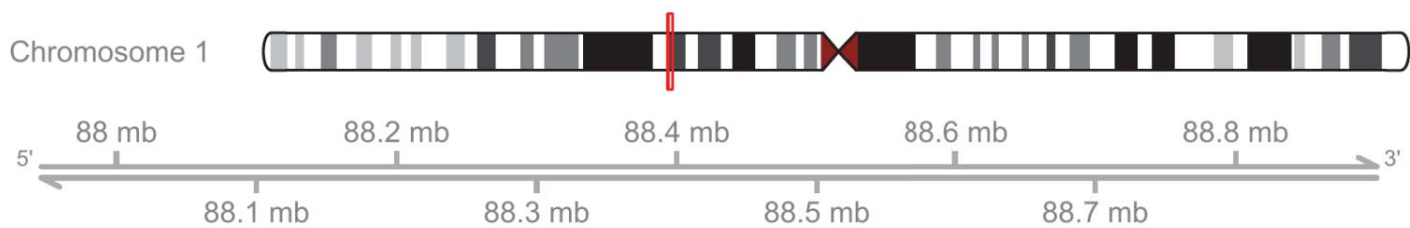
Uncorrected



Zscore of the effect allele: C



Inferior Parietal SA: rs1413536



GZ>CP

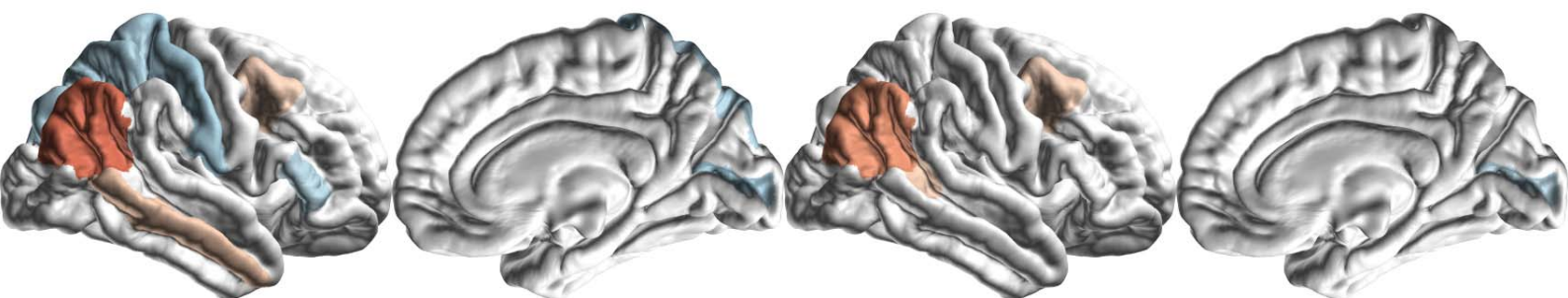


CP>GZ

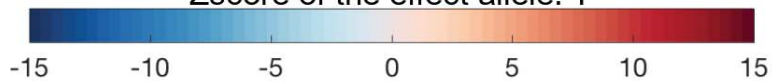


Corrected for total SA

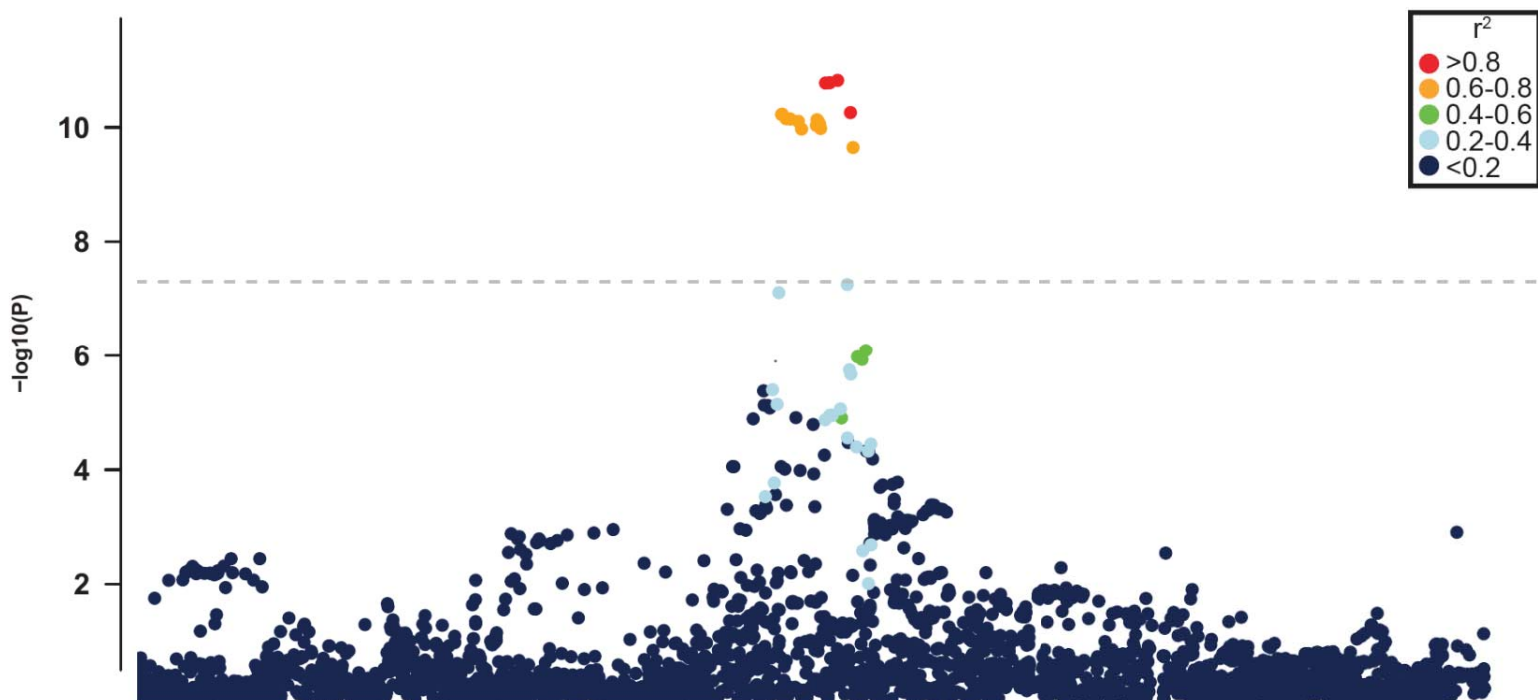
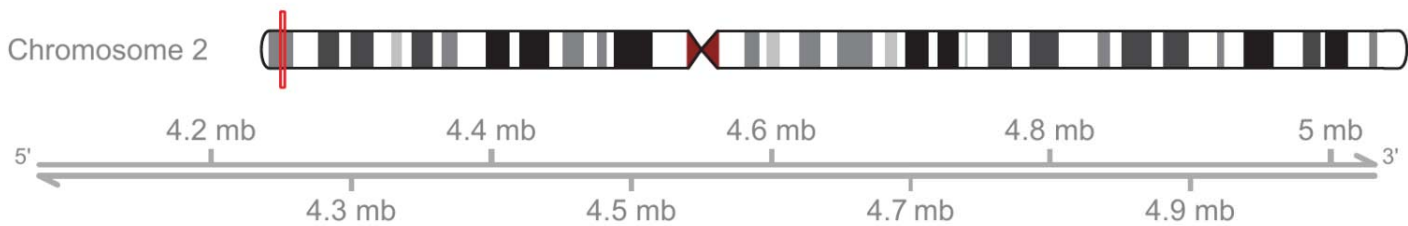
Uncorrected



Zscore of the effect allele: T



Inferior Parietal SA: rs639016

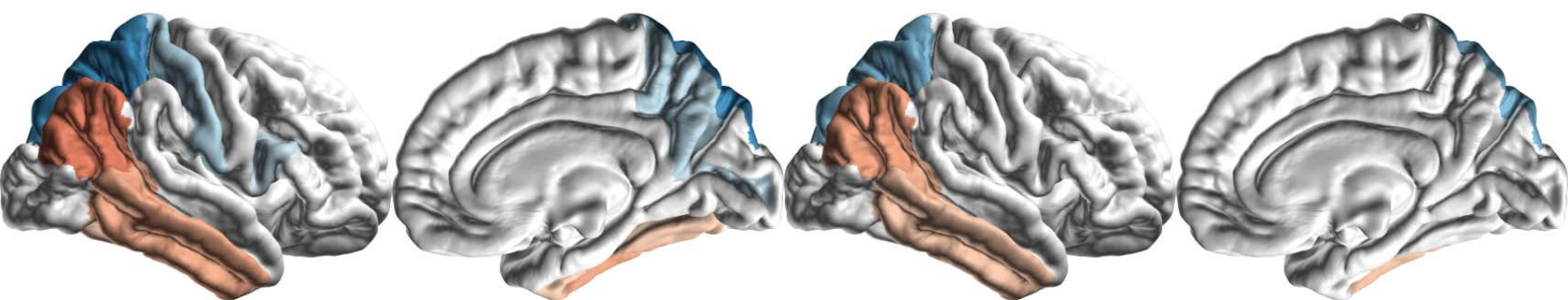


GZ>CP

CP>GZ

Corrected for total SA

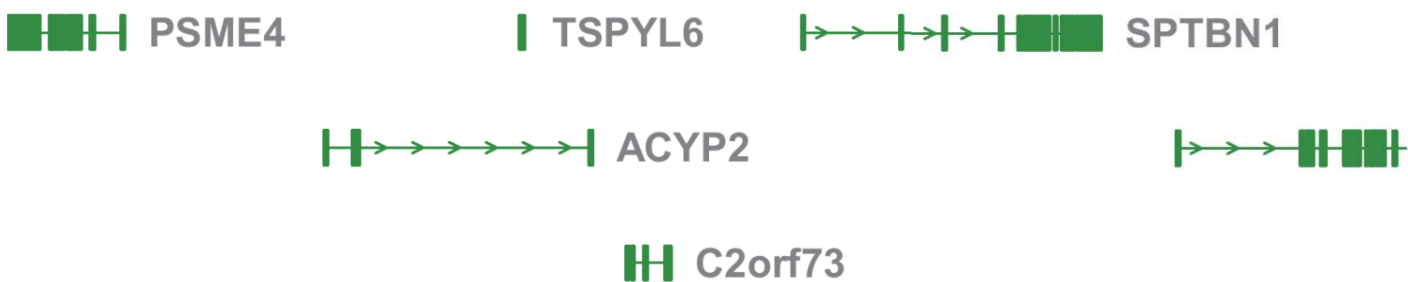
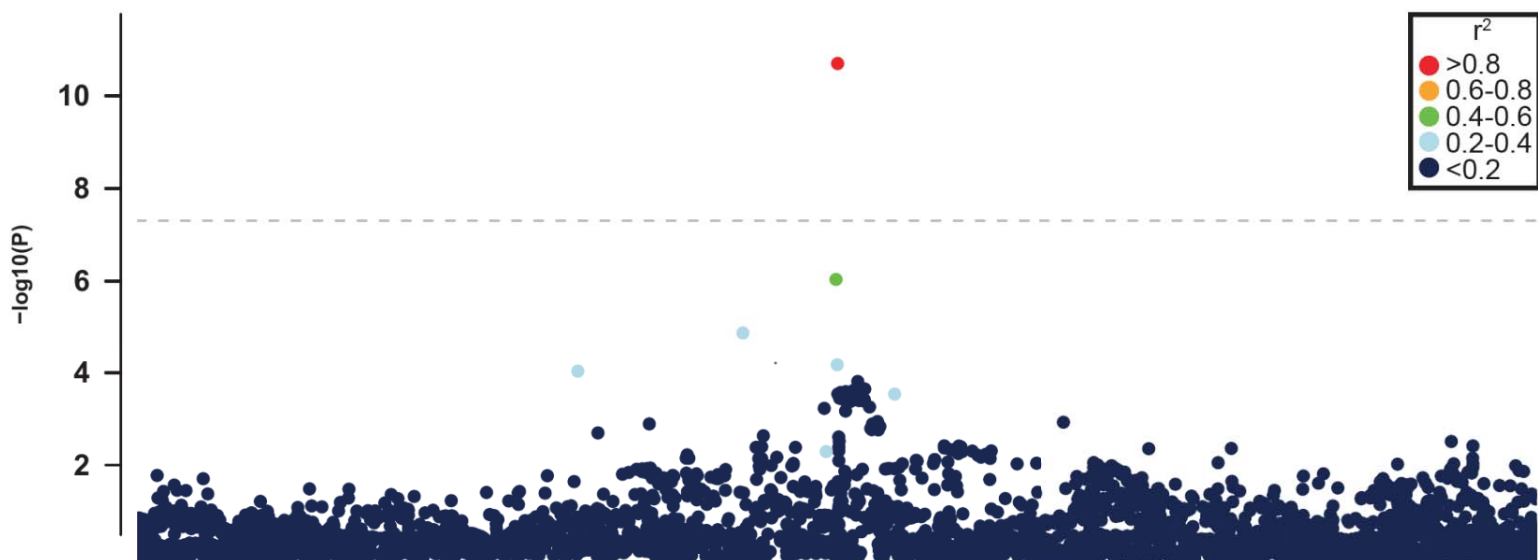
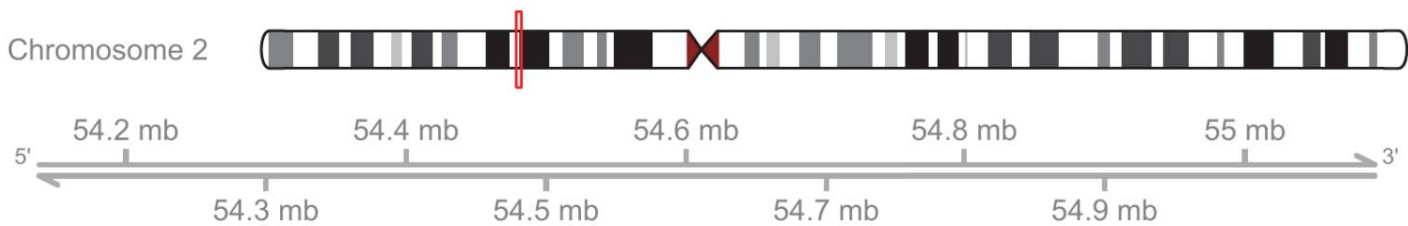
Uncorrected



Zscore of the effect allele: C

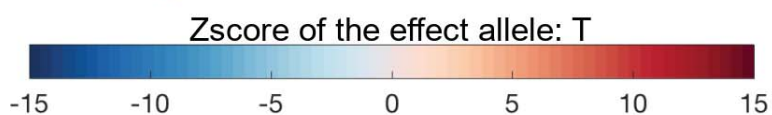
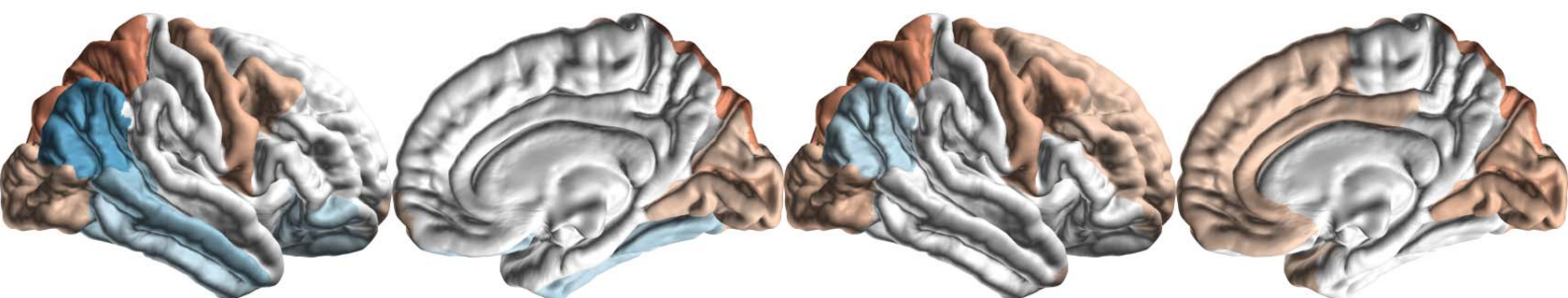


Inferior Parietal SA: rs79272390

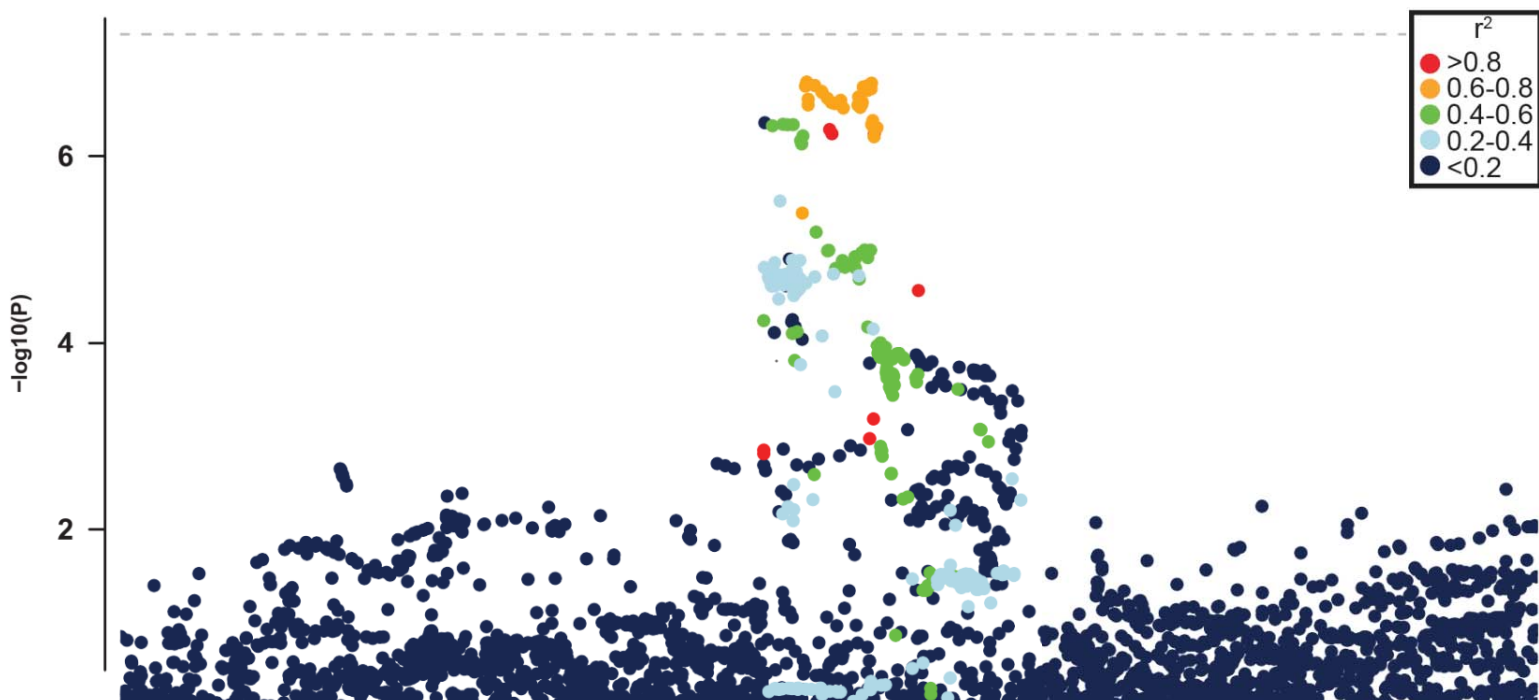
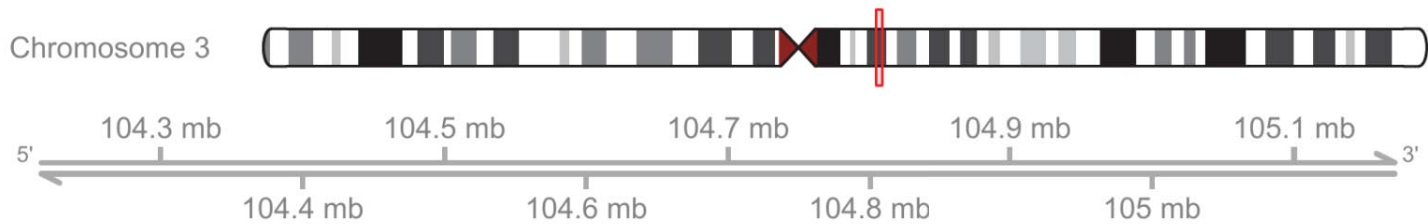


Corrected for total SA

Uncorrected



Inferior Parietal SA: rs13319705

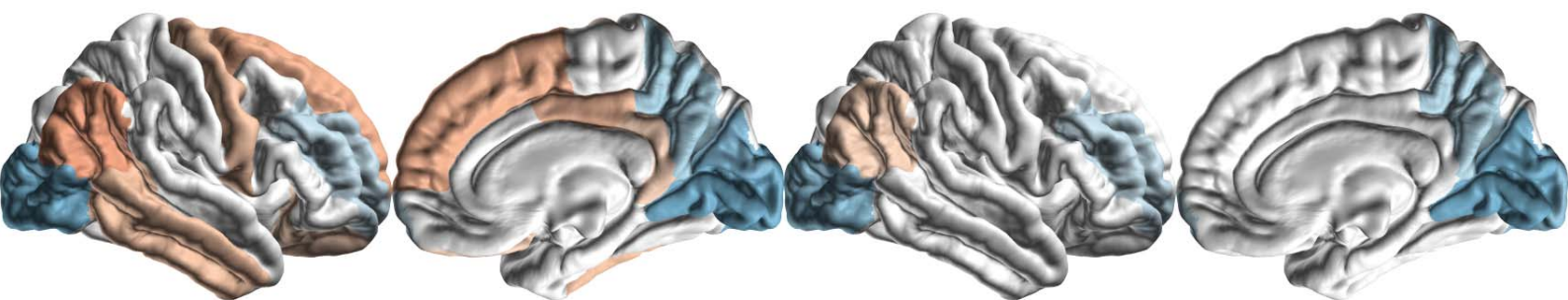


GZ>CP

CP>GZ

Corrected for total SA

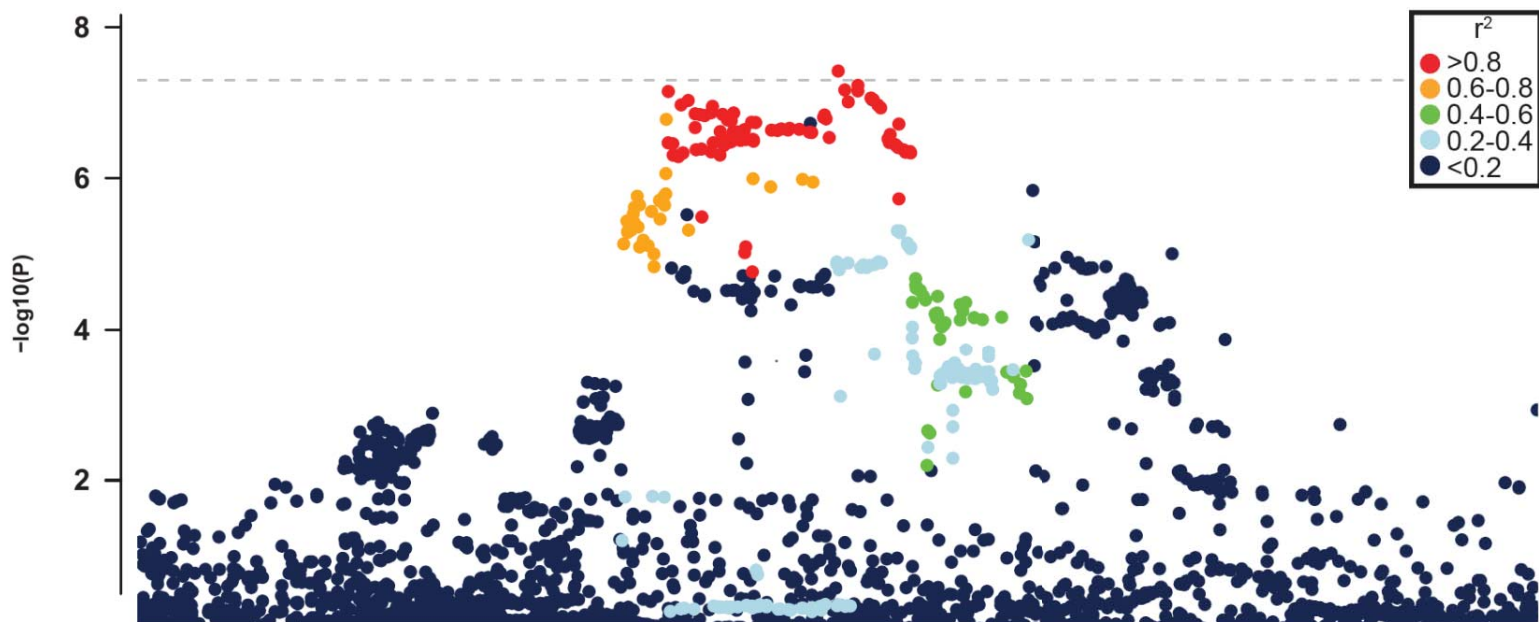
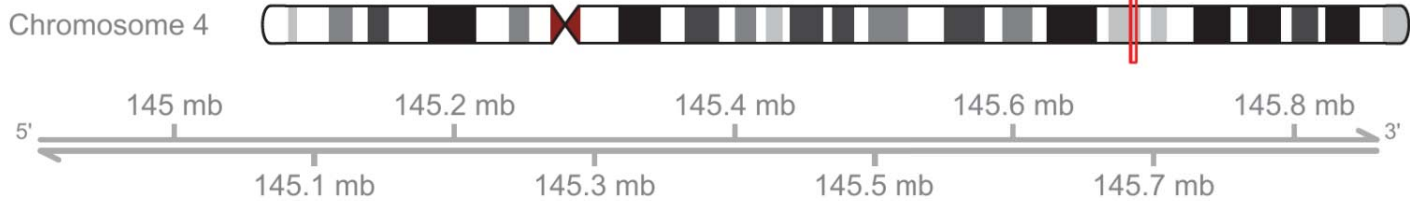
Uncorrected



Zscore of the effect allele: T



Inferior Parietal SA: rs17019370



GYPB

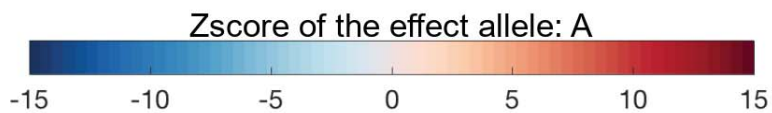
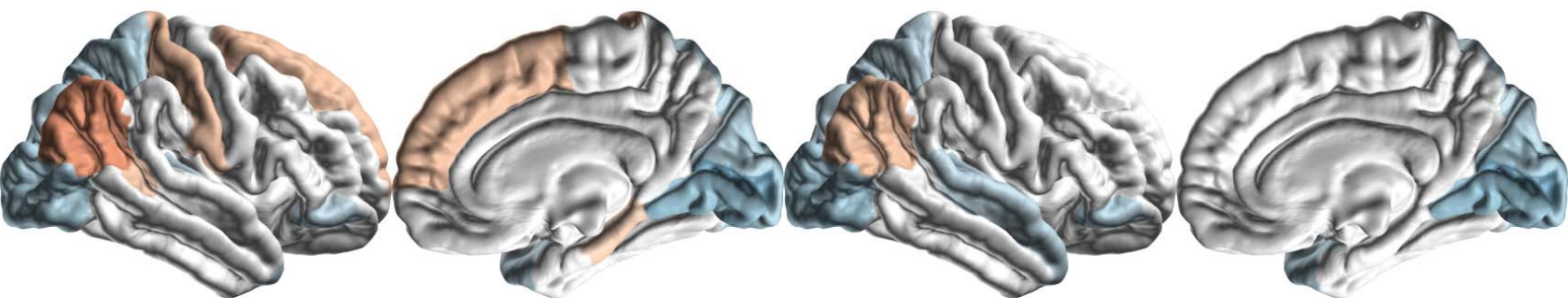
HHIP

GYPB

GZ>CP
CP>GZ

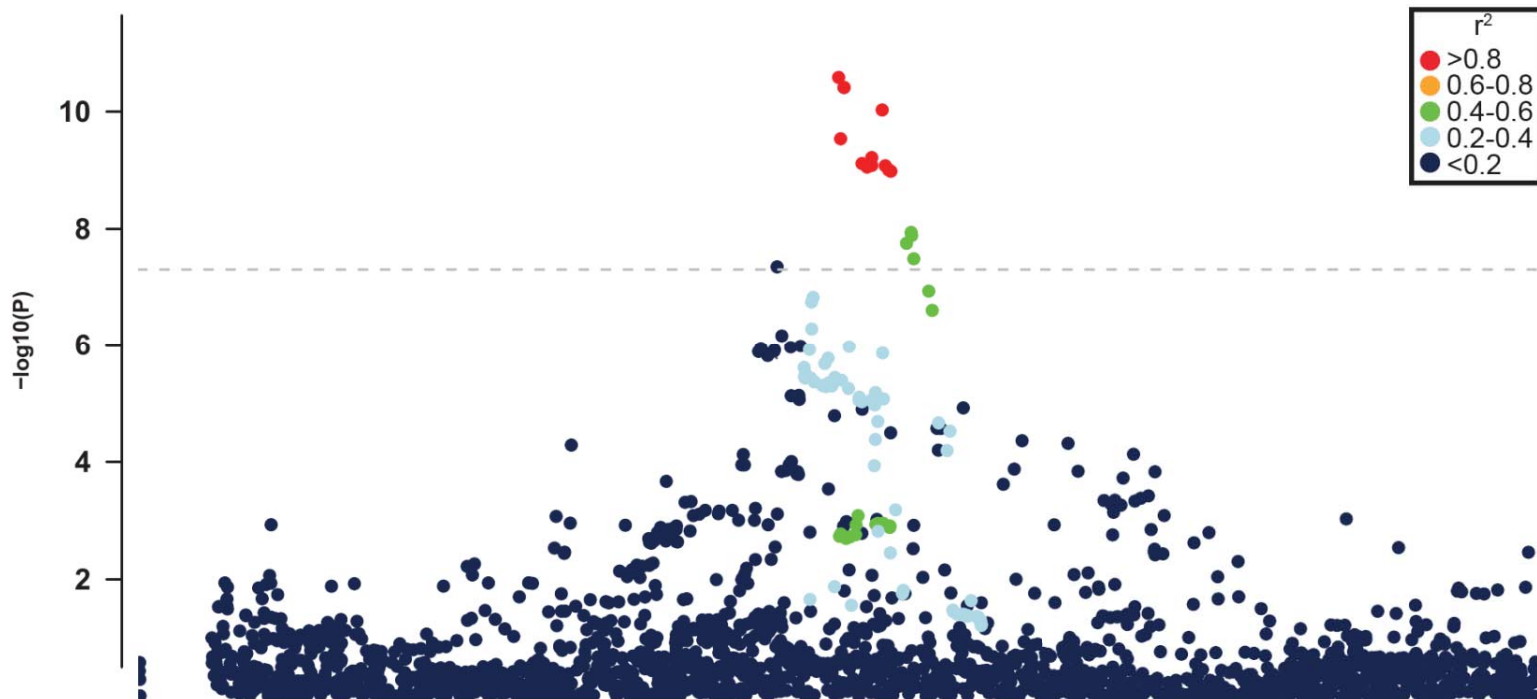
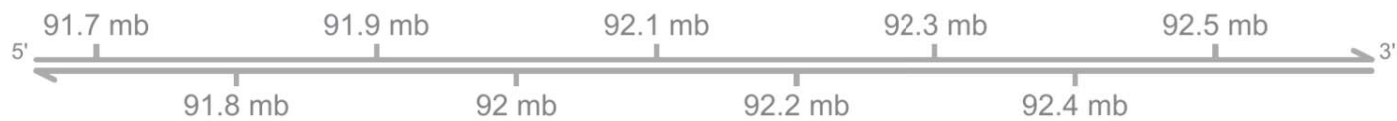
Corrected for total SA

Uncorrected



Inferior Parietal SA: rs27540

Chromosome 5



GZ>CP

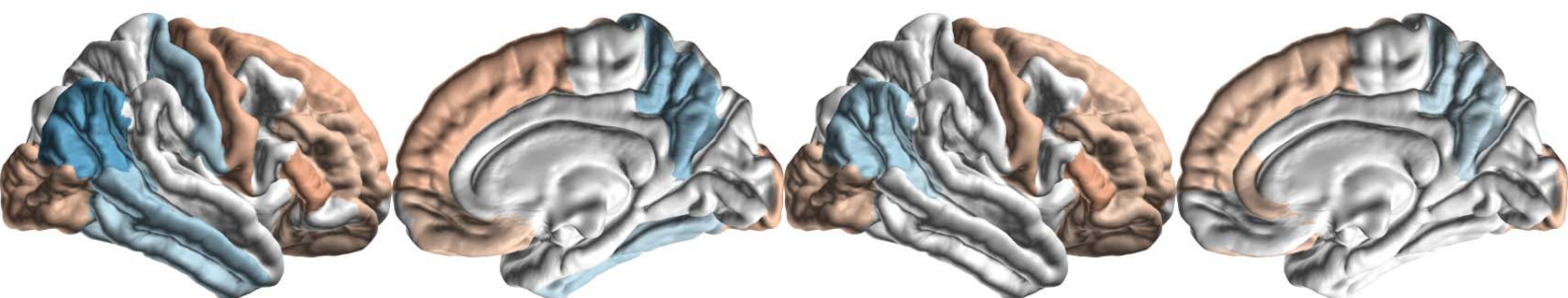


CP>GZ

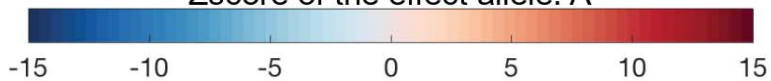


Corrected for total SA

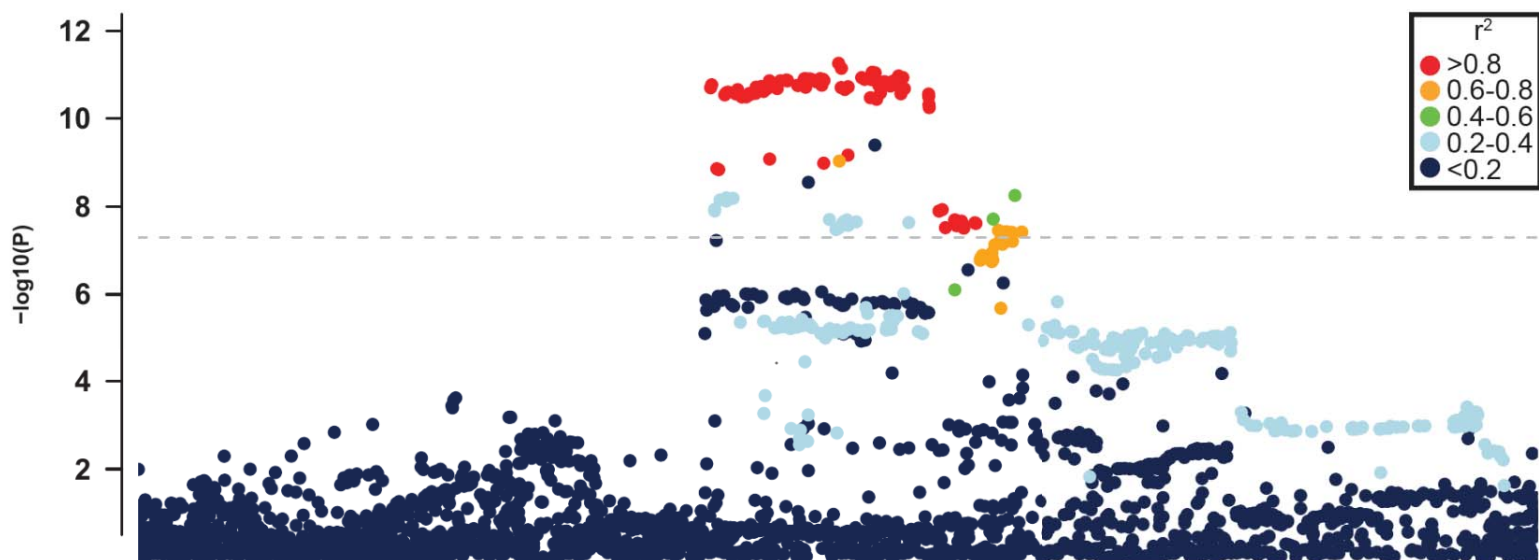
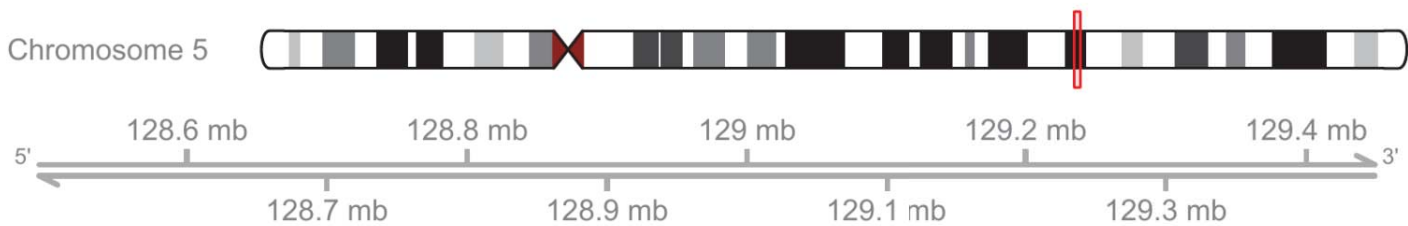
Uncorrected



Zscore of the effect allele: A



Inferior Parietal SA: rs62399042



ADAMTS19

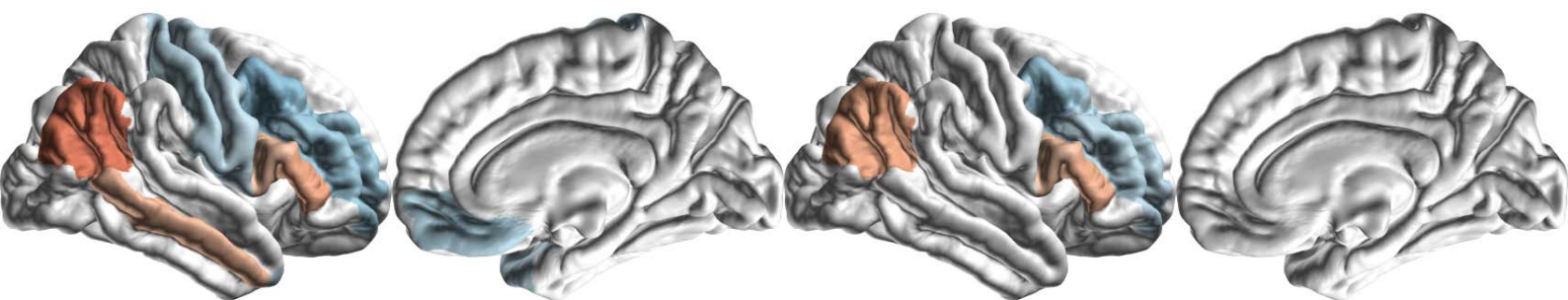
KIAA1024L

CP>GZ

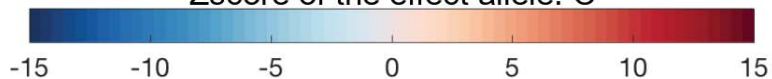
GZ>CP

Corrected for total SA

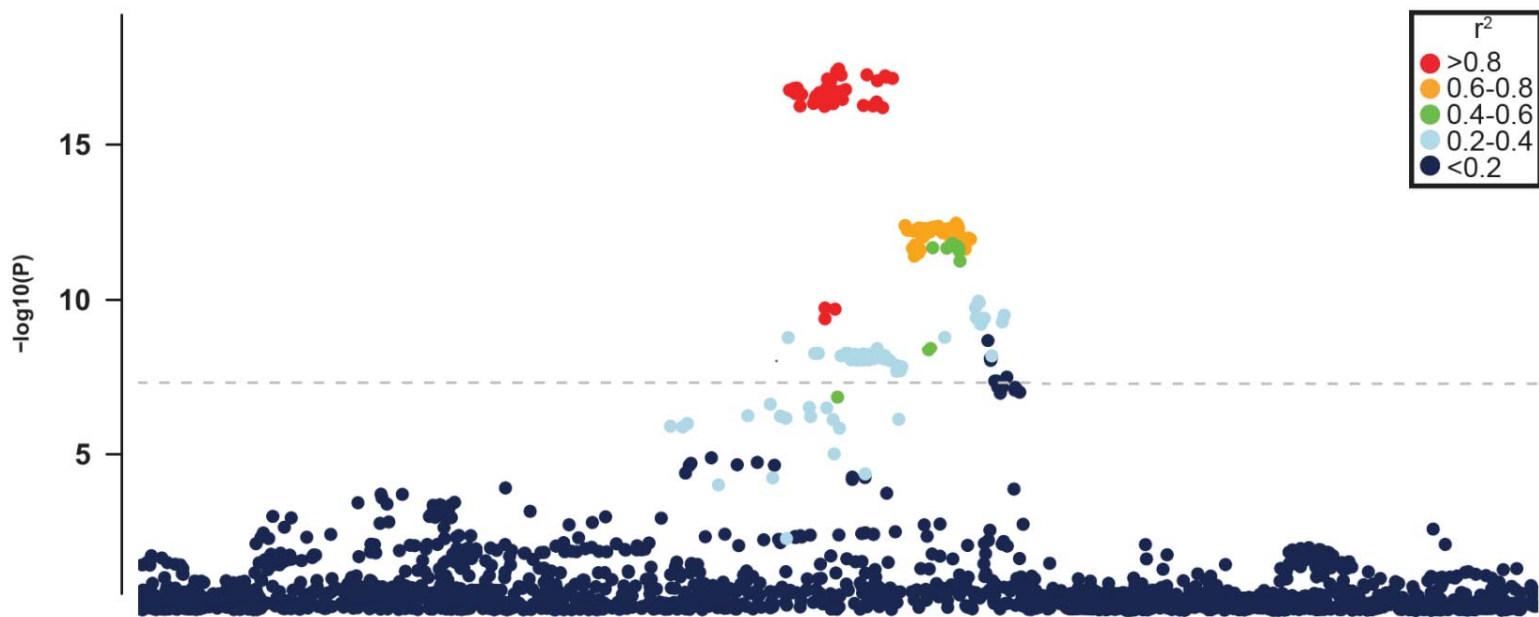
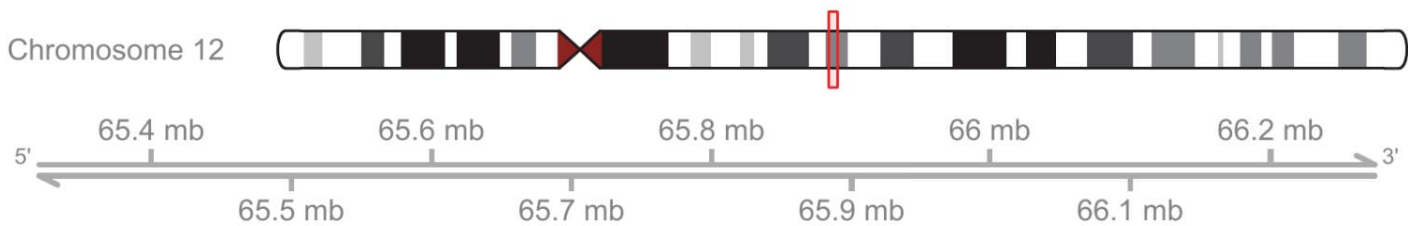
Uncorrected



Zscore of the effect allele: C

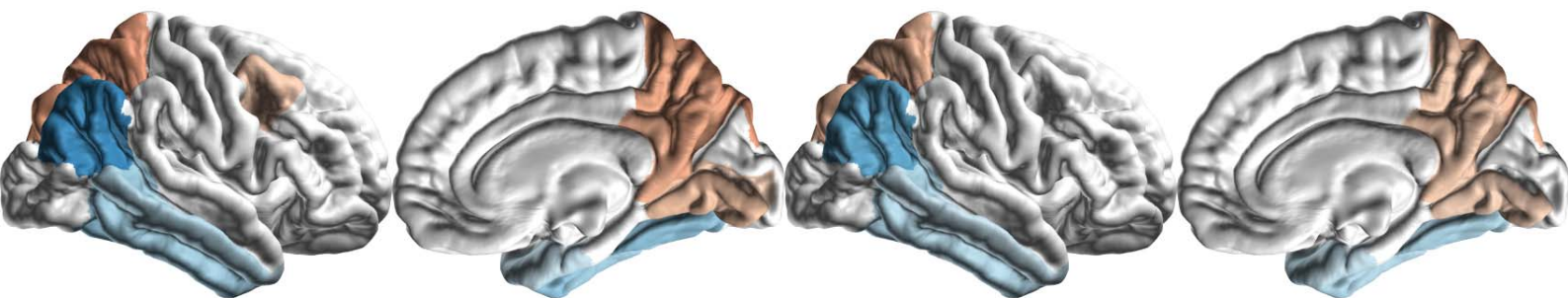


Inferior Parietal SA: rs2336714



Corrected for total SA

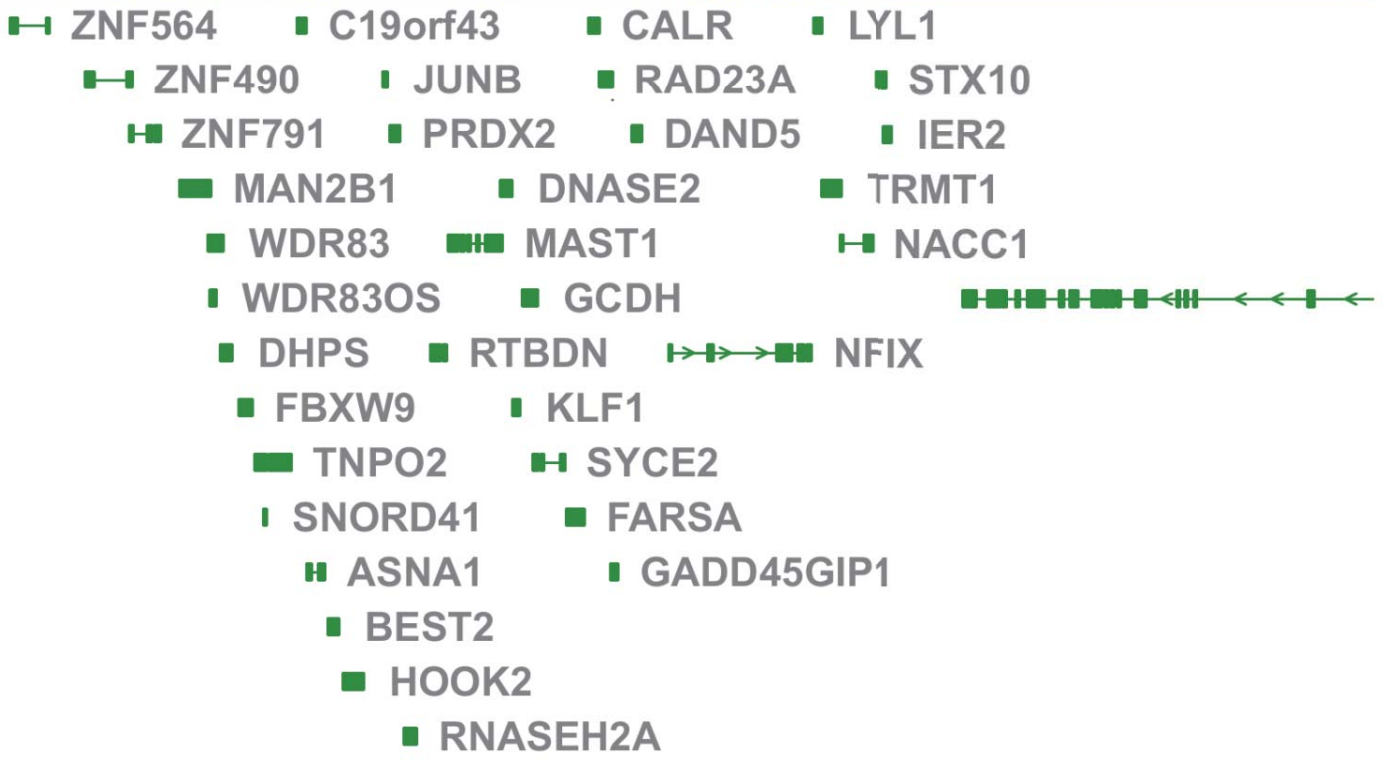
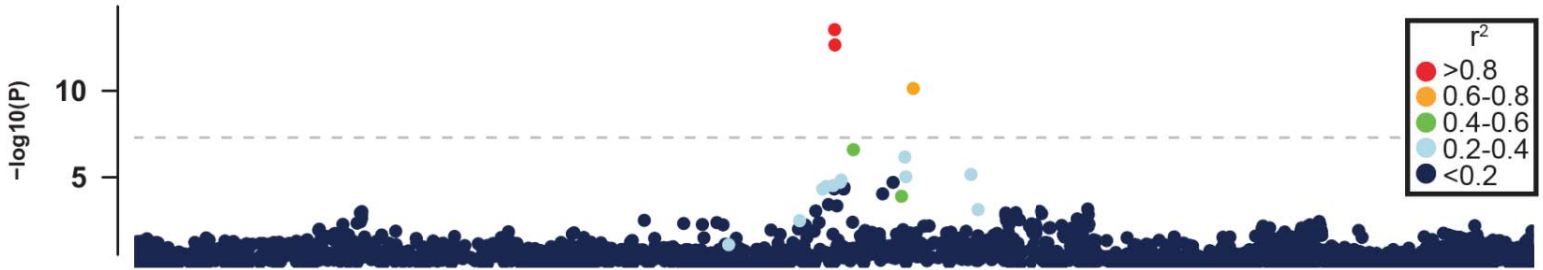
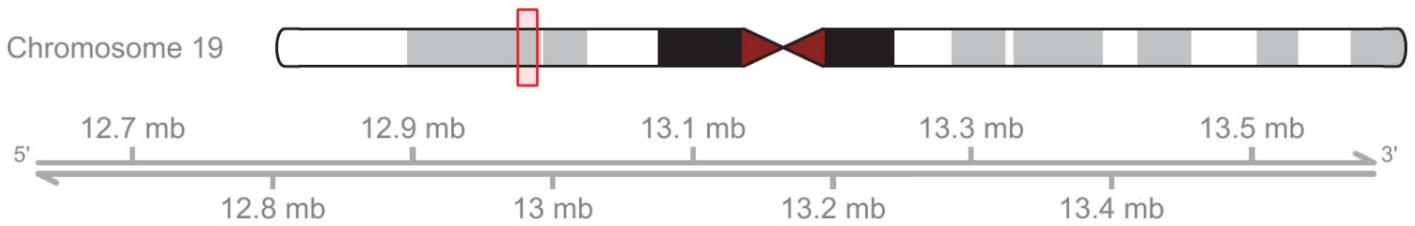
Uncorrected



Zscore of the effect allele: T



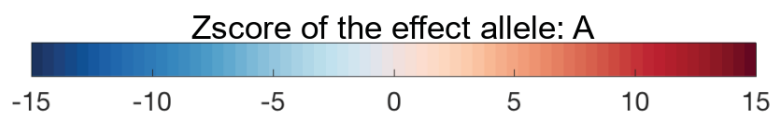
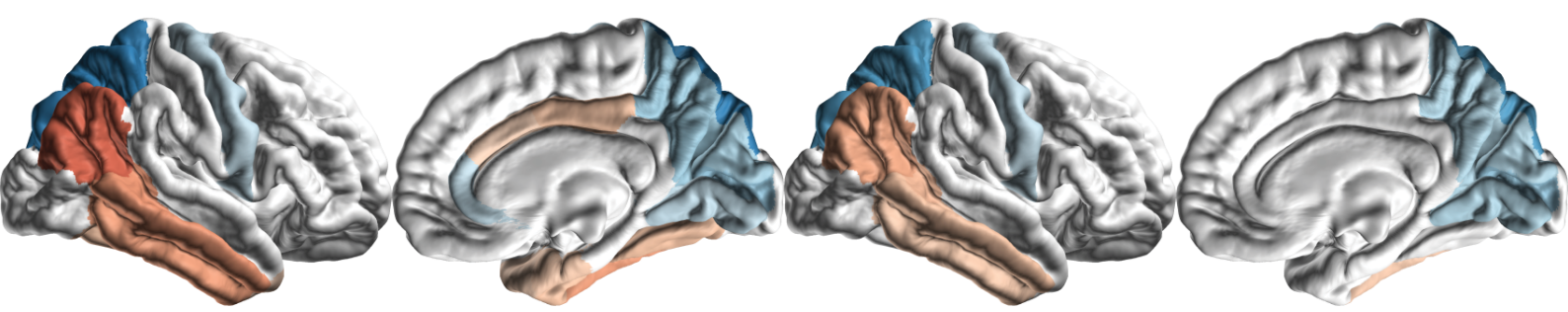
Inferior Parietal SA: rs68175985



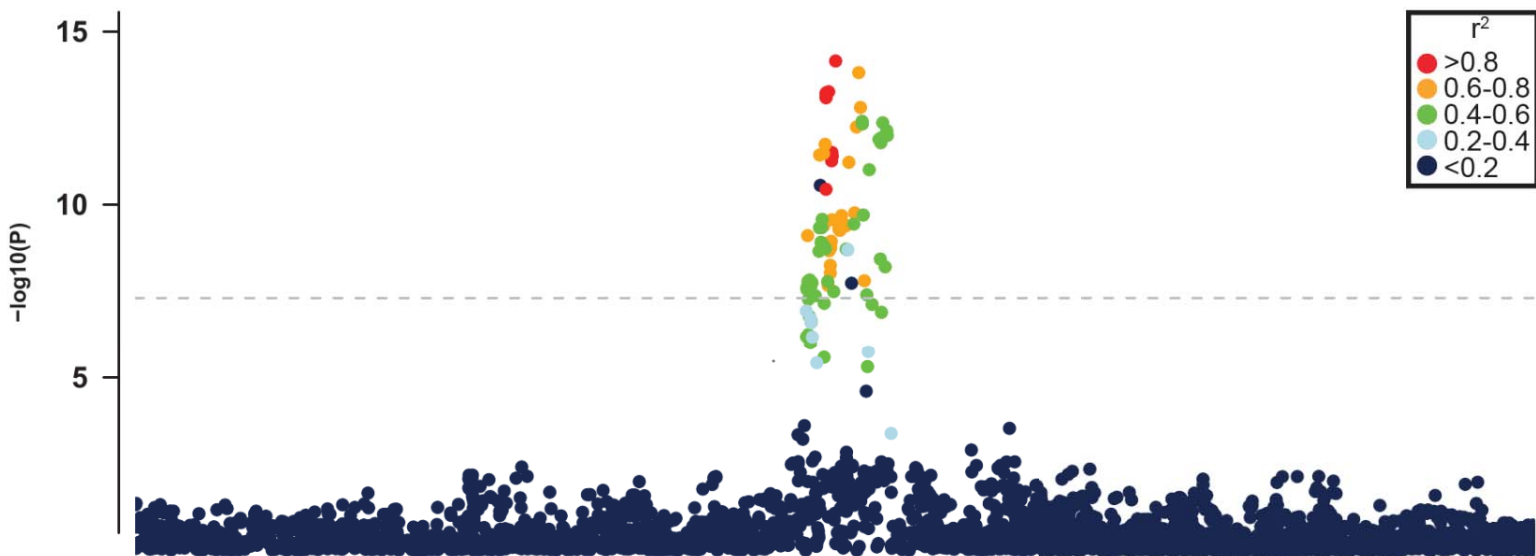
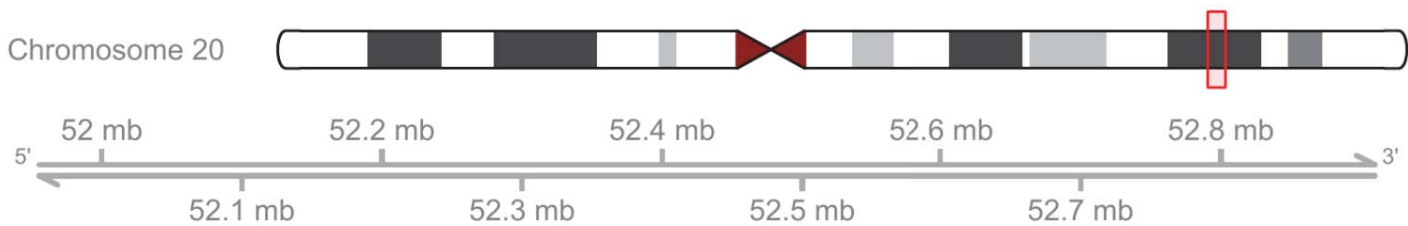
CP>GZZ>CP

Corrected for total SA

Uncorrected



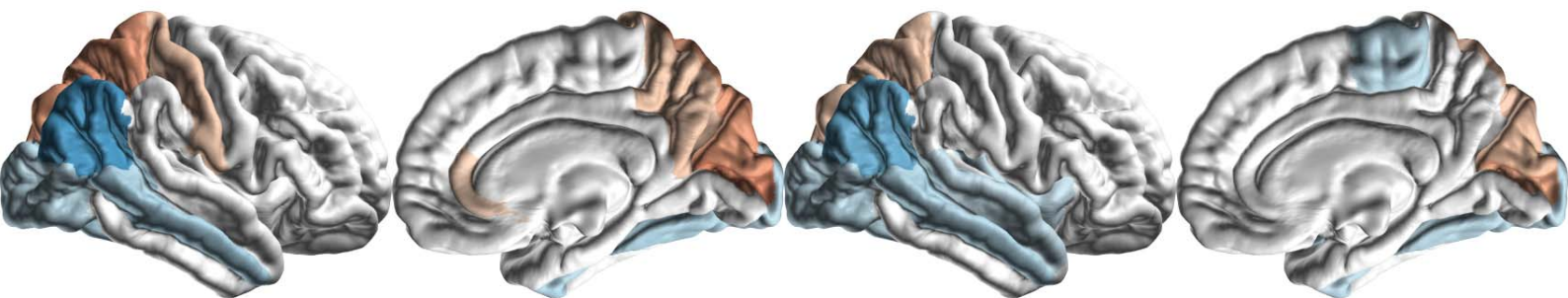
Inferior Parietal SA: rs4437022



CP>GZ
GZ>CP

Corrected for total SA

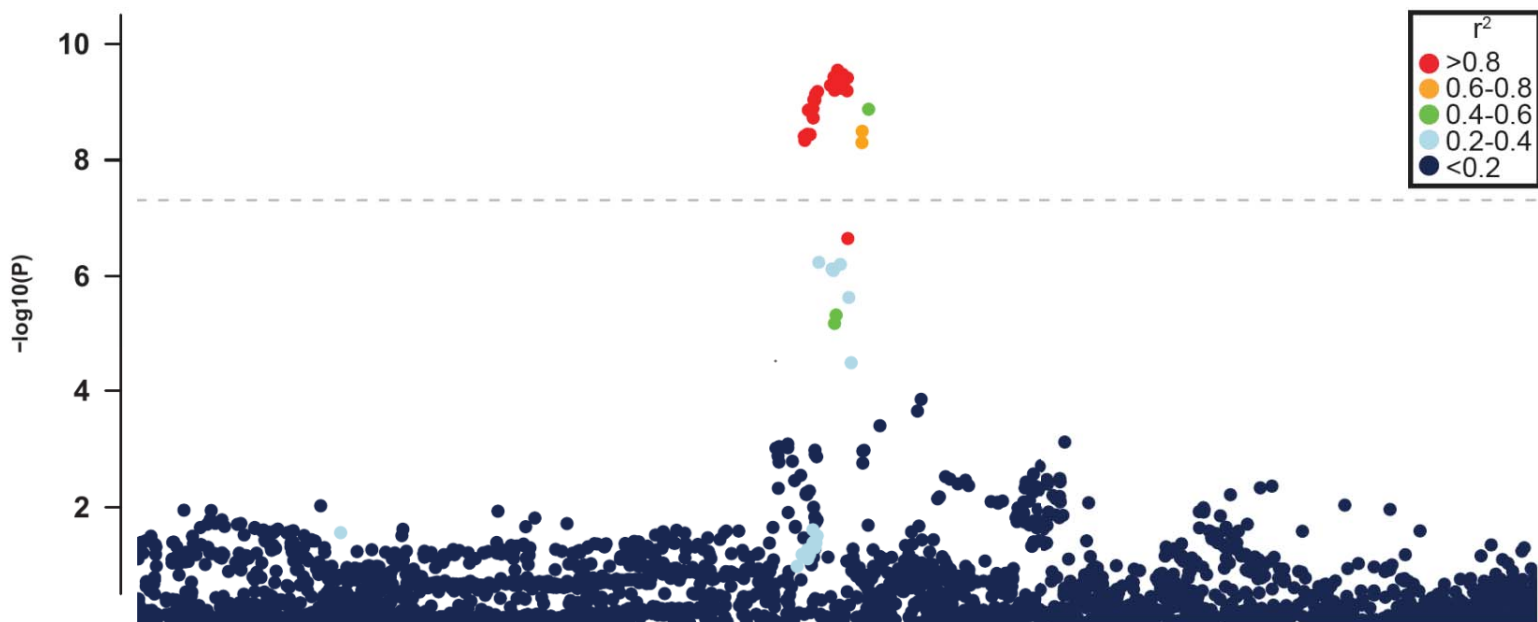
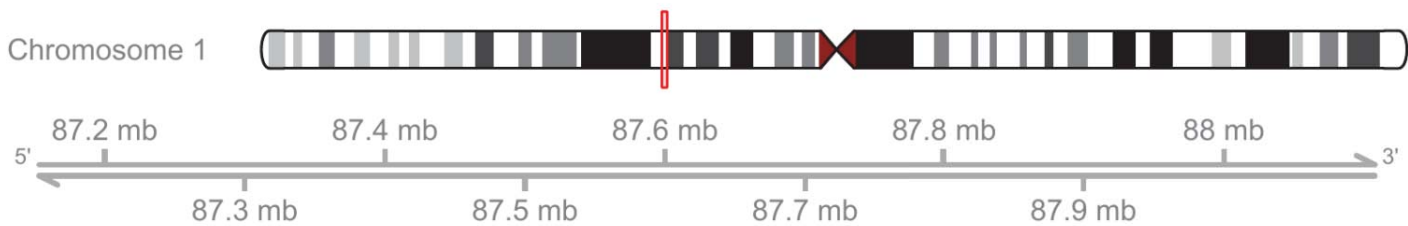
Uncorrected



Zscore of the effect allele: A



Posterior Cingulate SA: rs11161942



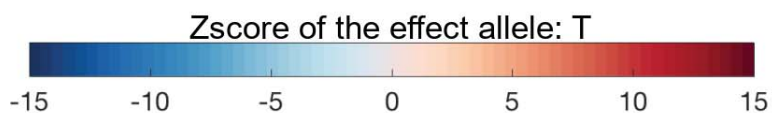
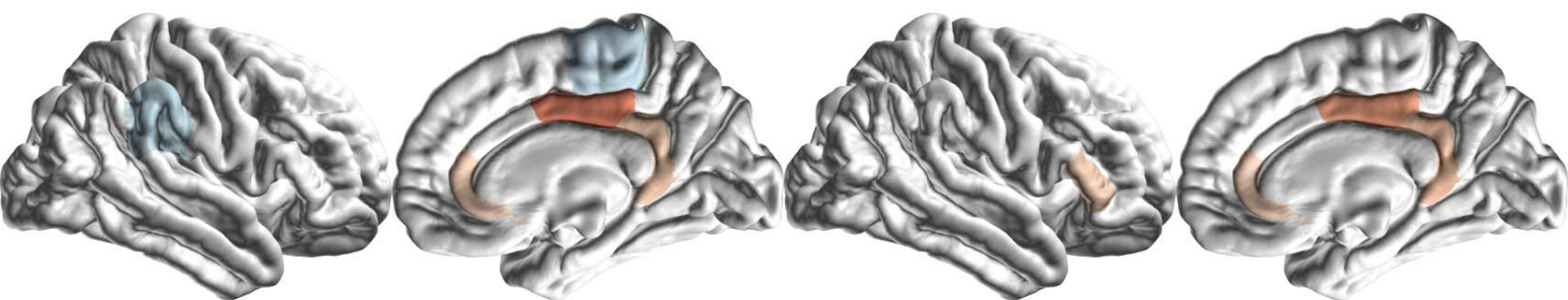
SH3GLB1 HS2ST1 LMO4

SEP15

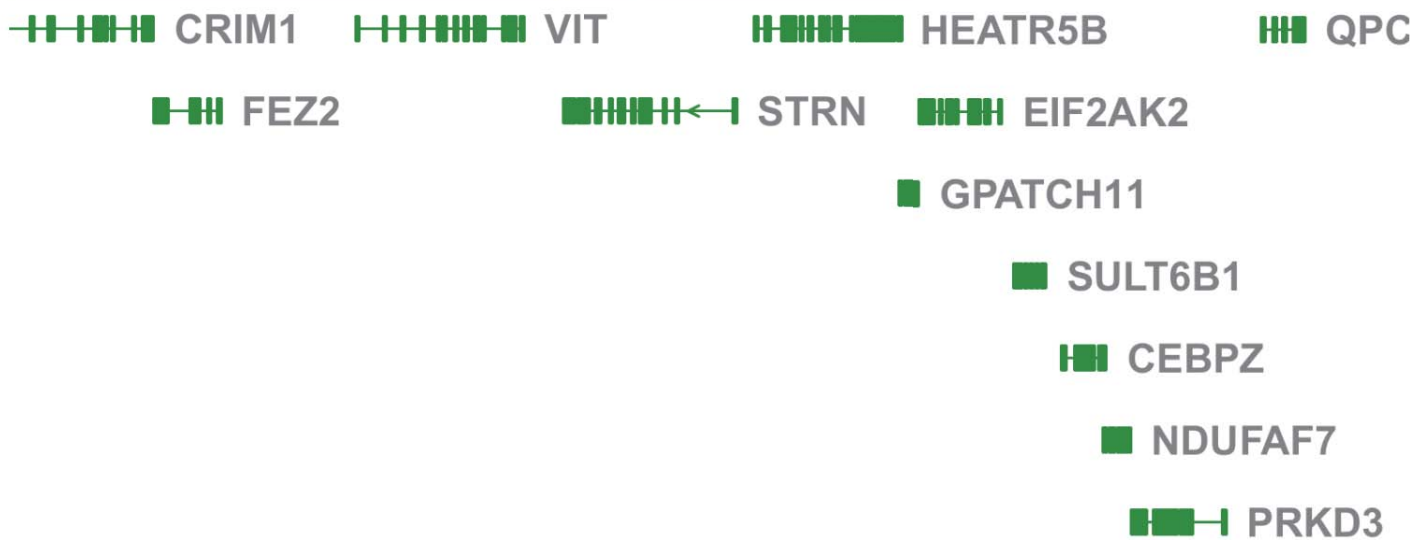
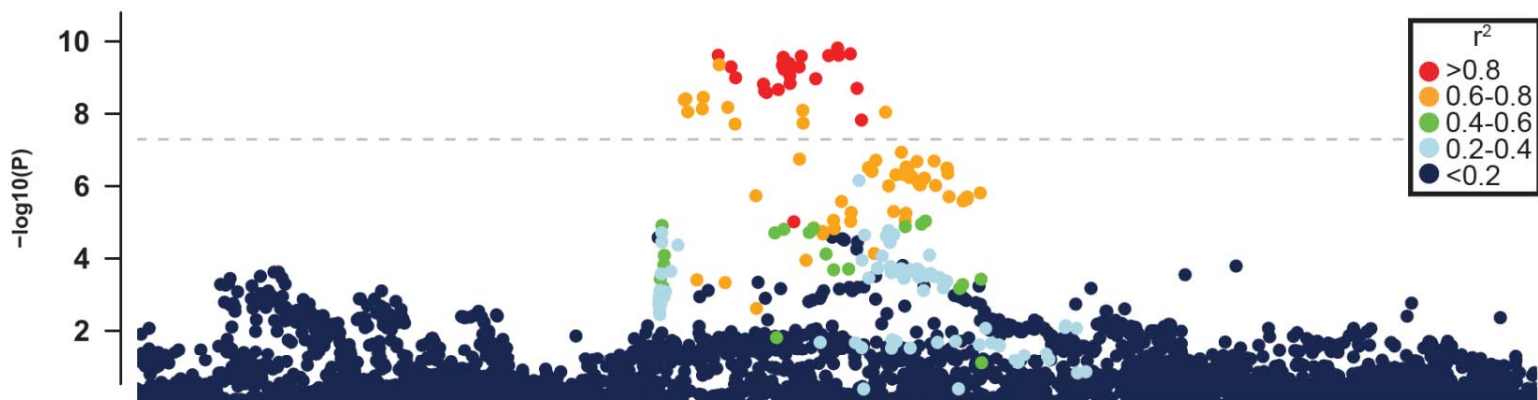
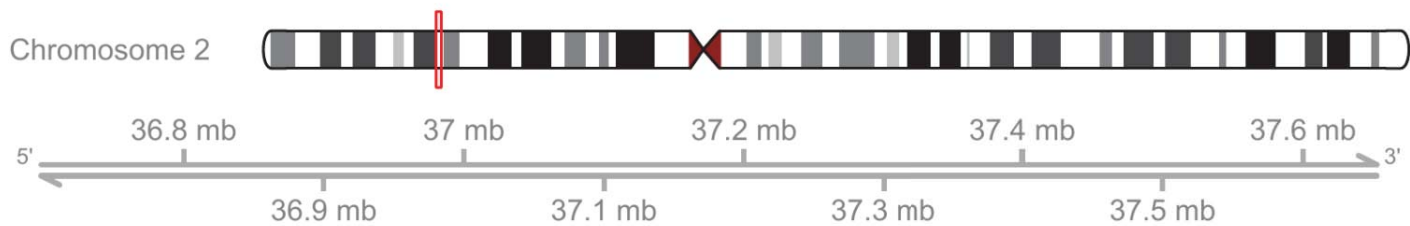


Corrected for total SA

Uncorrected

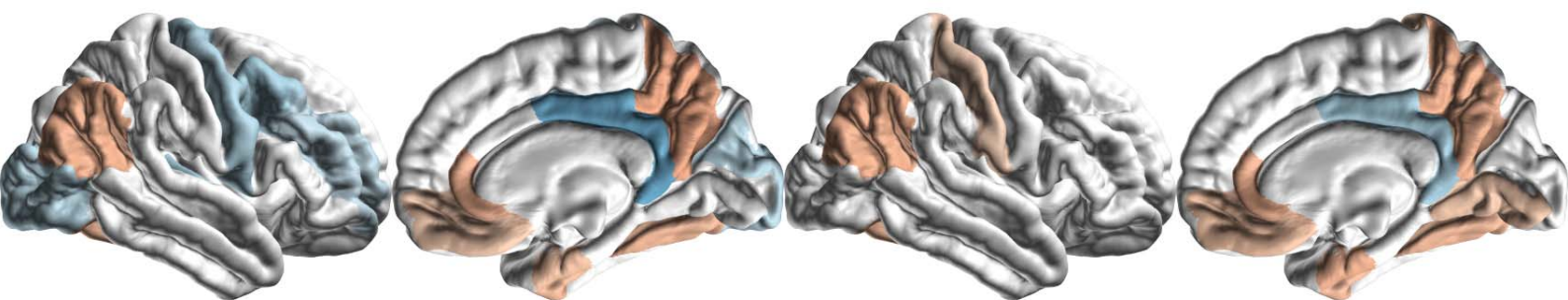


Posterior Cingulate SA: rs11695609



Corrected for total SA

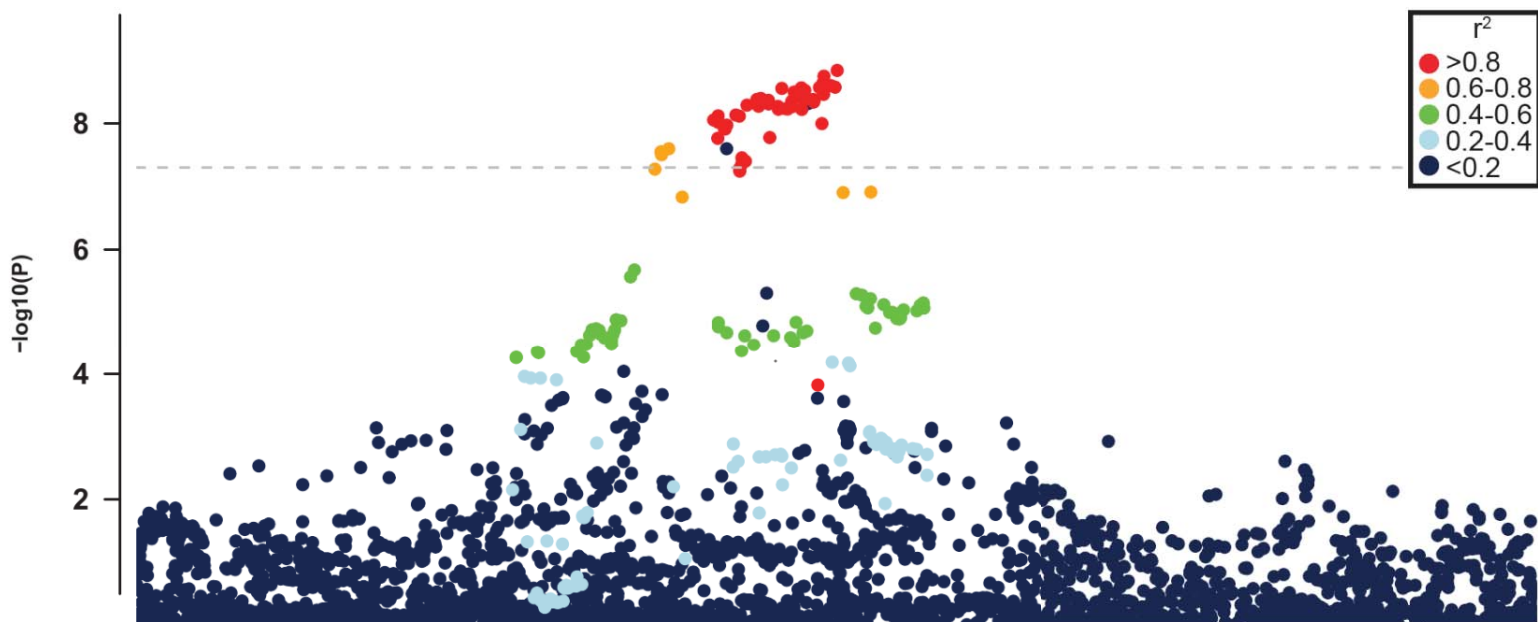
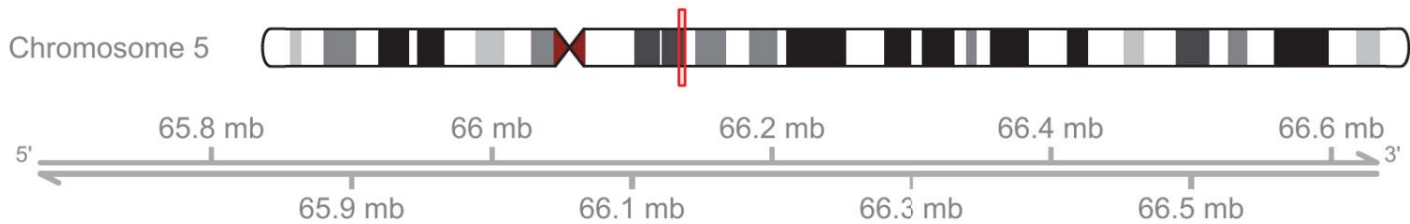
Uncorrected



Zscore of the effect allele: T

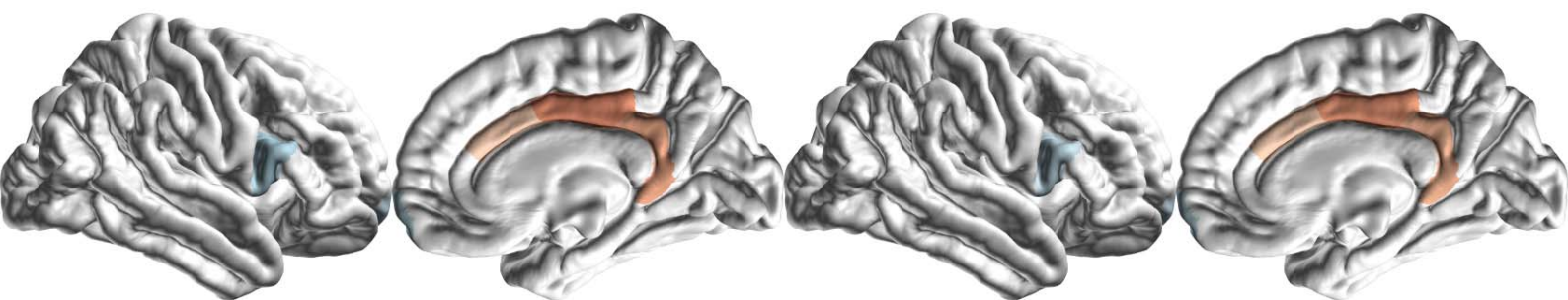


Posterior Cingulate SA: rs72761270



Corrected for total SA

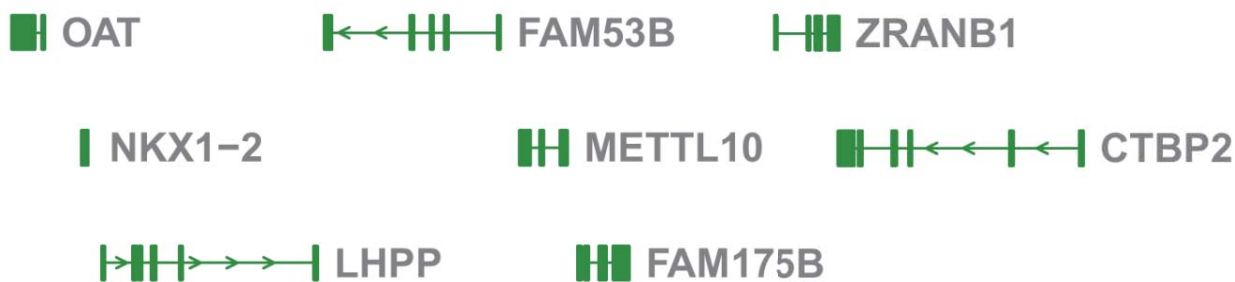
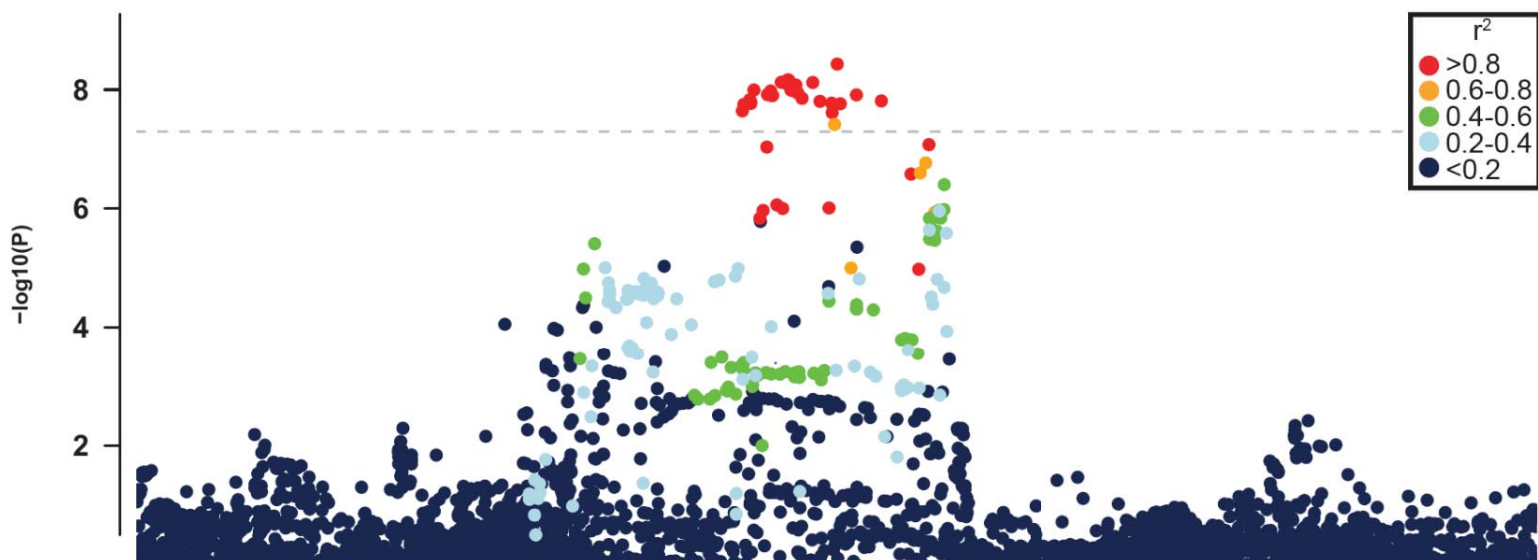
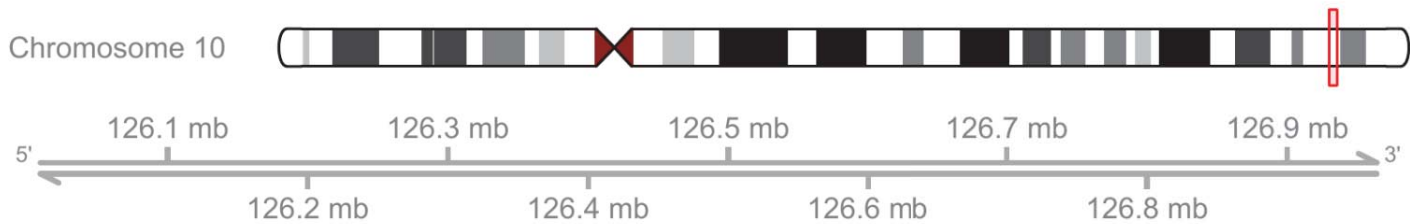
Uncorrected



Zscore of the effect allele: T



Posterior Cingulate SA: rs12764880

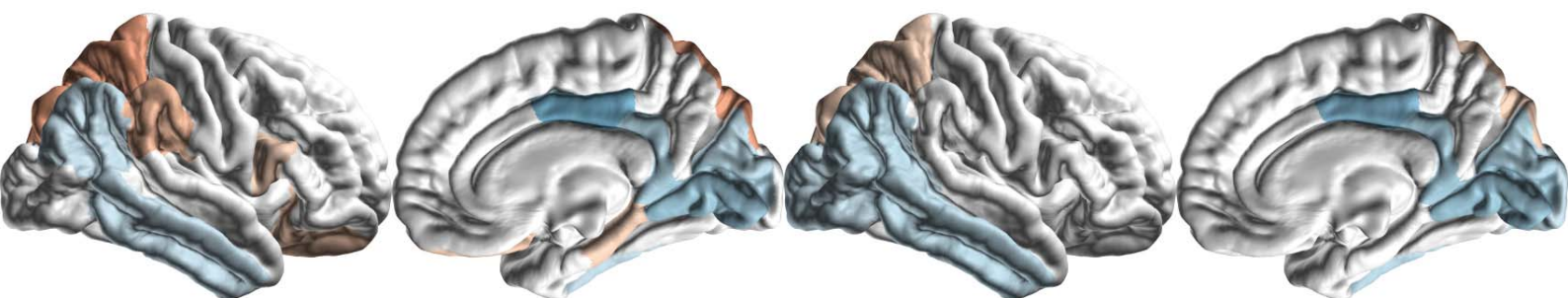


GZ>CP
CP>GZ



Corrected for total SA

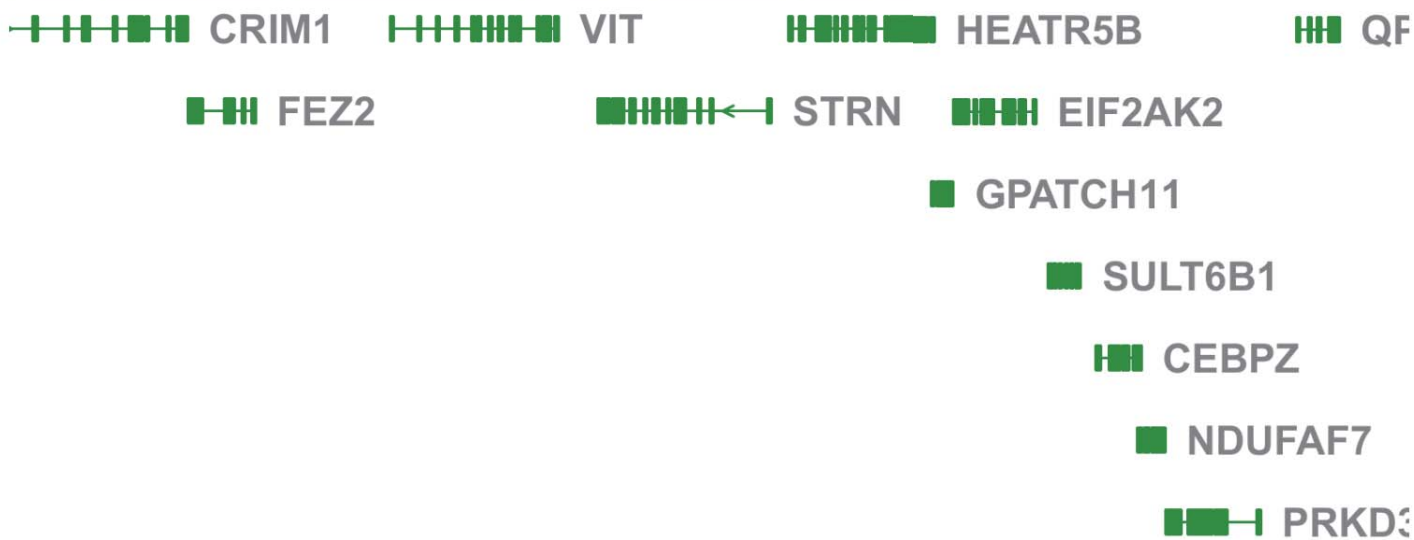
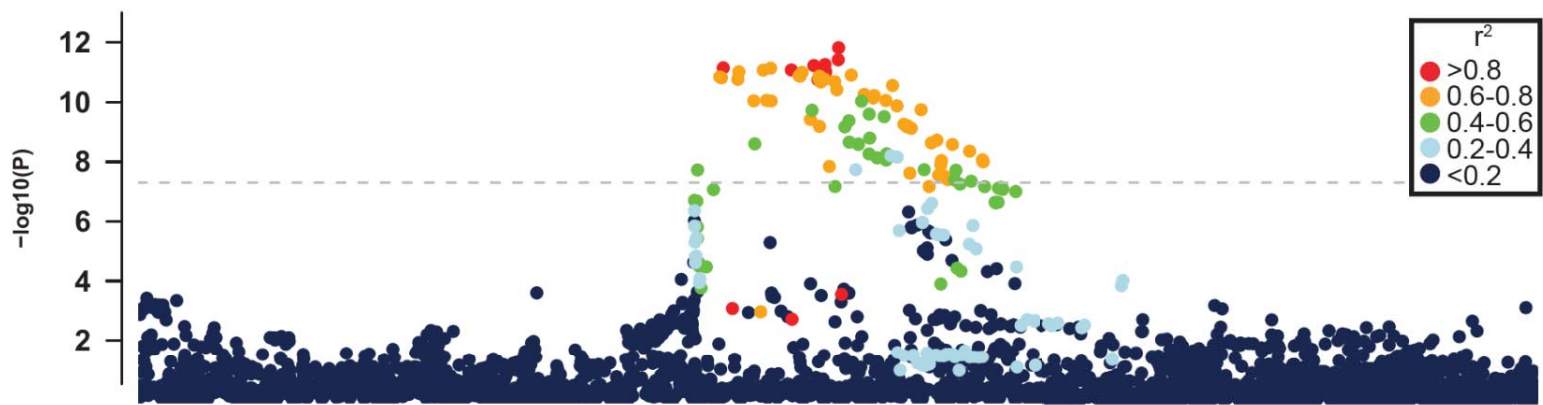
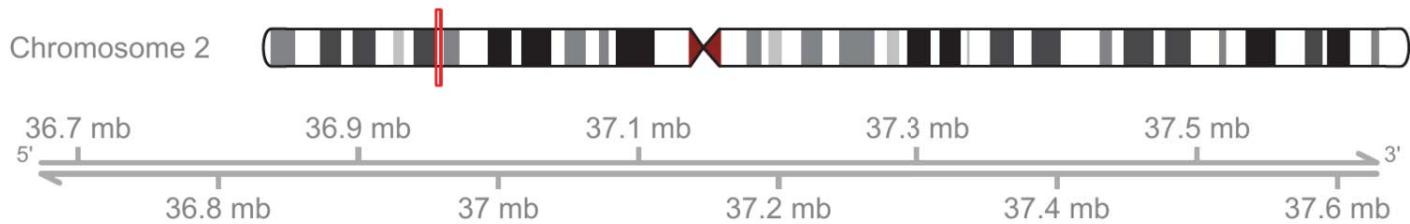
Uncorrected



Zscore of the effect allele: T

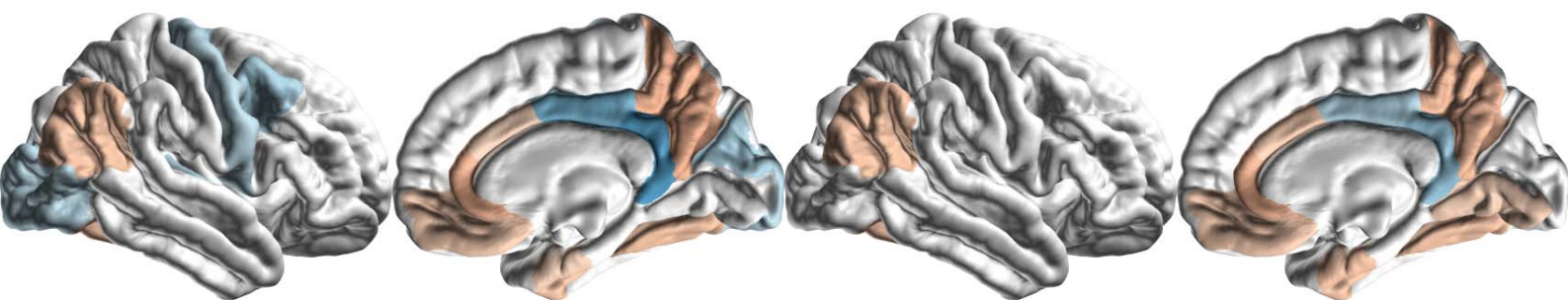


Isthmus Cingulate SA: rs3770776



Corrected for total SA

Uncorrected

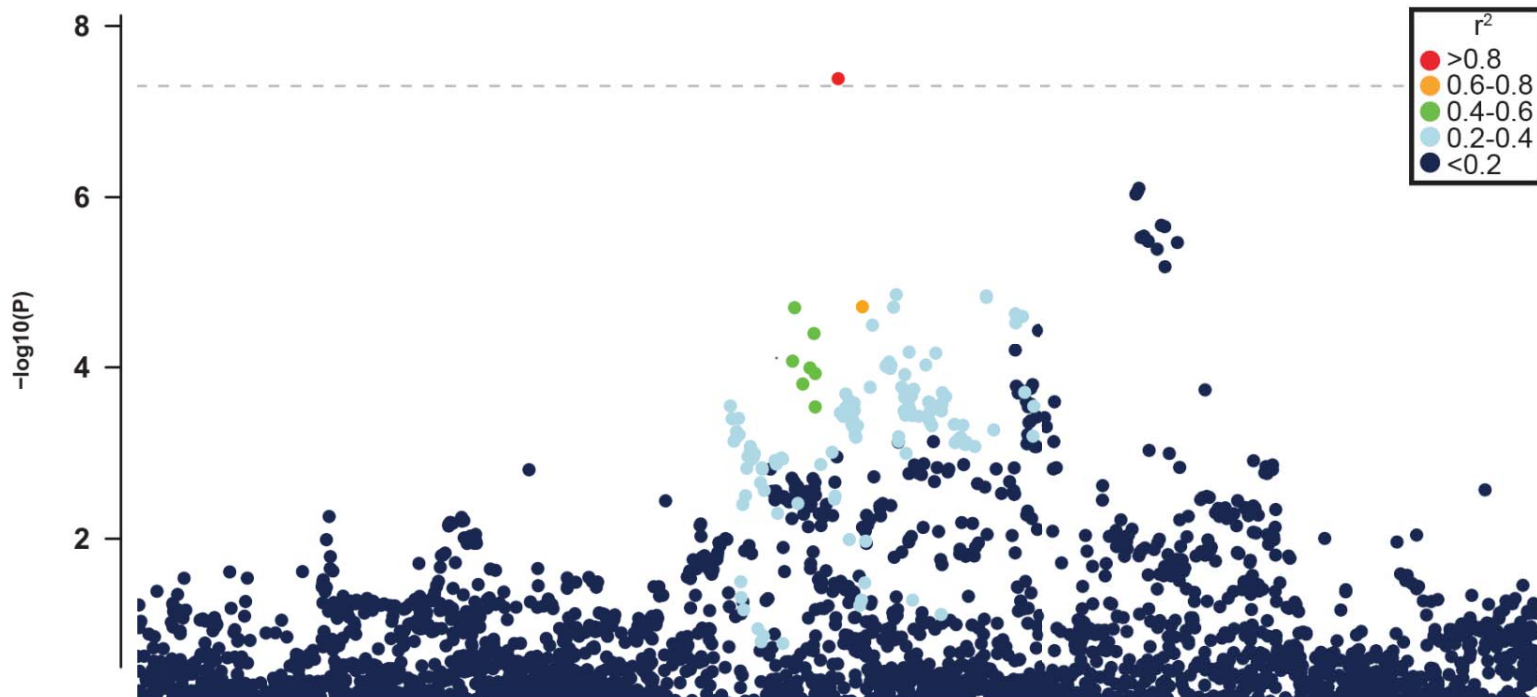
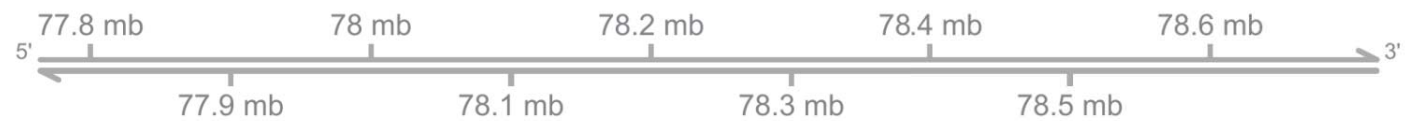


Zscore of the effect allele: A



Isthmus Cingulate SA: rs1123680

Chromosome 12



GZ>CP

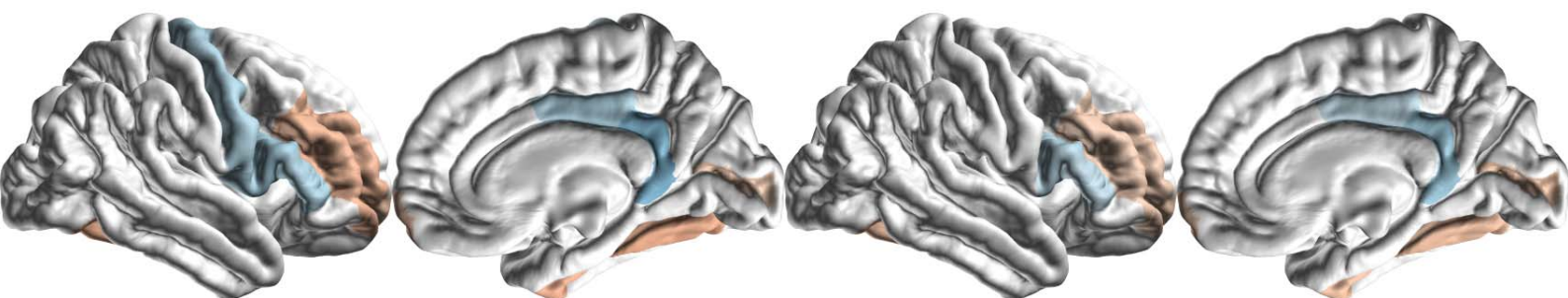


CP>GZ

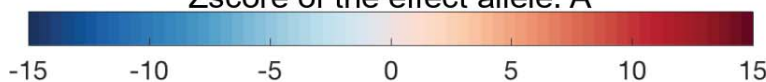


Corrected for total SA

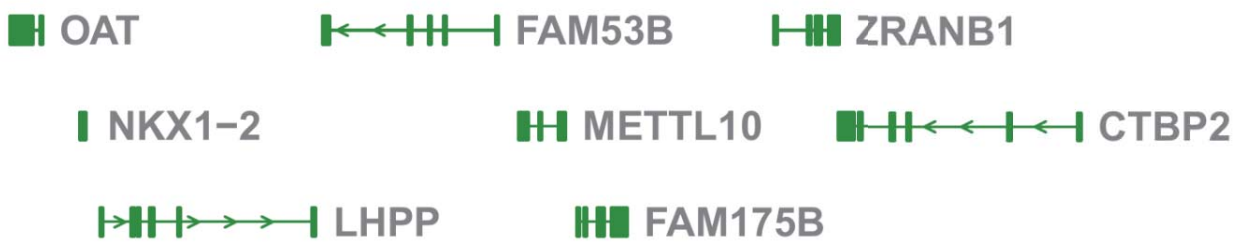
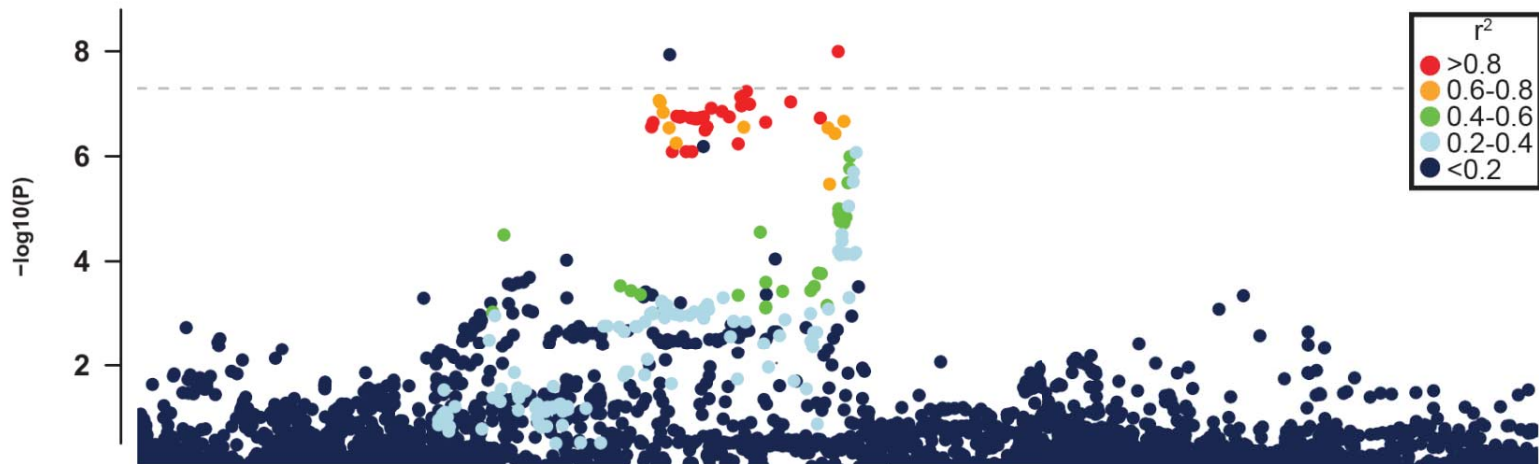
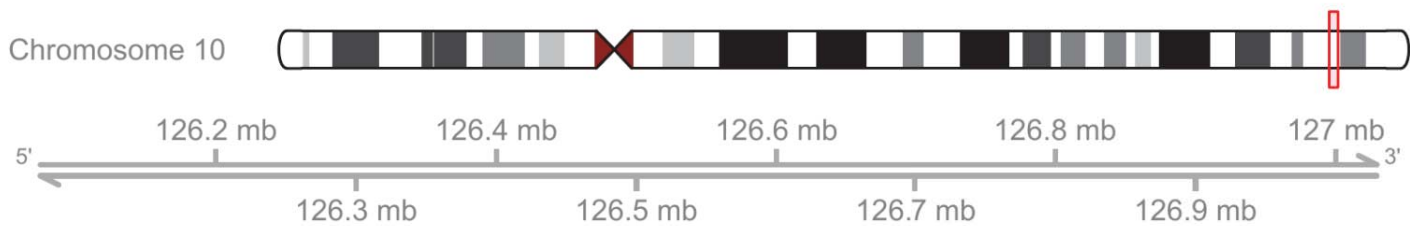
Uncorrected



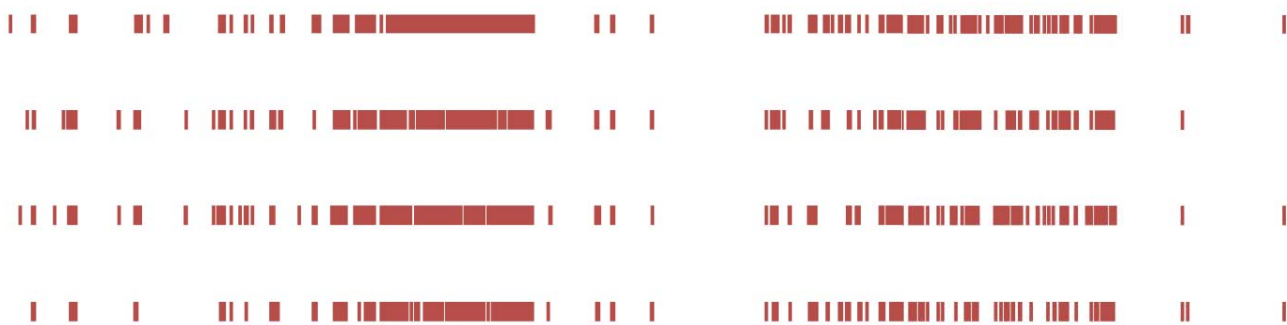
Zscore of the effect allele: A



Isthmus Cingulate TH: rs2559509

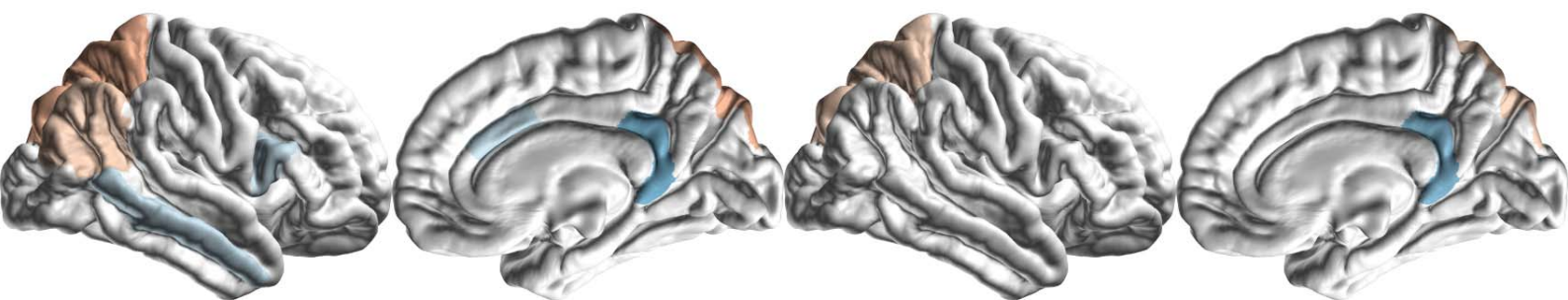


Angular Temporal dIPFC Cingulate



Corrected for average TH

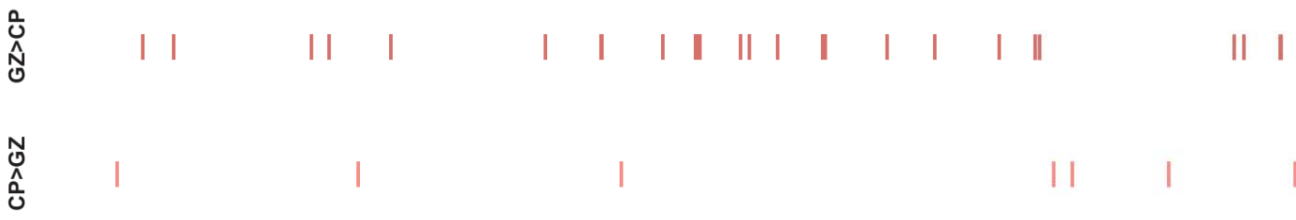
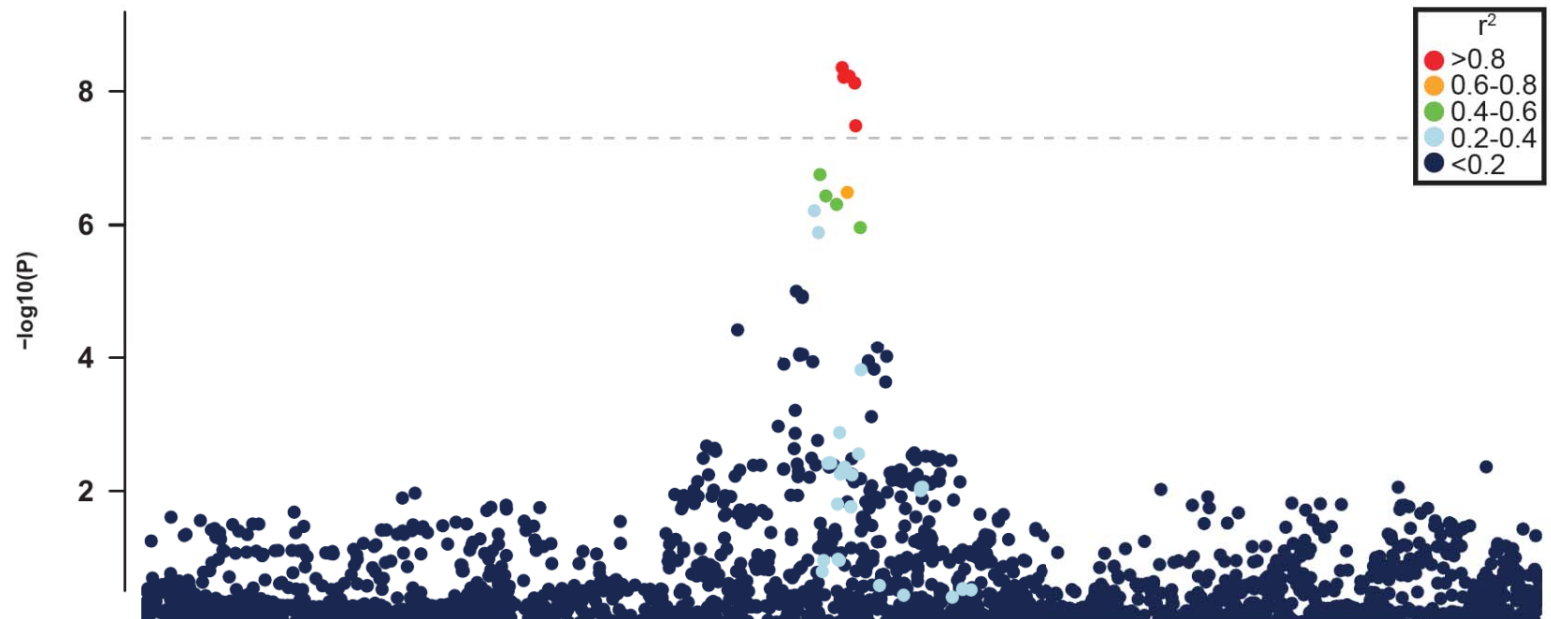
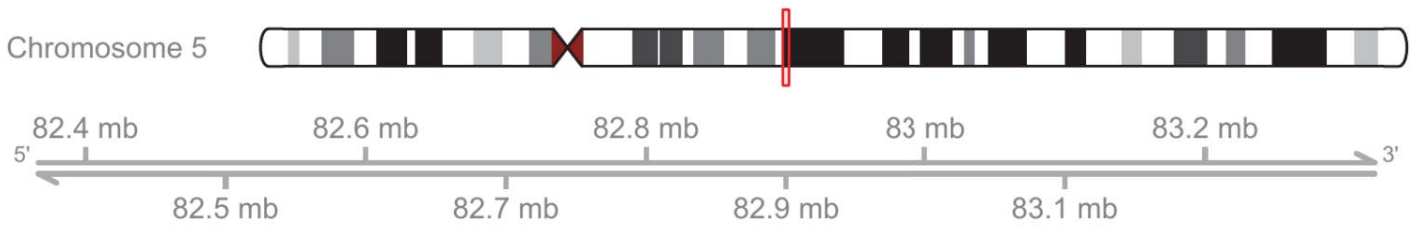
Uncorrected



Zscore of the effect allele: A

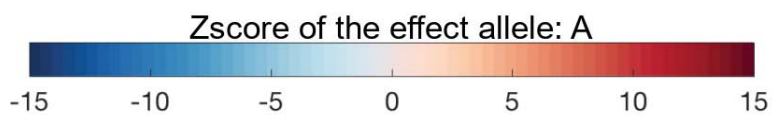
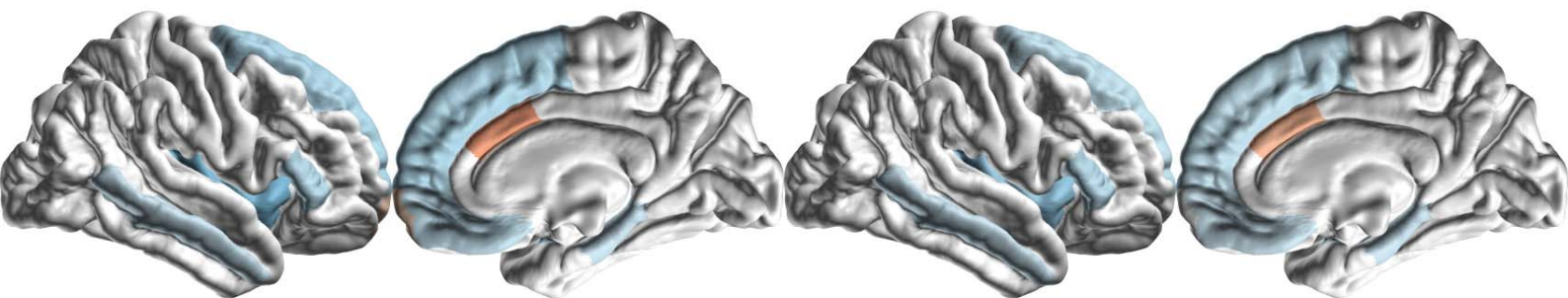


Insula SA: rs7728751

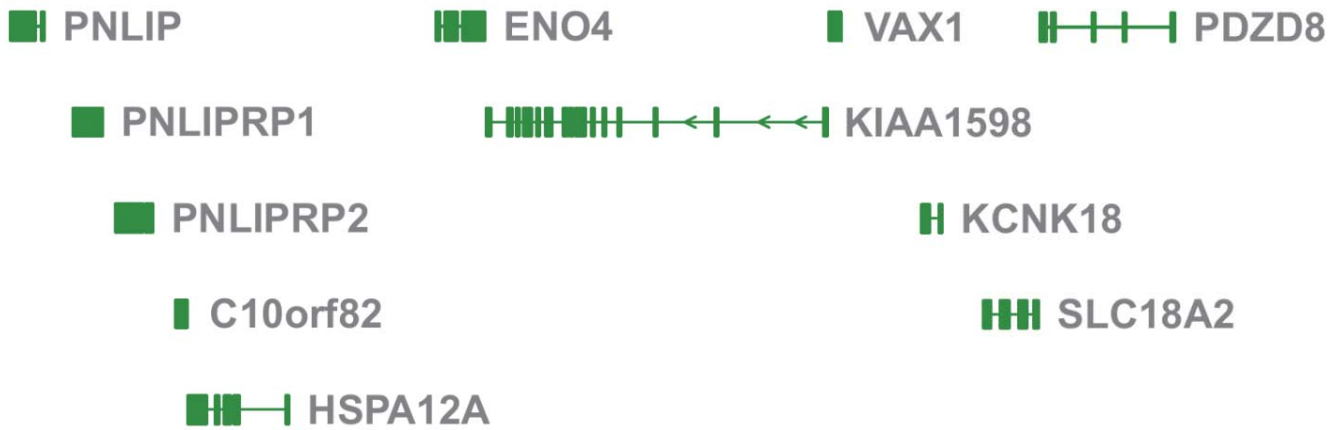
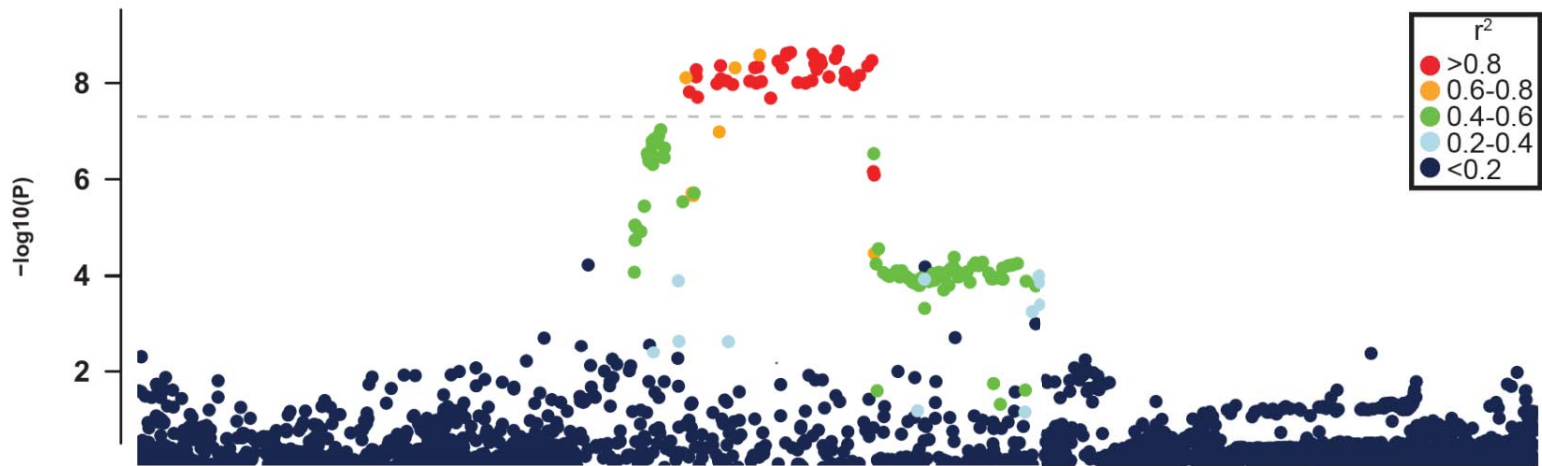
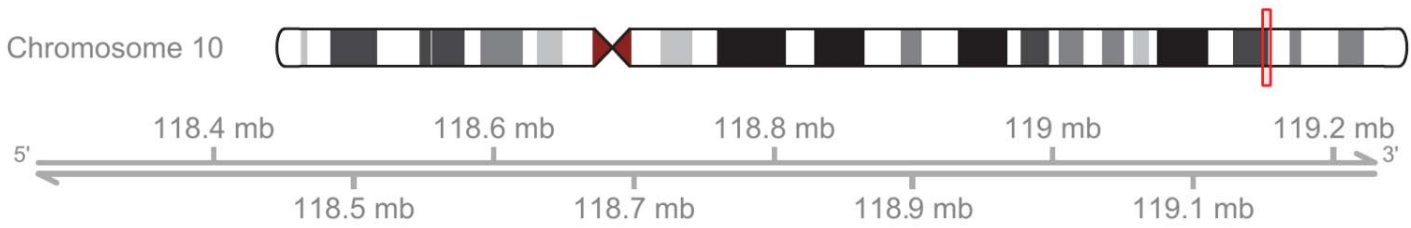


Corrected for total SA

Uncorrected



Insula SA: rs1122688

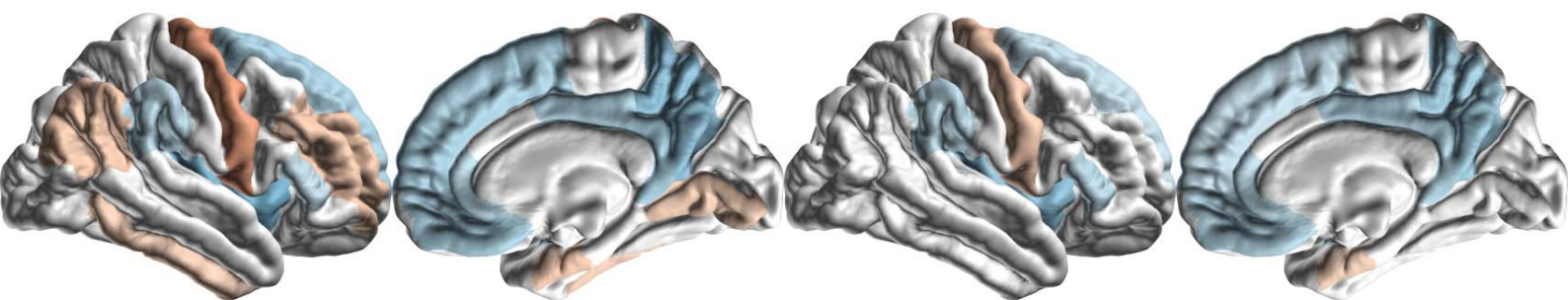


CP>GZ GZ>CP

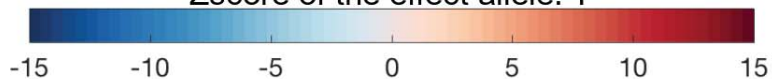


Corrected for total SA

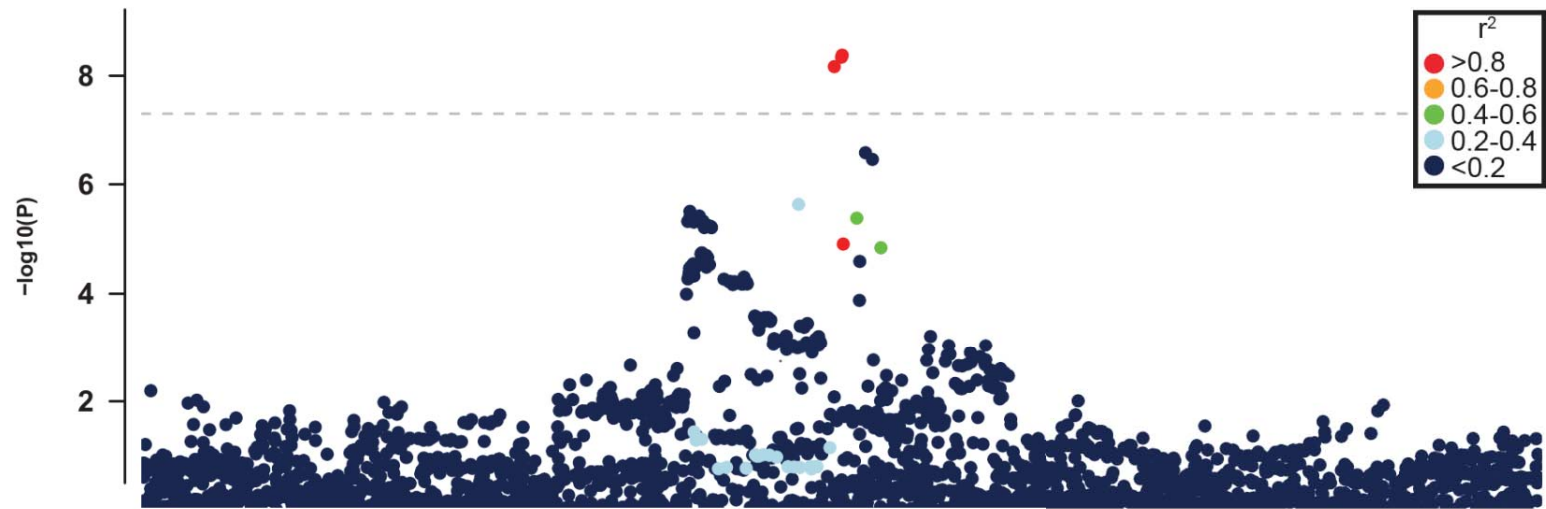
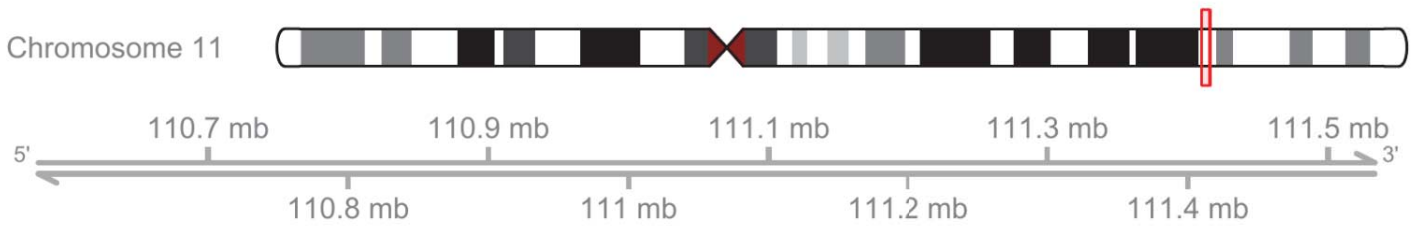
Uncorrected



Zscore of the effect allele: T



Insula SA: rs58066679



ARHGAP20

C11orf53

C11orf88

COLCA2

LAYN

POU2AF1

→

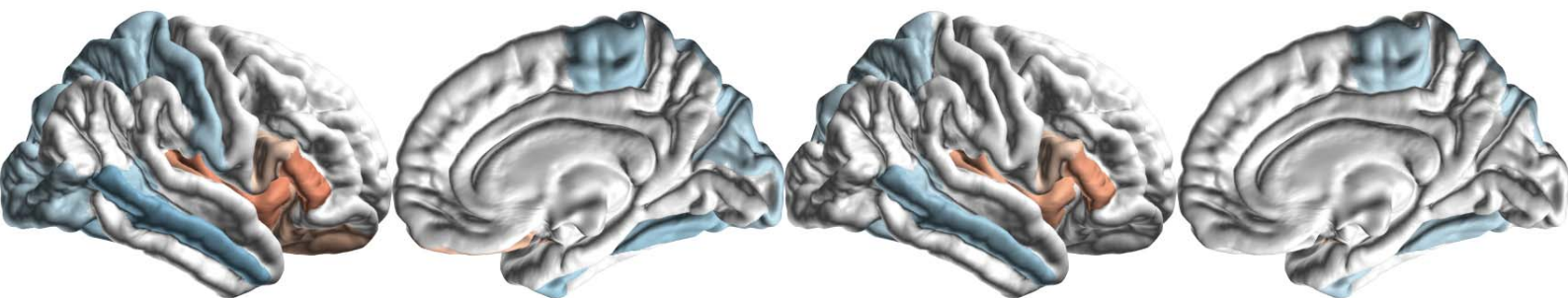
← BTG4

CP>GZ

GZ>CP

Corrected for total SA

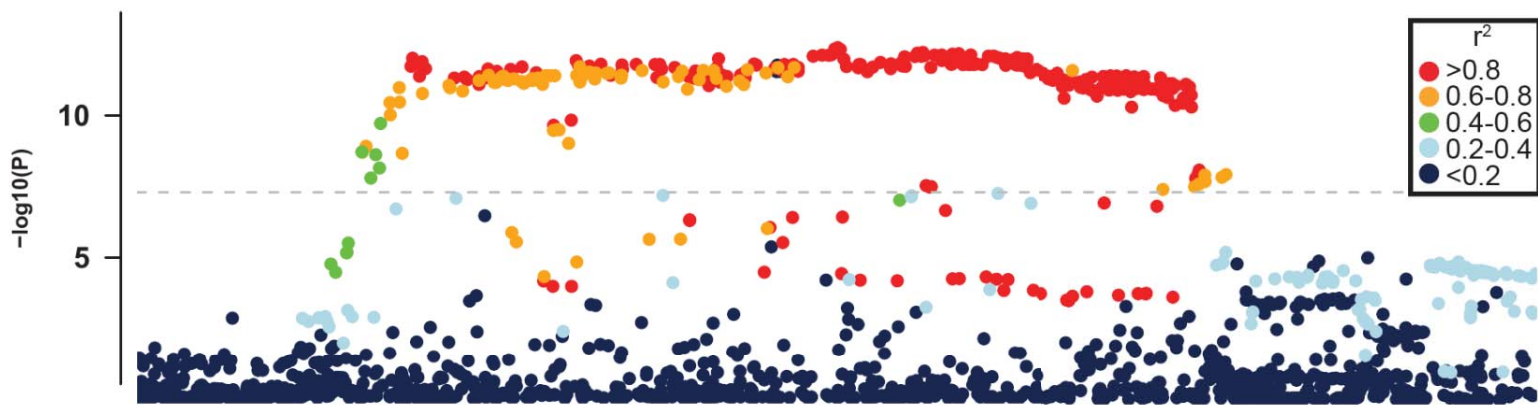
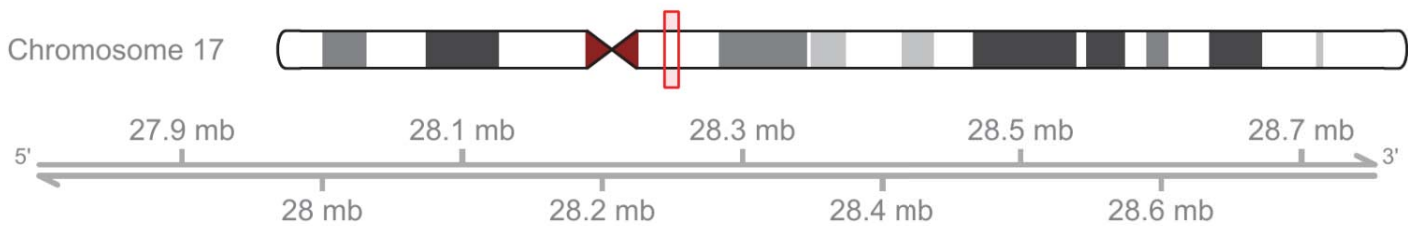
Uncorrected



Zscore of the effect allele: A

-15 -10 -5 0 5 10 15

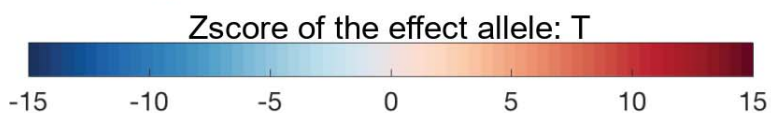
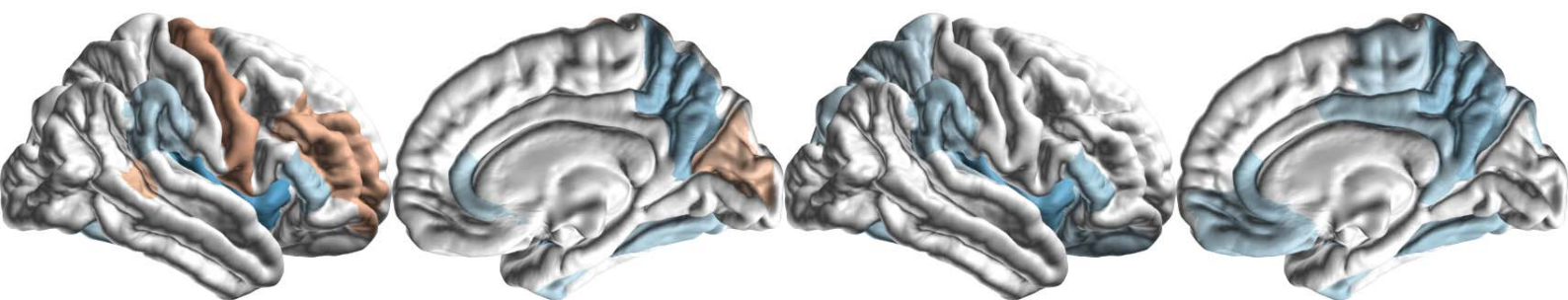
Insula SA: rs6505147



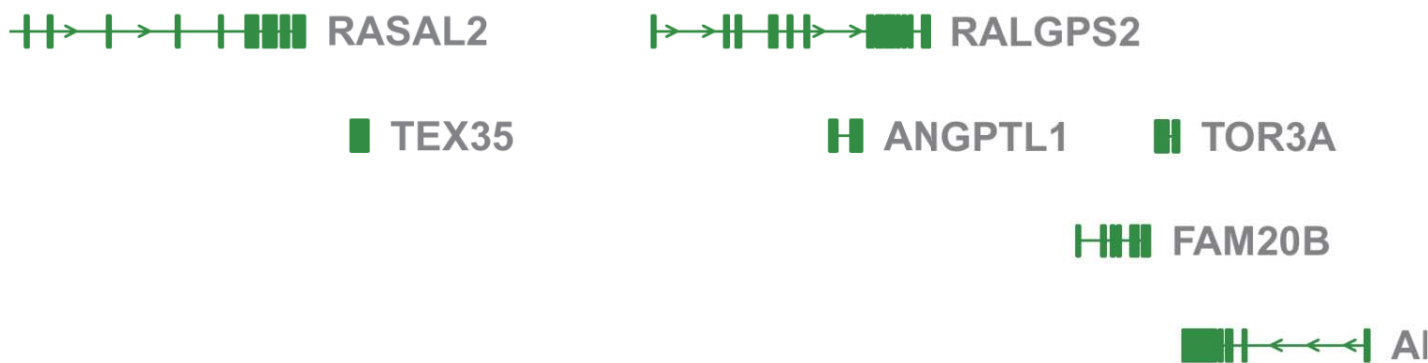
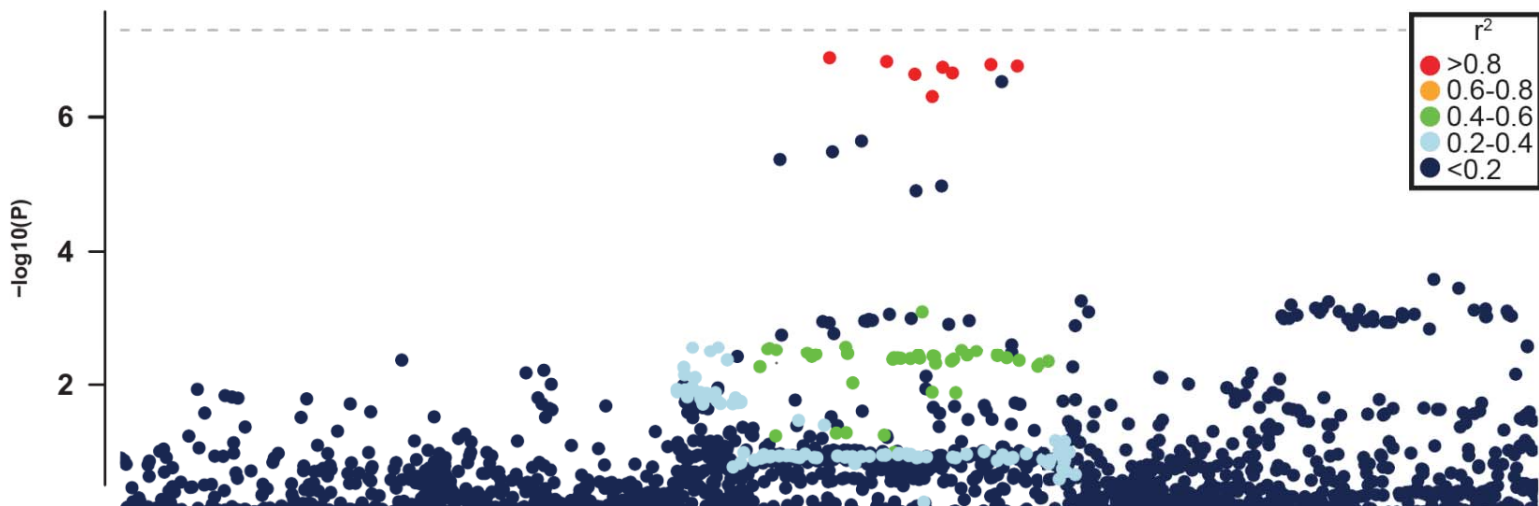
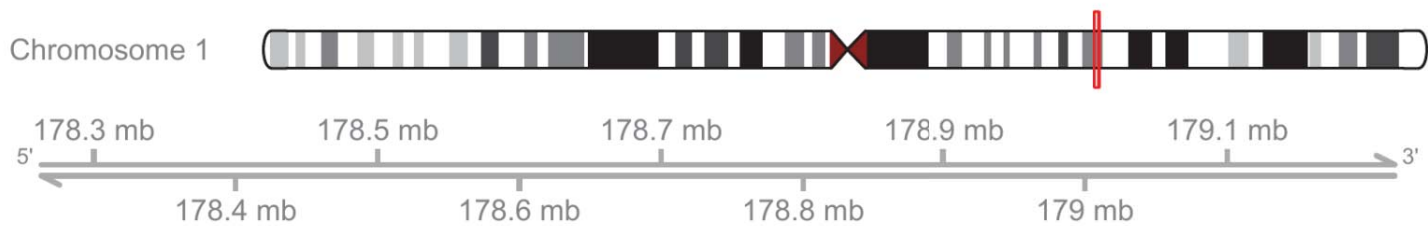
CP>GZ GZ>CP

Corrected for total SA

Uncorrected

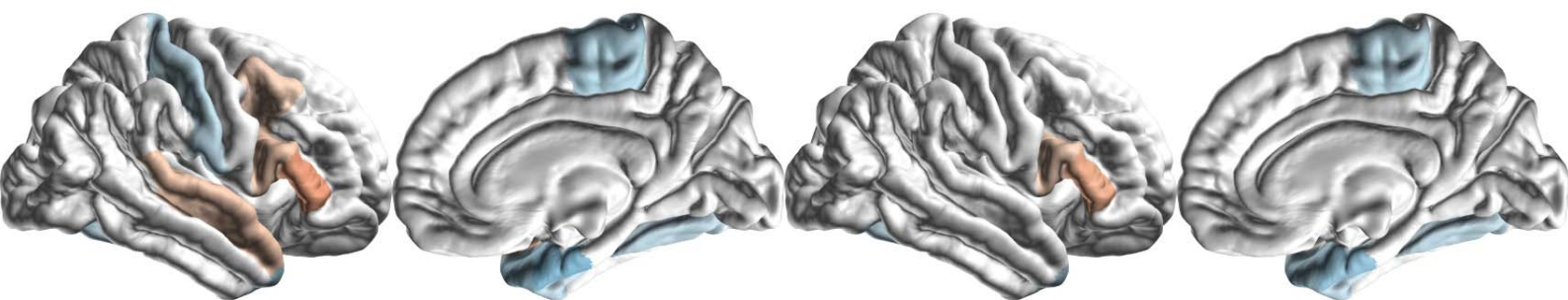


Entorhinal SA: rs6693283



Corrected for total SA

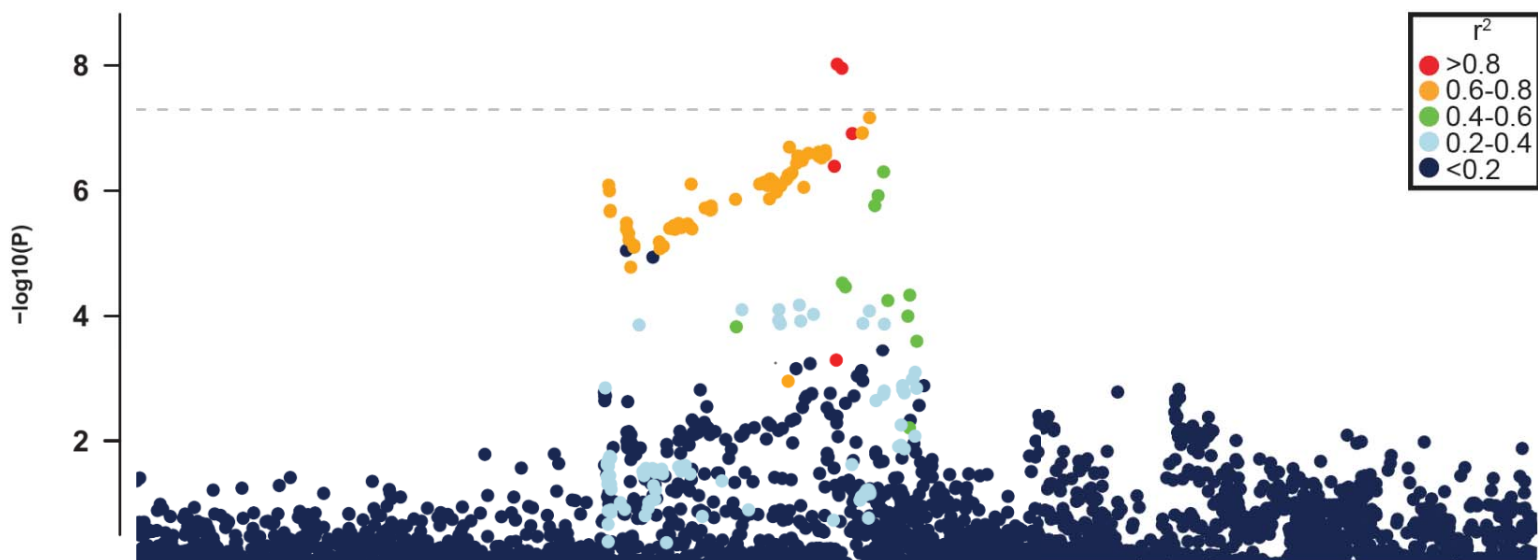
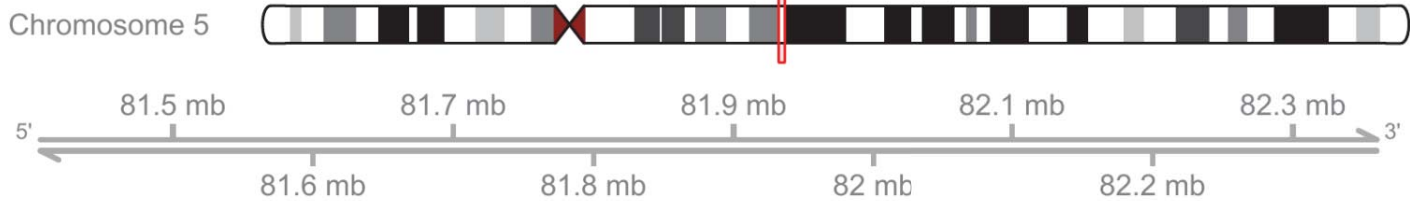
Uncorrected



Zscore of the effect allele: A

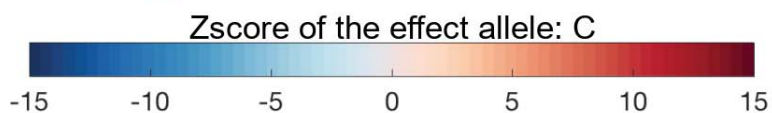
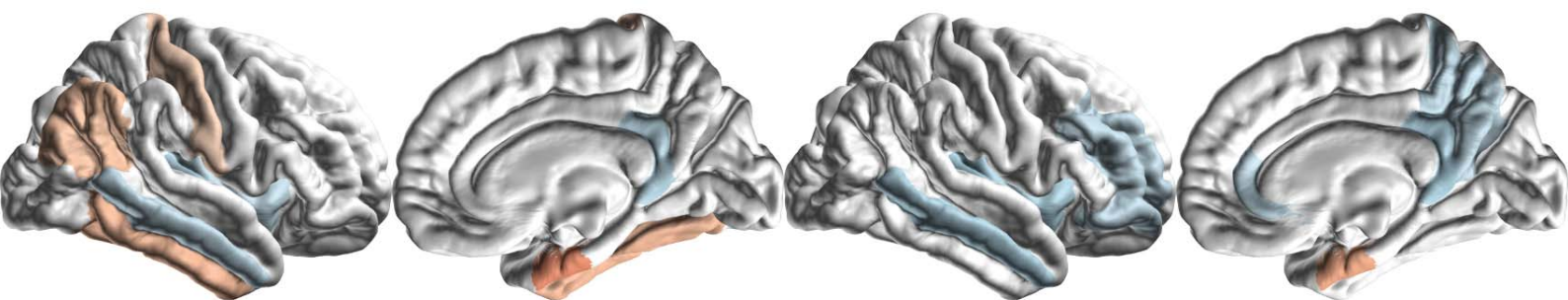


Entorhinal SA: rs4147321



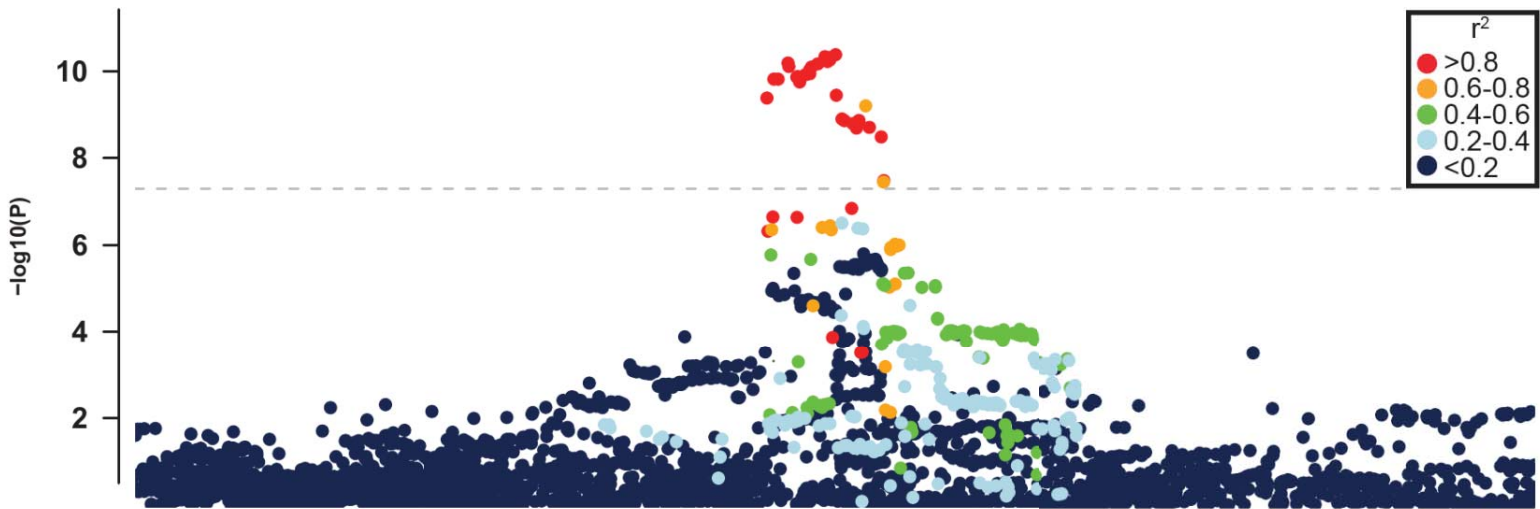
Corrected for total SA

Uncorrected



Entorhinal SA: rs7141150

Chromosome 14



DAAM1 CCDC175

GPR135 F

L3HYPDH

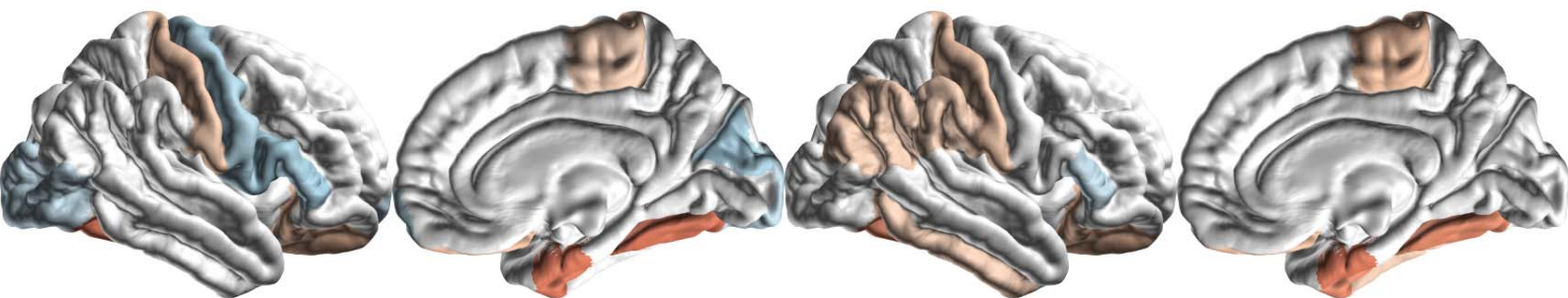
JKAMP

CP>GZ GZ>CP



Corrected for total SA

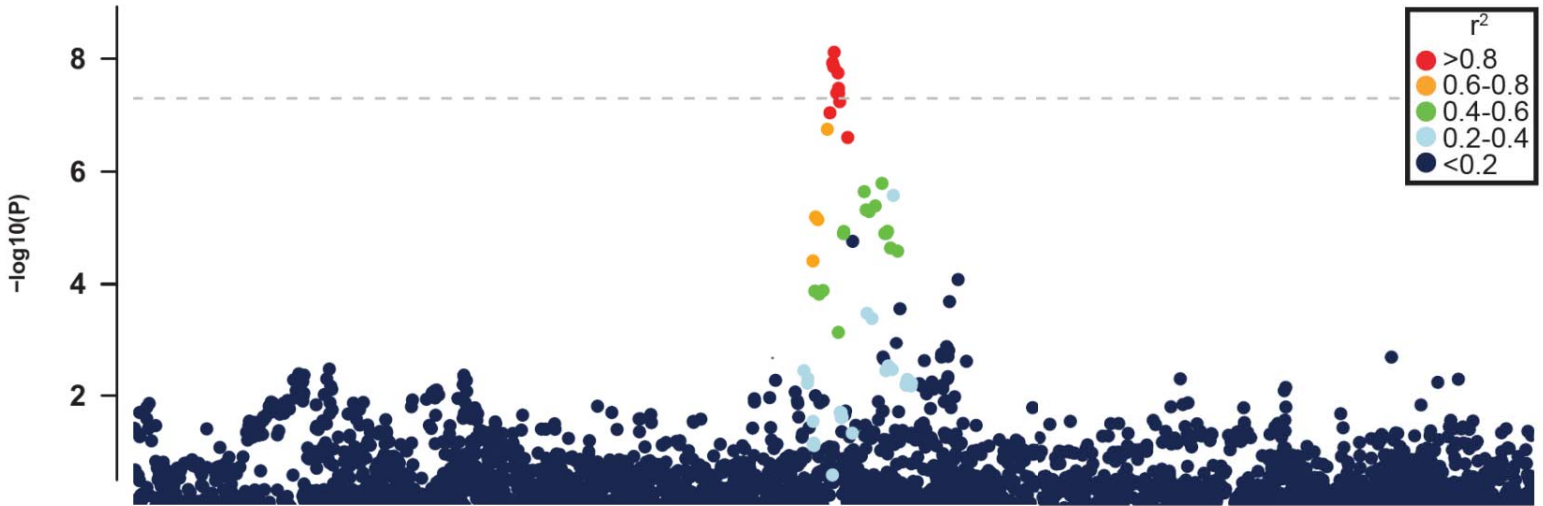
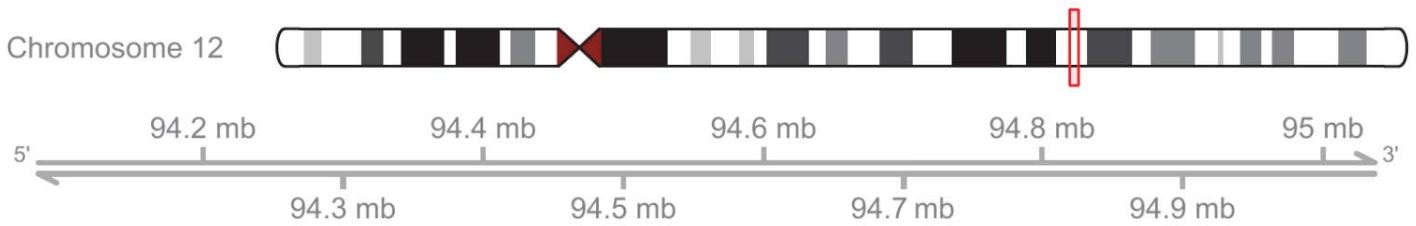
Uncorrected



Zscore of the effect allele: A

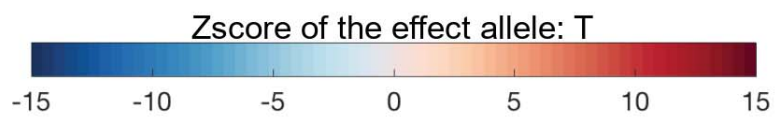
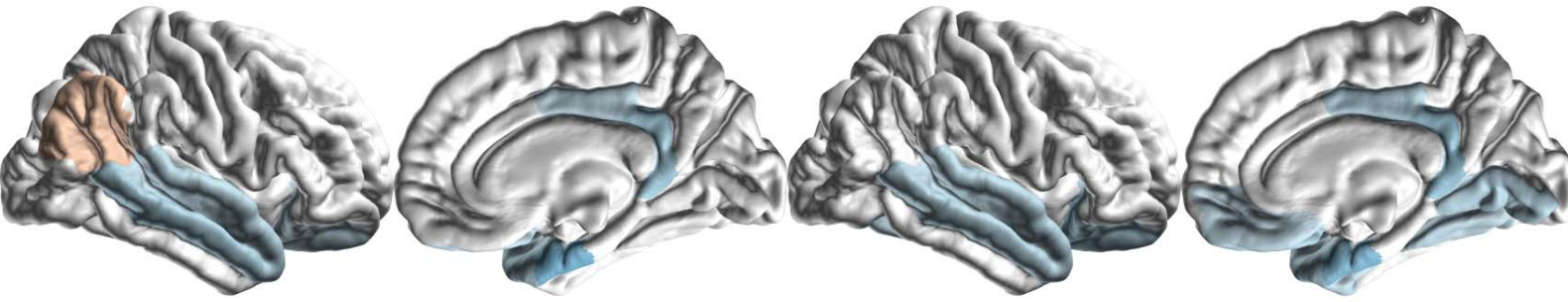


Entorhinal TH: rs2885259

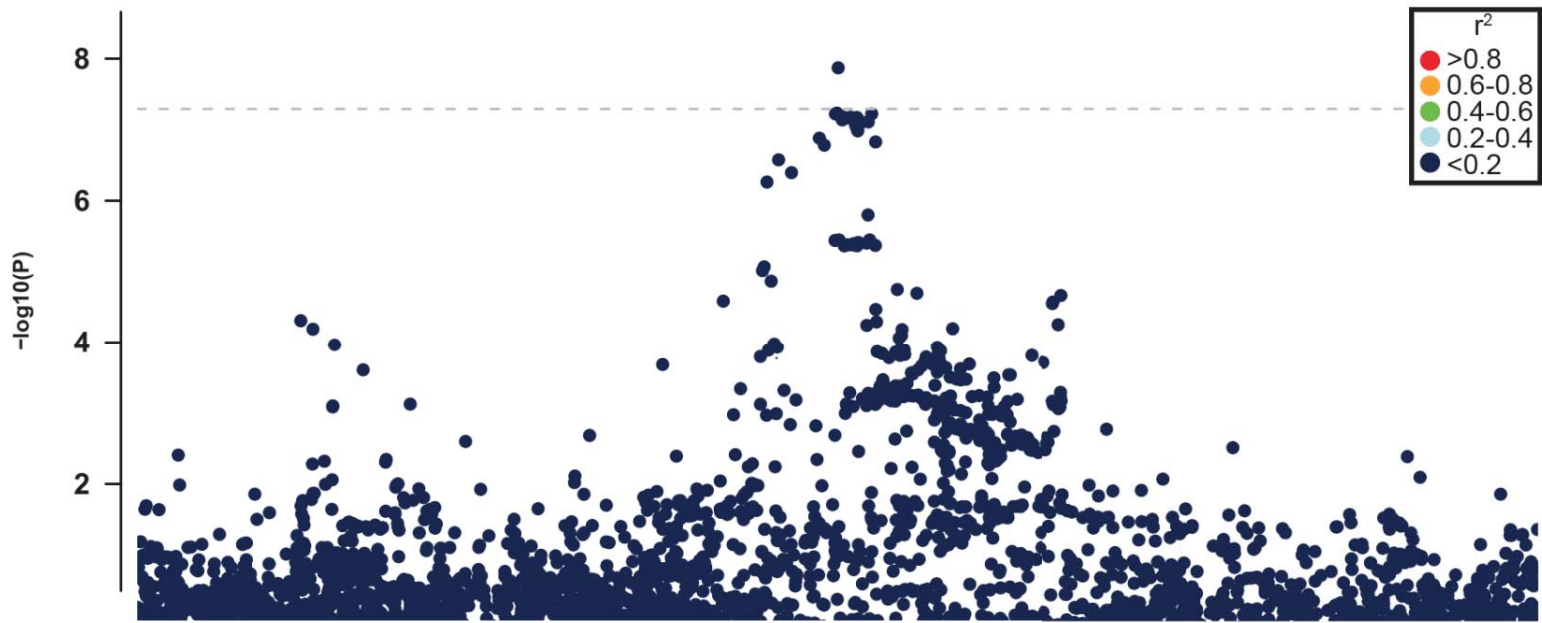
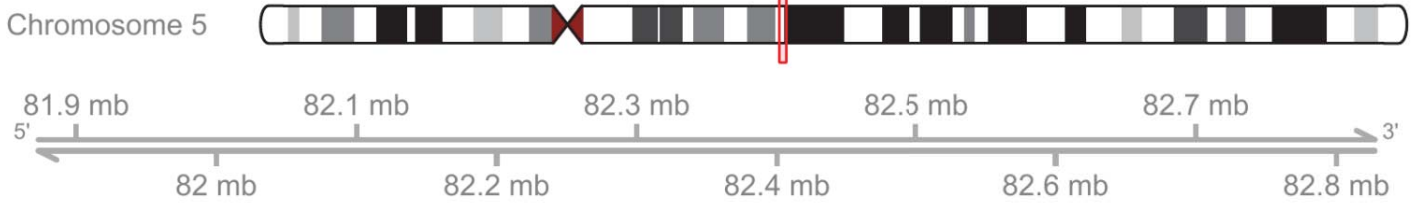


Corrected for average TH

Uncorrected



Parahippocampal SA: rs27493



TMEM167A

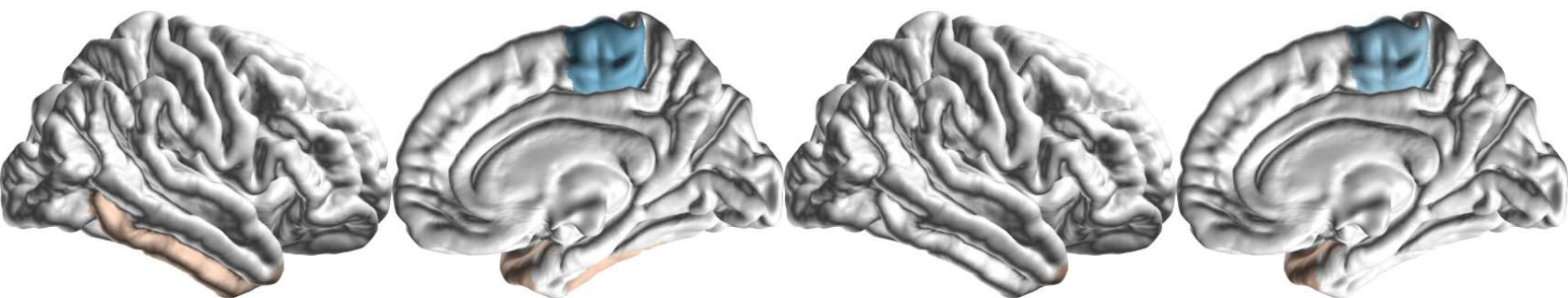
XRCC4

GZ>CP

CP>GZ

Corrected for total SA

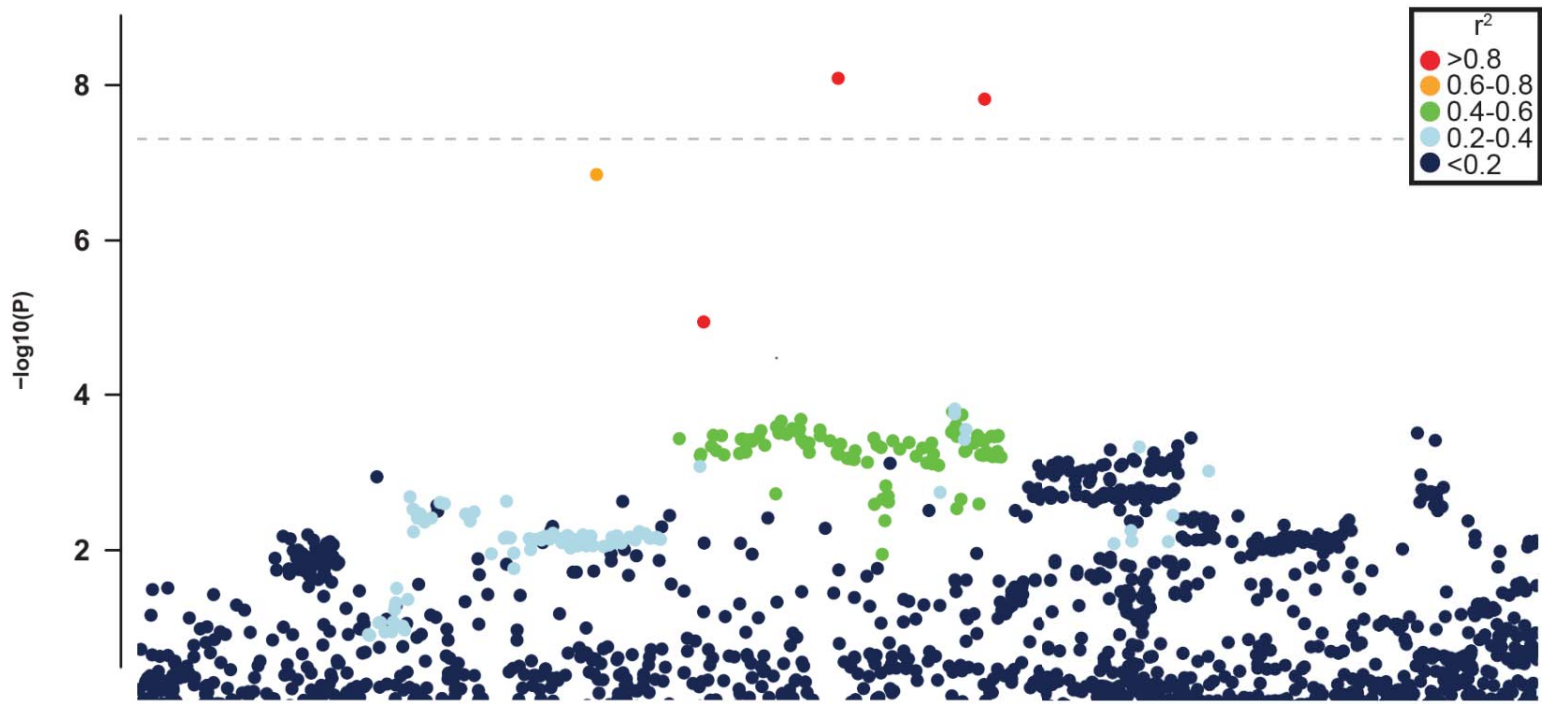
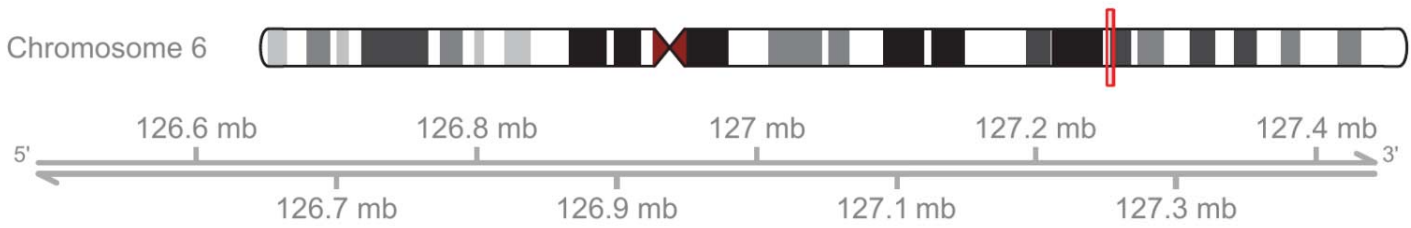
Uncorrected



Zscore of the effect allele: A



Parahippocampal SA: rs4273712



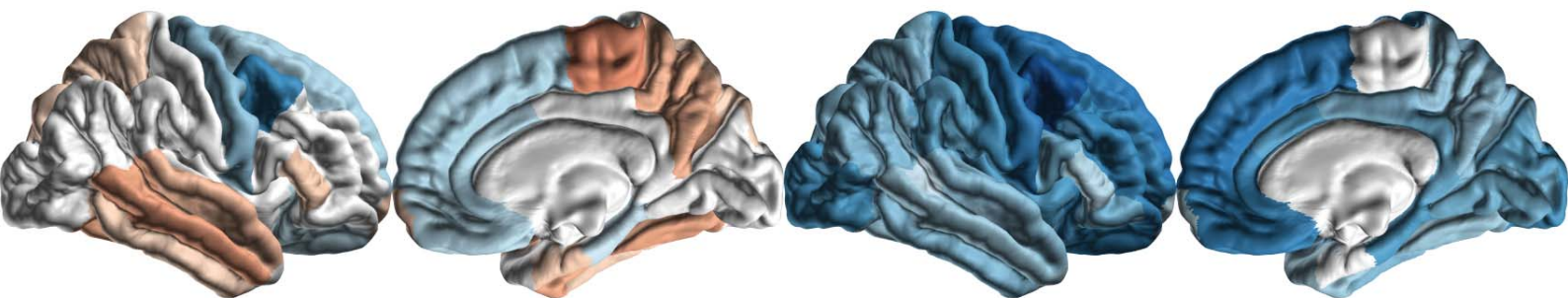
CENPW

GZ>CP

CP>GZ

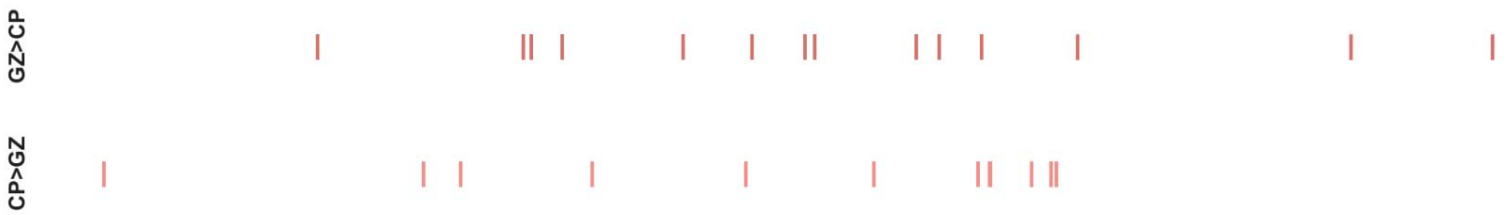
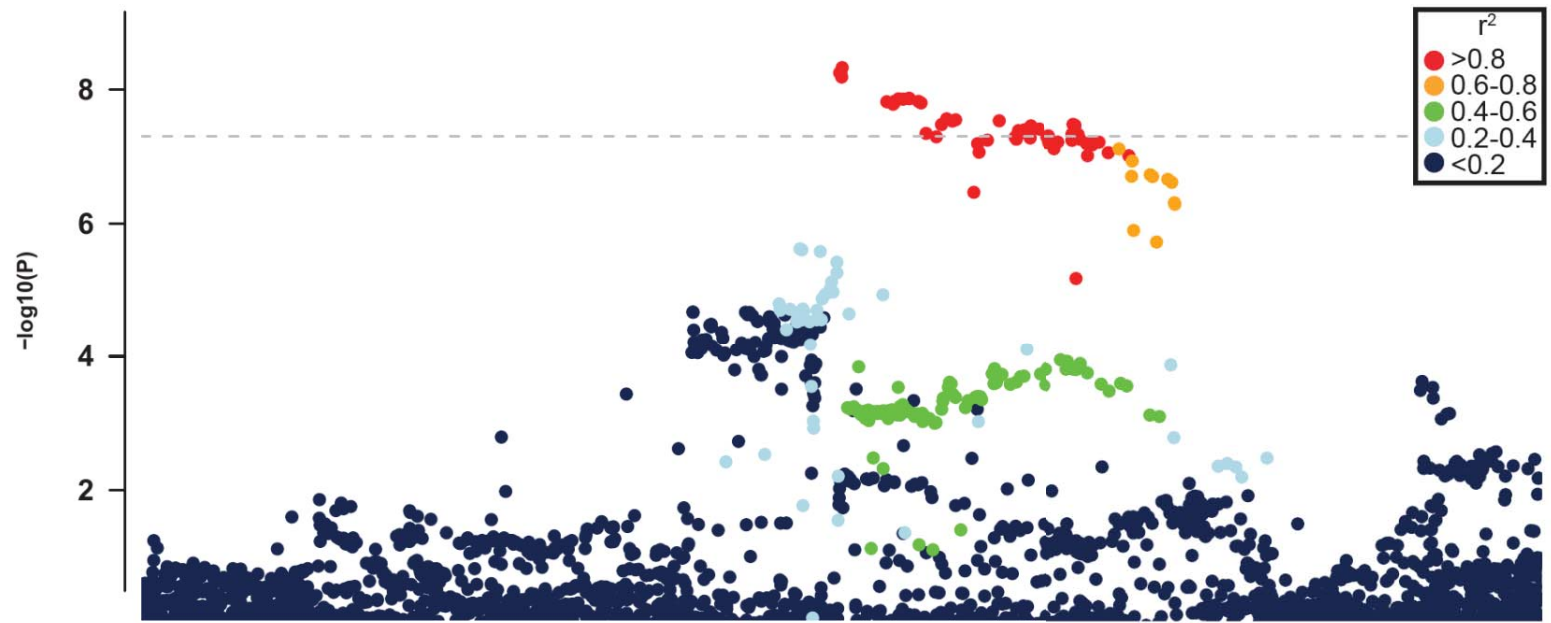
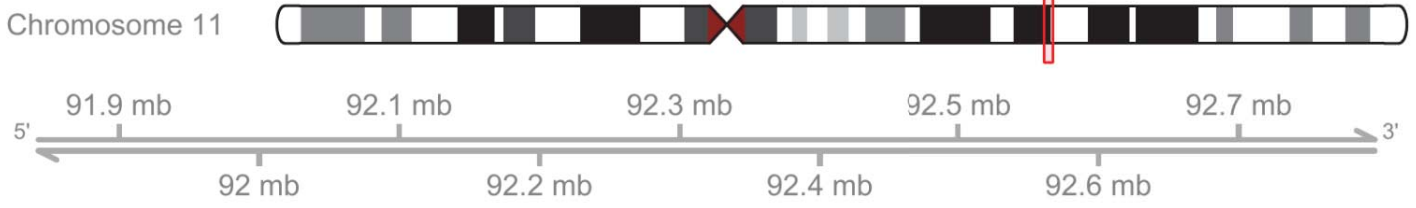
Corrected for total SA

Uncorrected



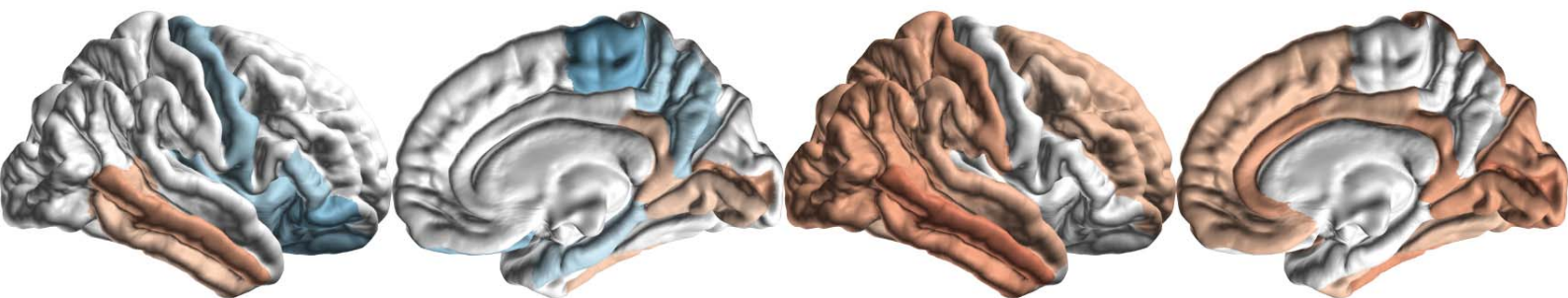
-15 -10 -5 0 5 10 15

Parahippocampal SA: rs1792354



Corrected for total SA

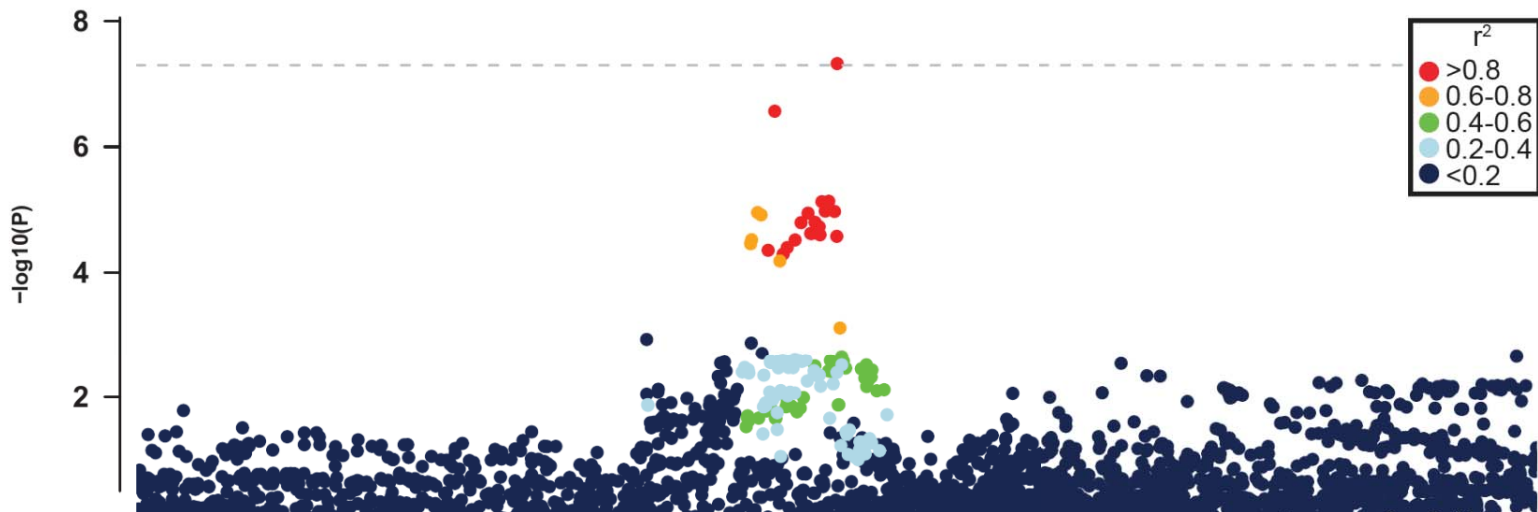
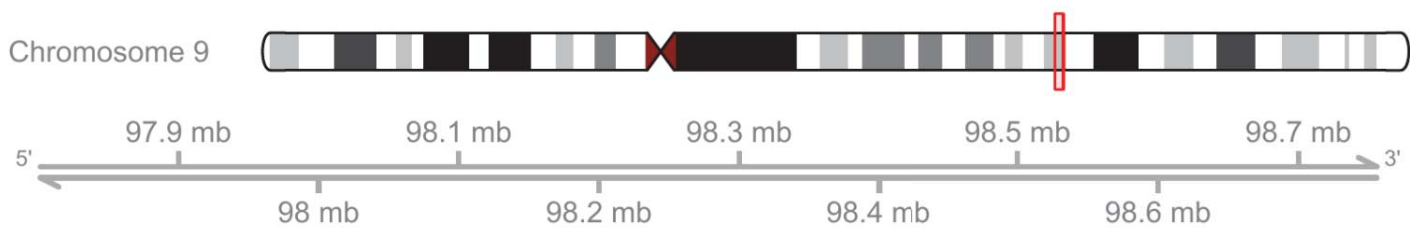
Uncorrected



Zscore of the effect allele: T

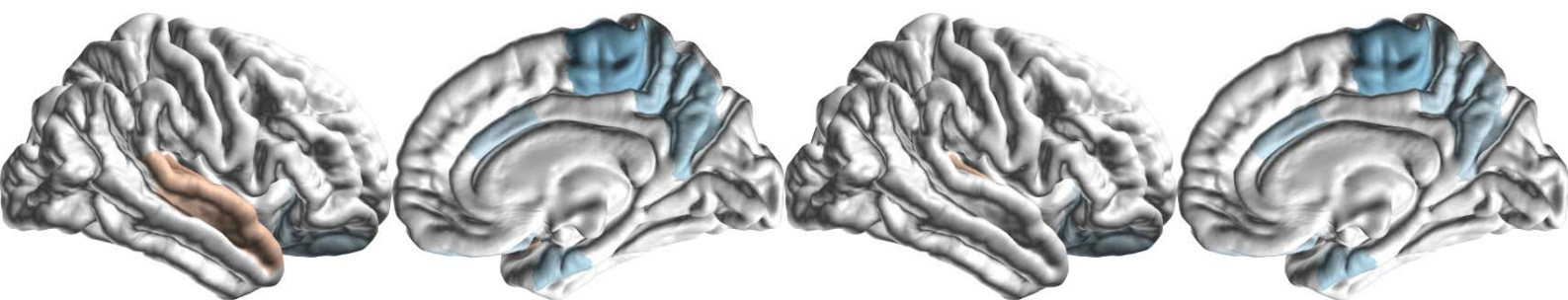


Parahippocampal TH: rs28591501



Corrected for average TH

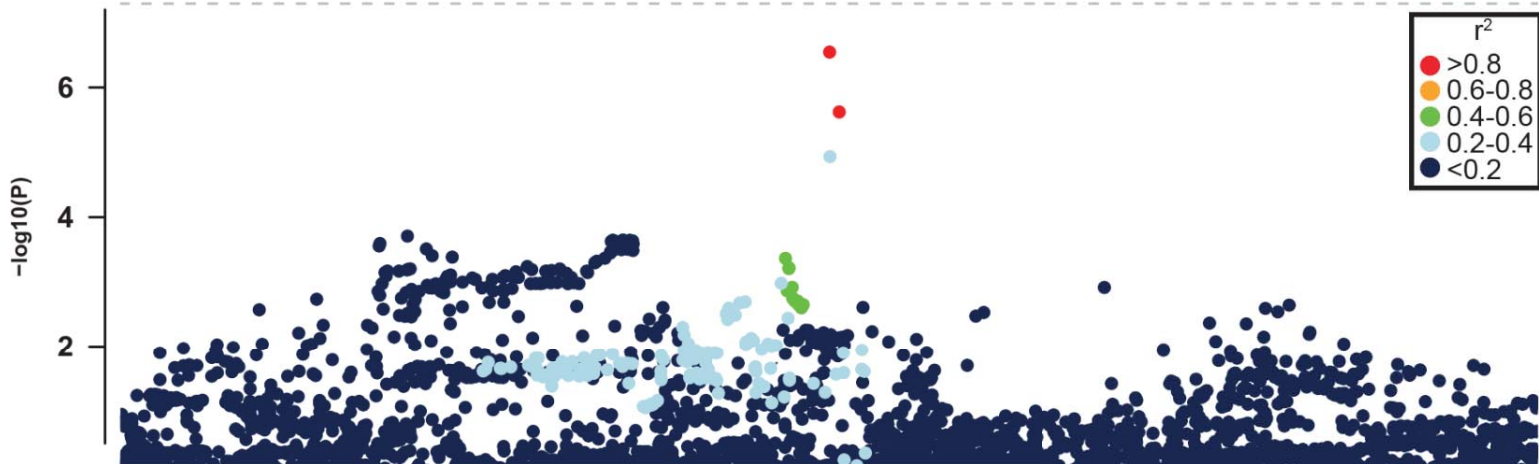
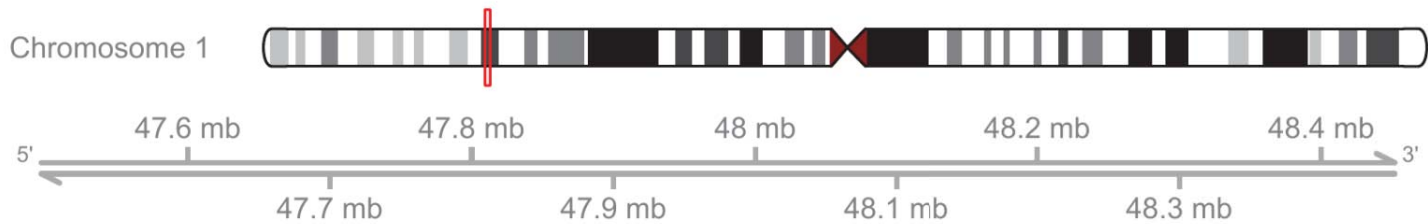
Uncorrected



Zscore of the effect allele: A

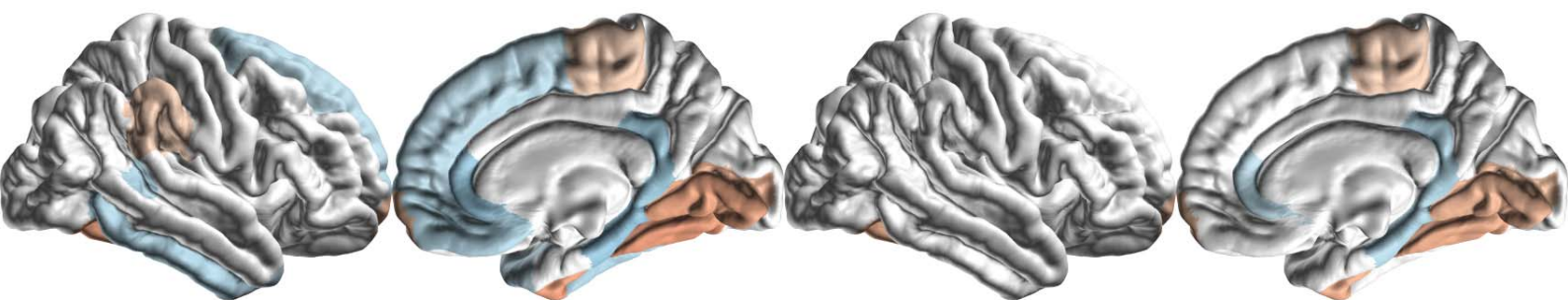


Fusiform SA: rs61784835

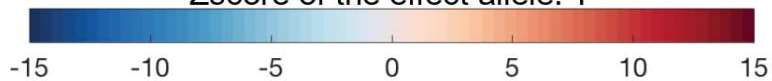


Corrected for total SA

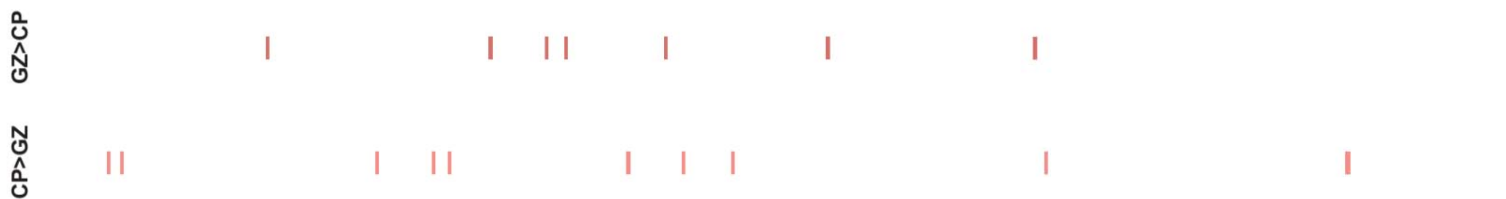
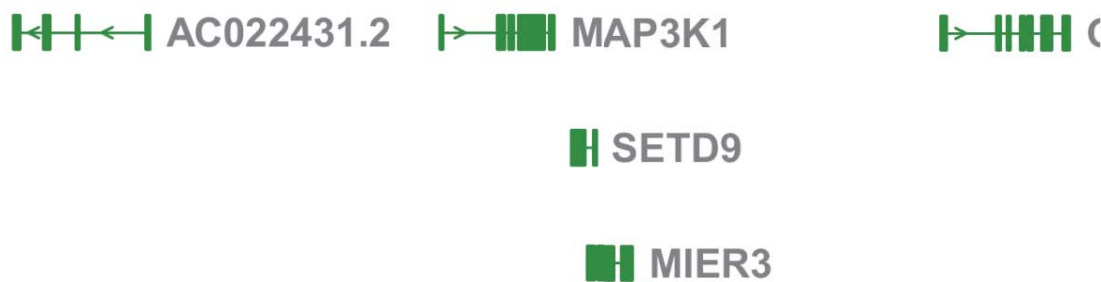
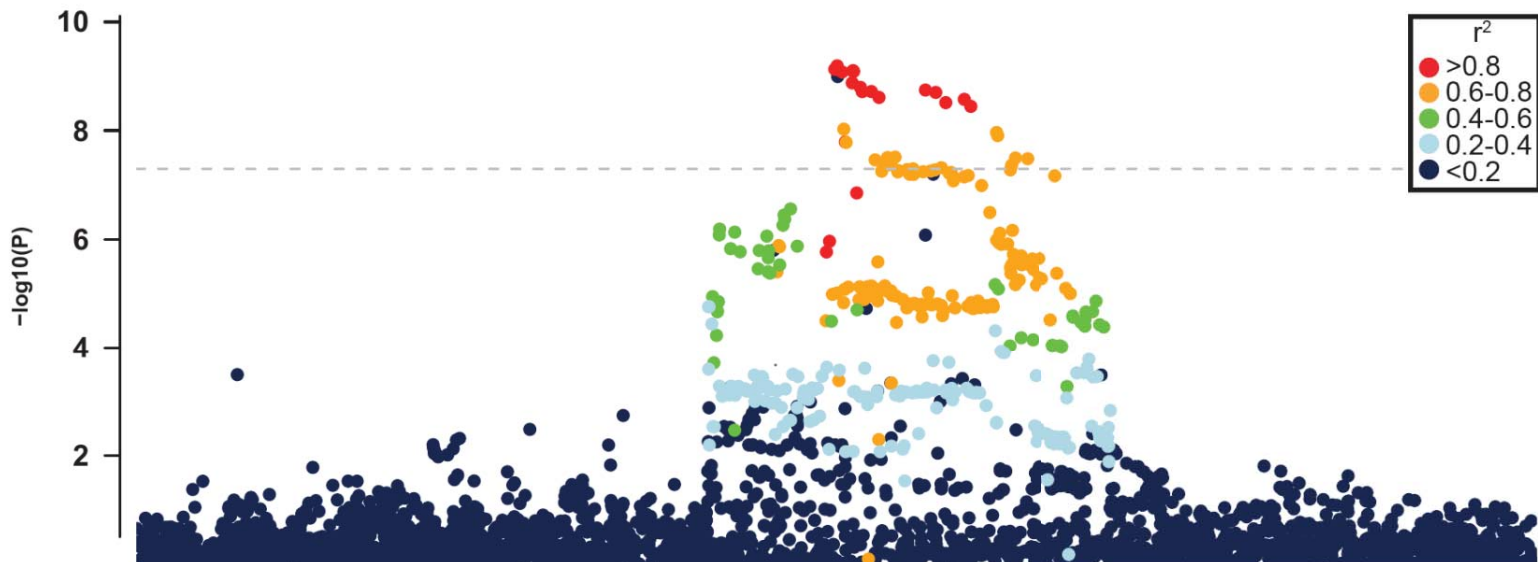
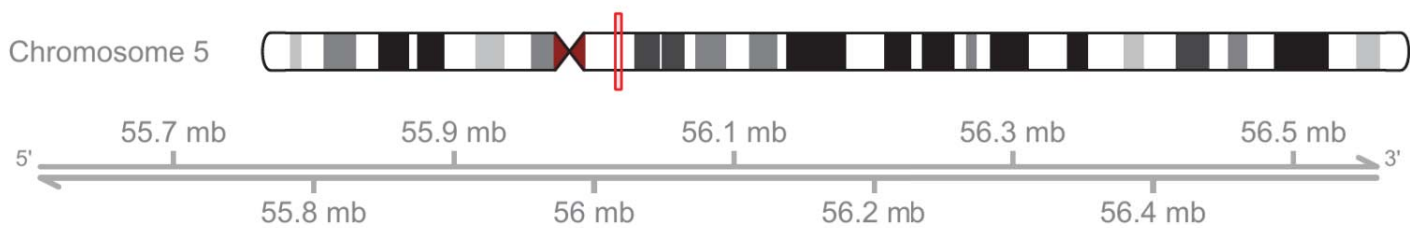
Uncorrected



Zscore of the effect allele: T

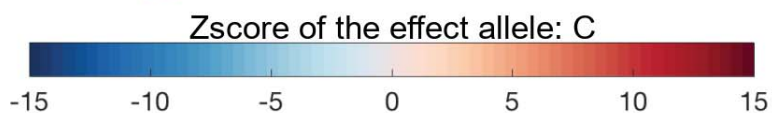
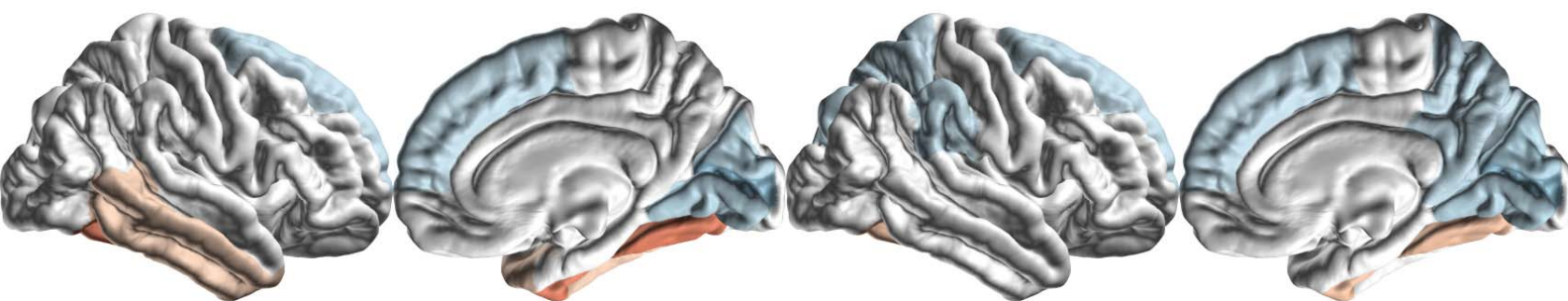


Fusiform SA: rs10940512

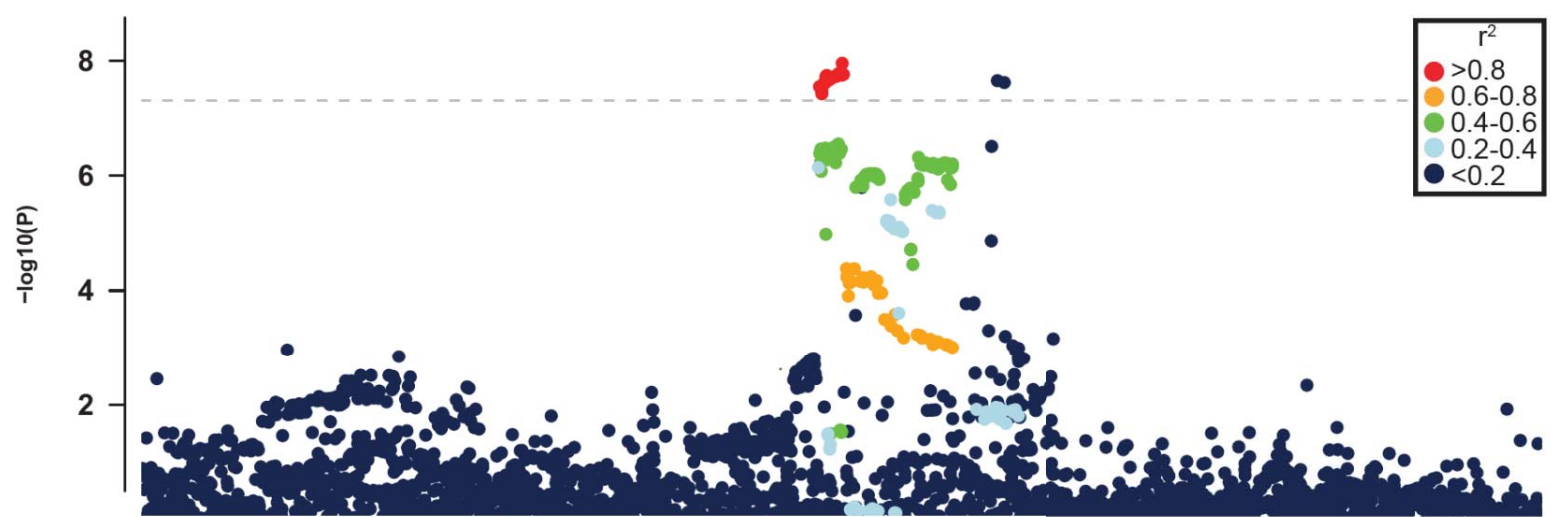
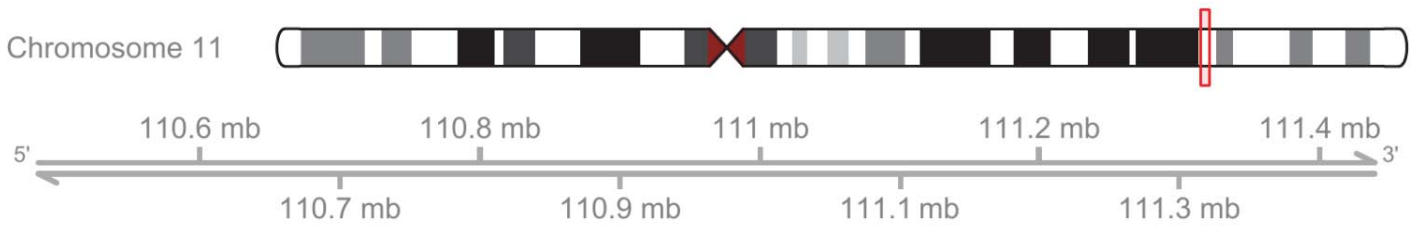


Corrected for total SA

Uncorrected

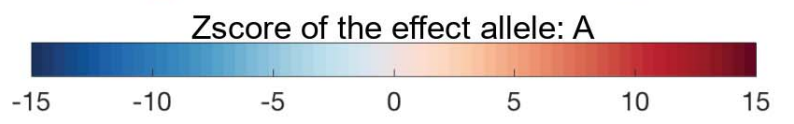
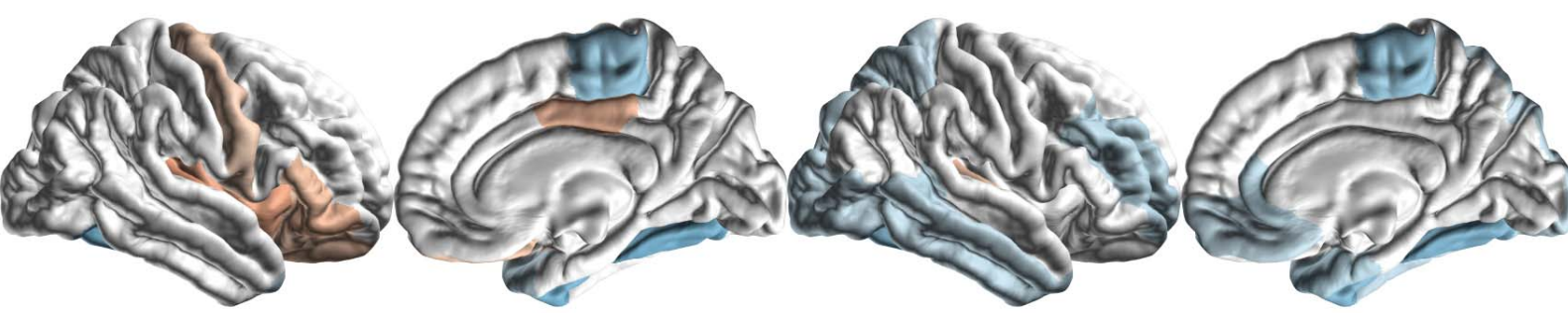


Fusiform SA: rs7123402



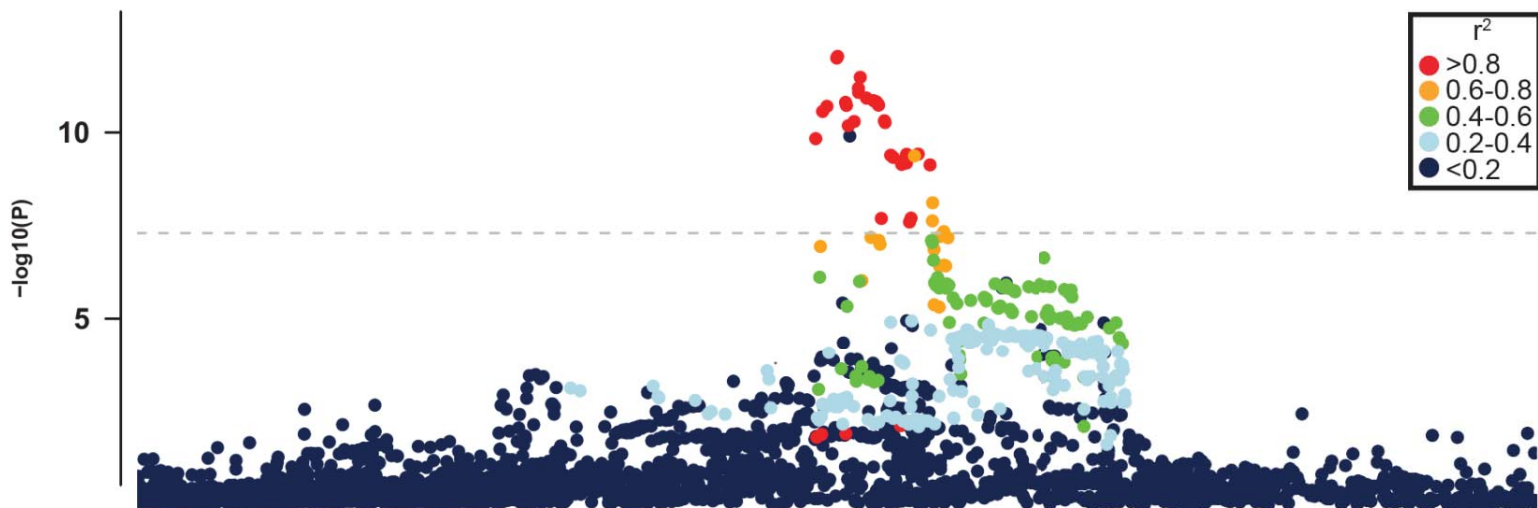
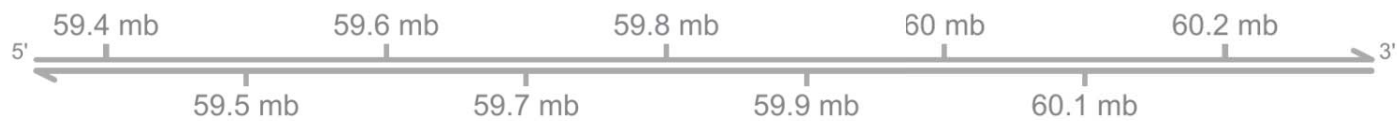
Corrected for total SA

Uncorrected



Fusiform SA: rs17834032

Chromosome 14



DAAM1 JKAMP

GPR135

L3HYPDH

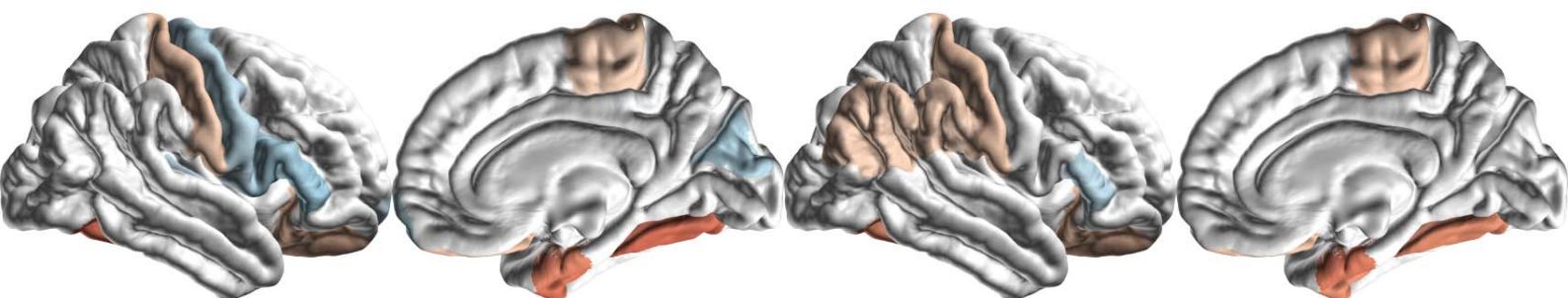
CCDC175

CP>GZ GZ>CP

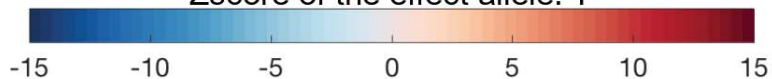


Corrected for total SA

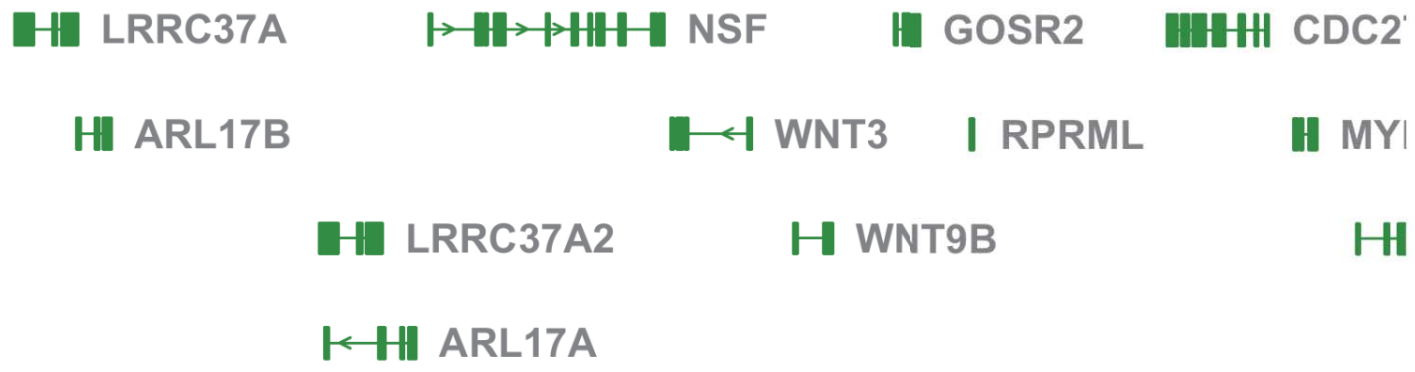
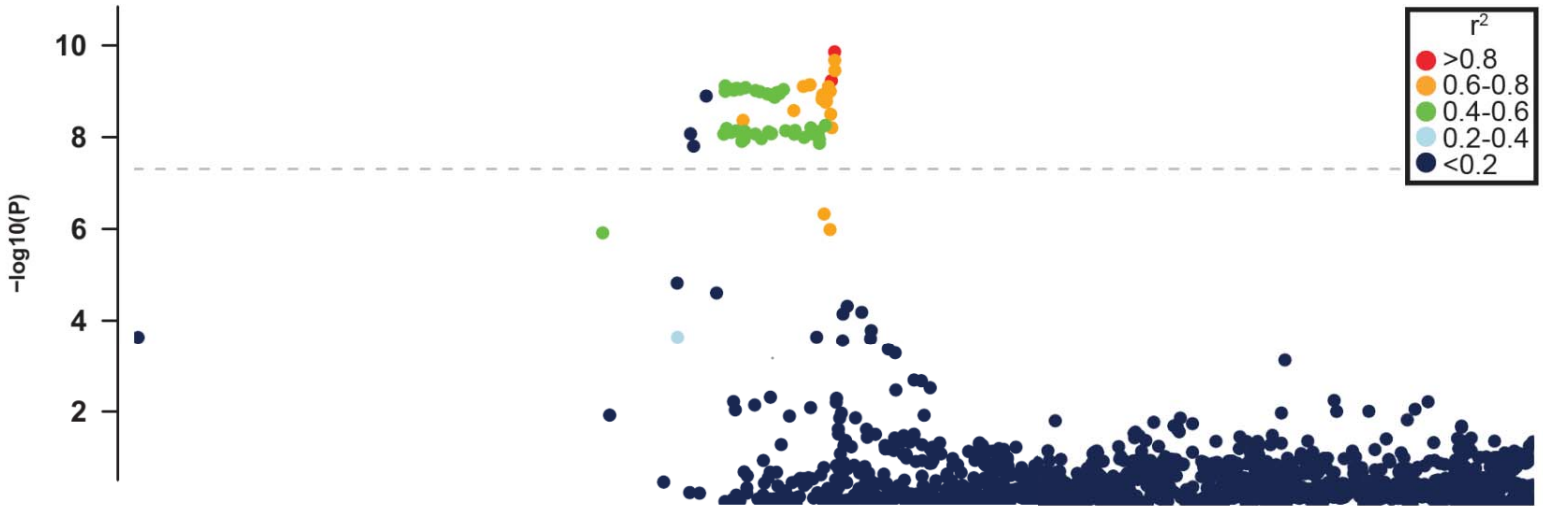
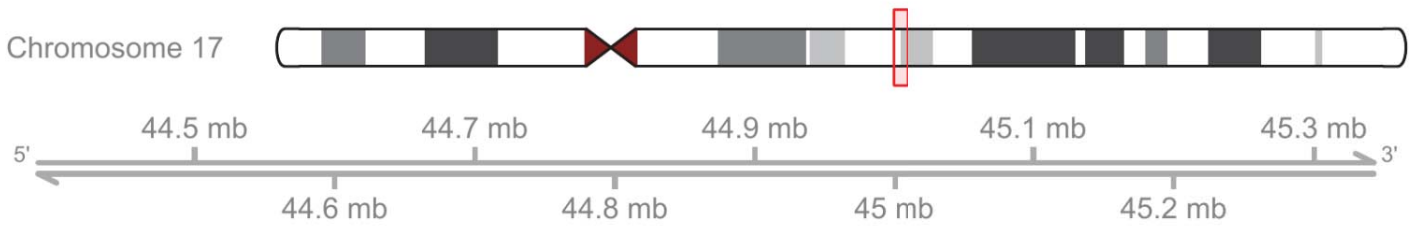
Uncorrected



Zscore of the effect allele: T



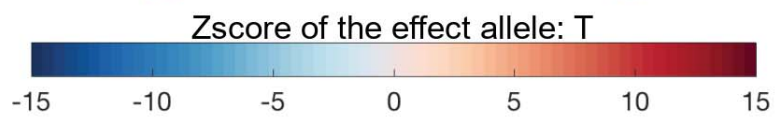
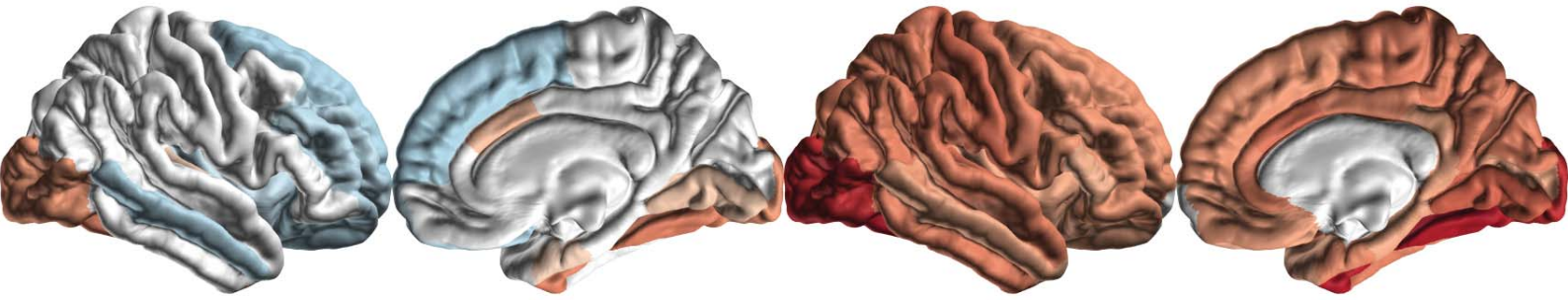
Fusiform SA: rs2074404



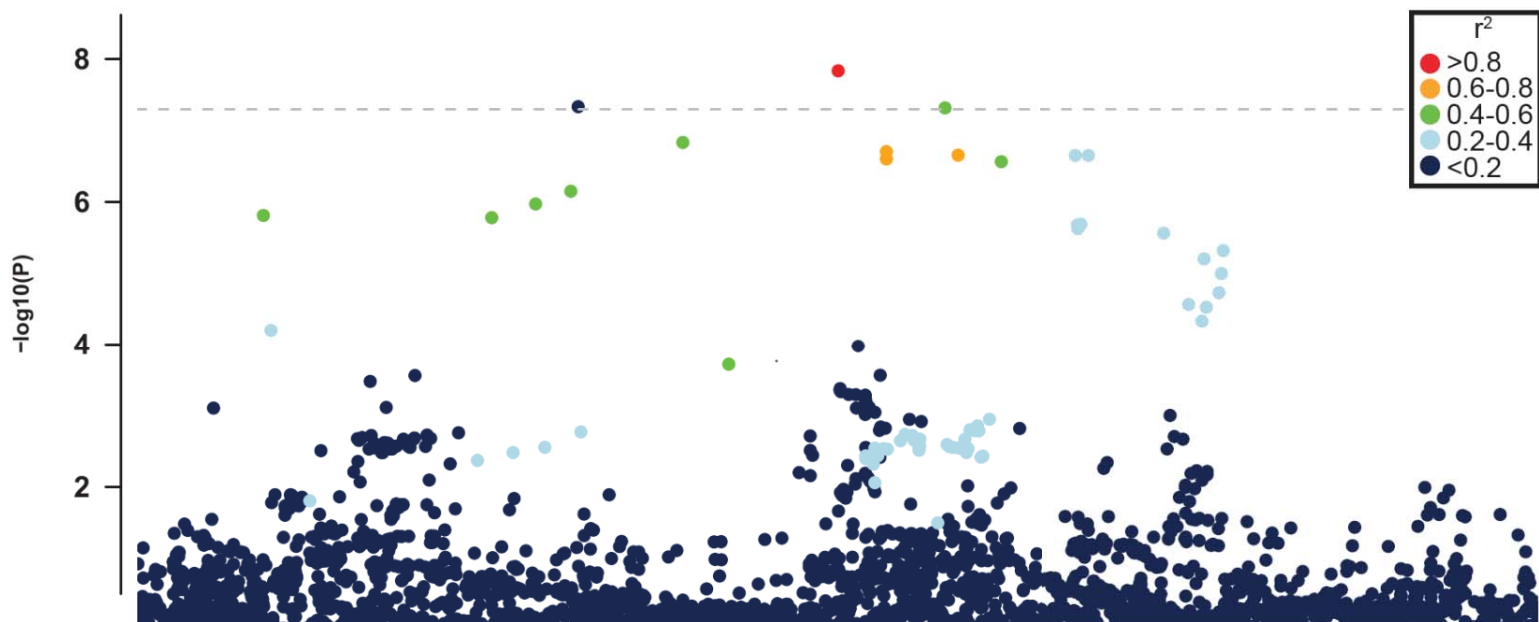
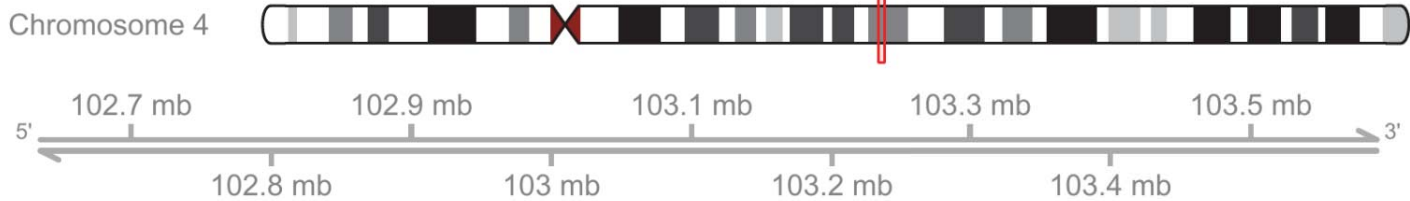
CP>GZ GZ>CP

Corrected for total SA

Uncorrected



Temporal Pole SA: rs6855246

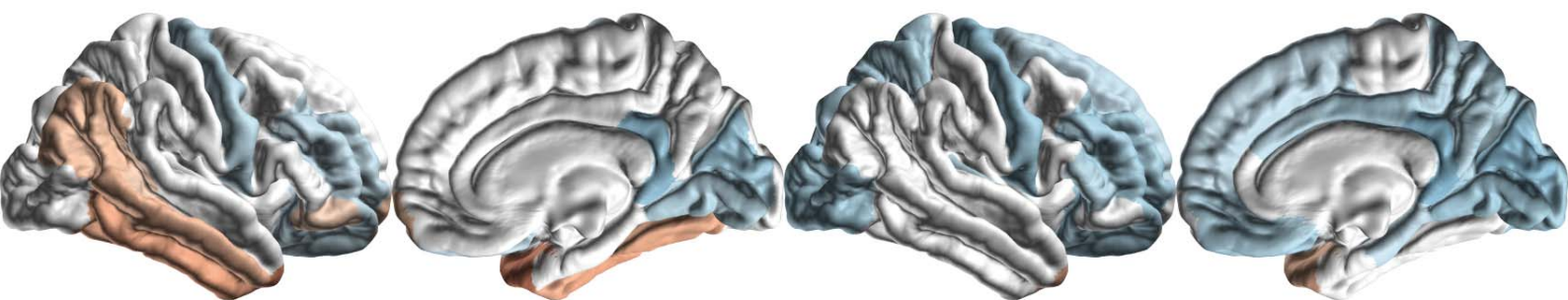


GZ>CP

CP>GZ

Corrected for total SA

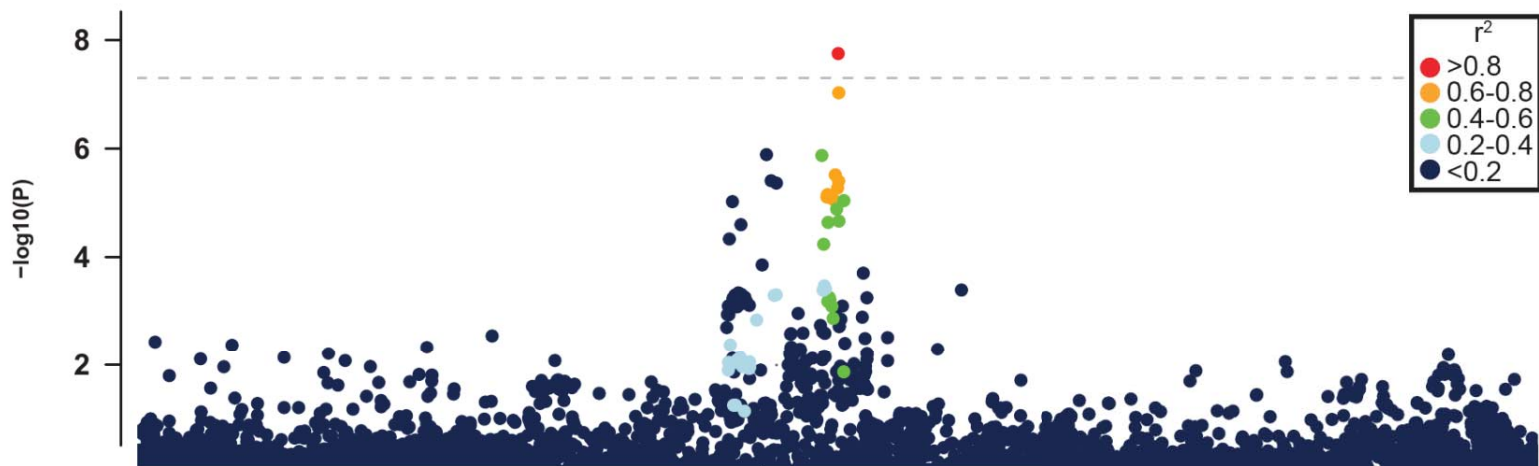
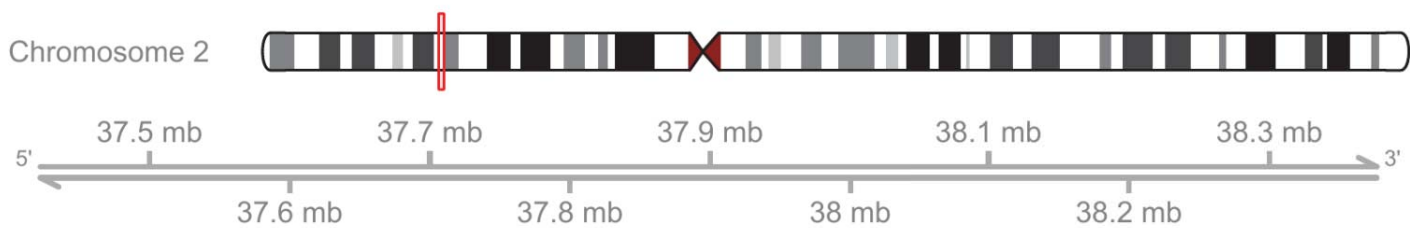
Uncorrected



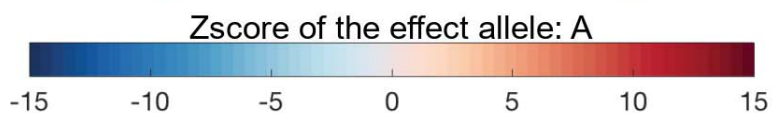
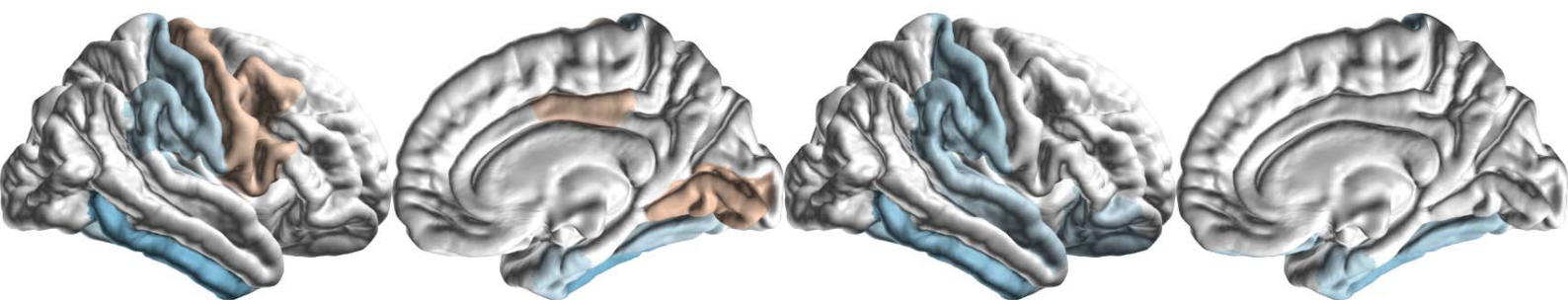
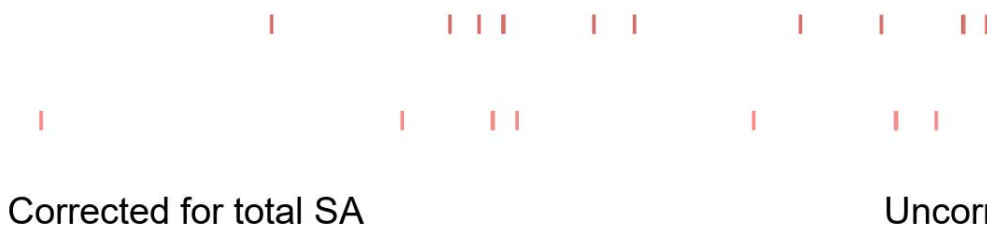
Zscore of the effect allele: A



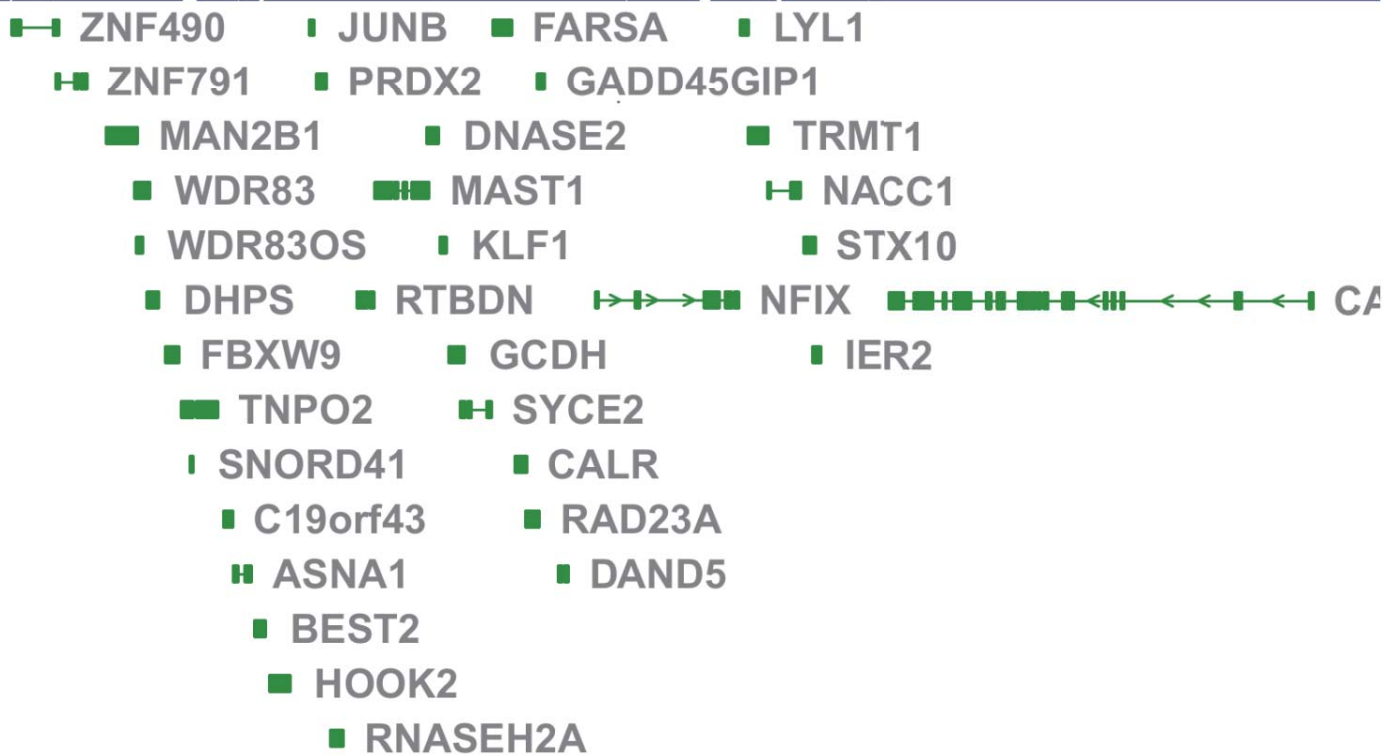
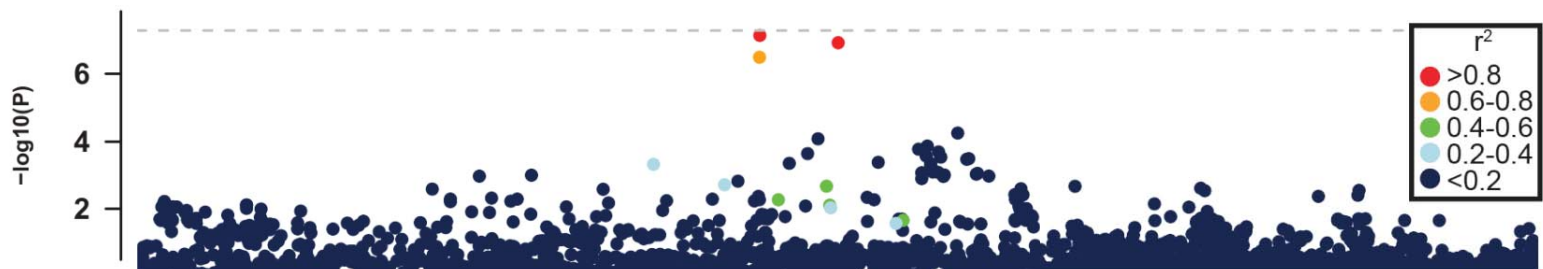
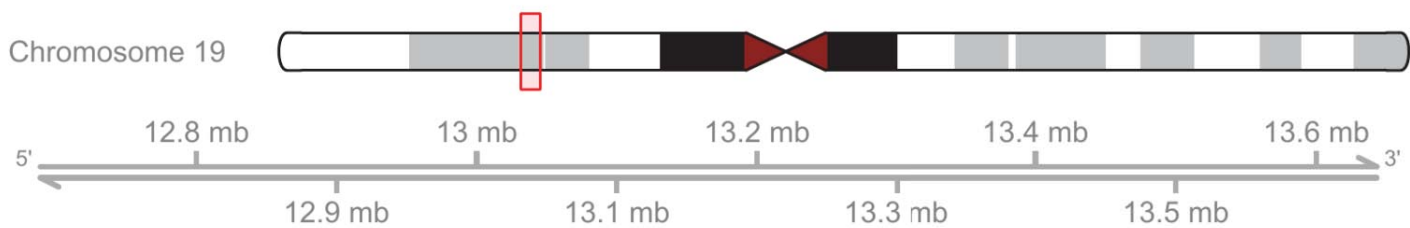
Inferior Temporal SA: rs9309013



CP>GZ GZ>CP



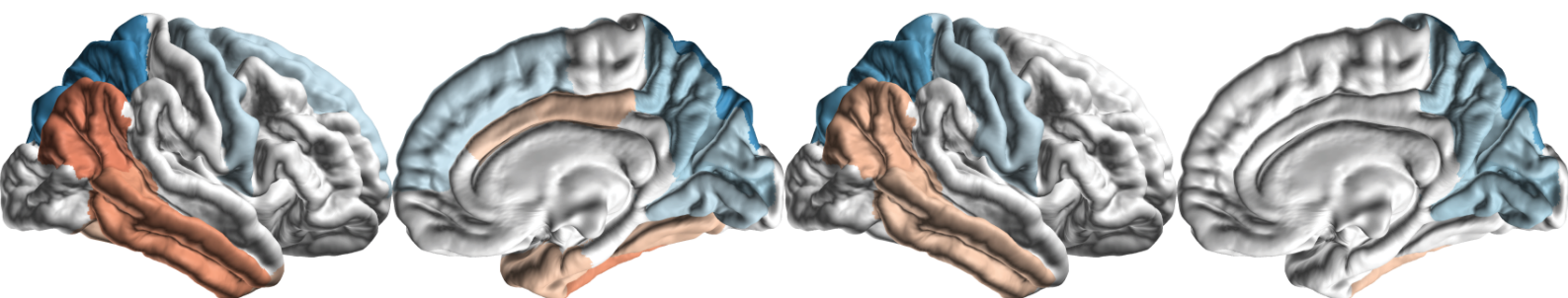
Inferior Temporal SA: rs73006837



CP>GZZ>CP

Corrected for total SA

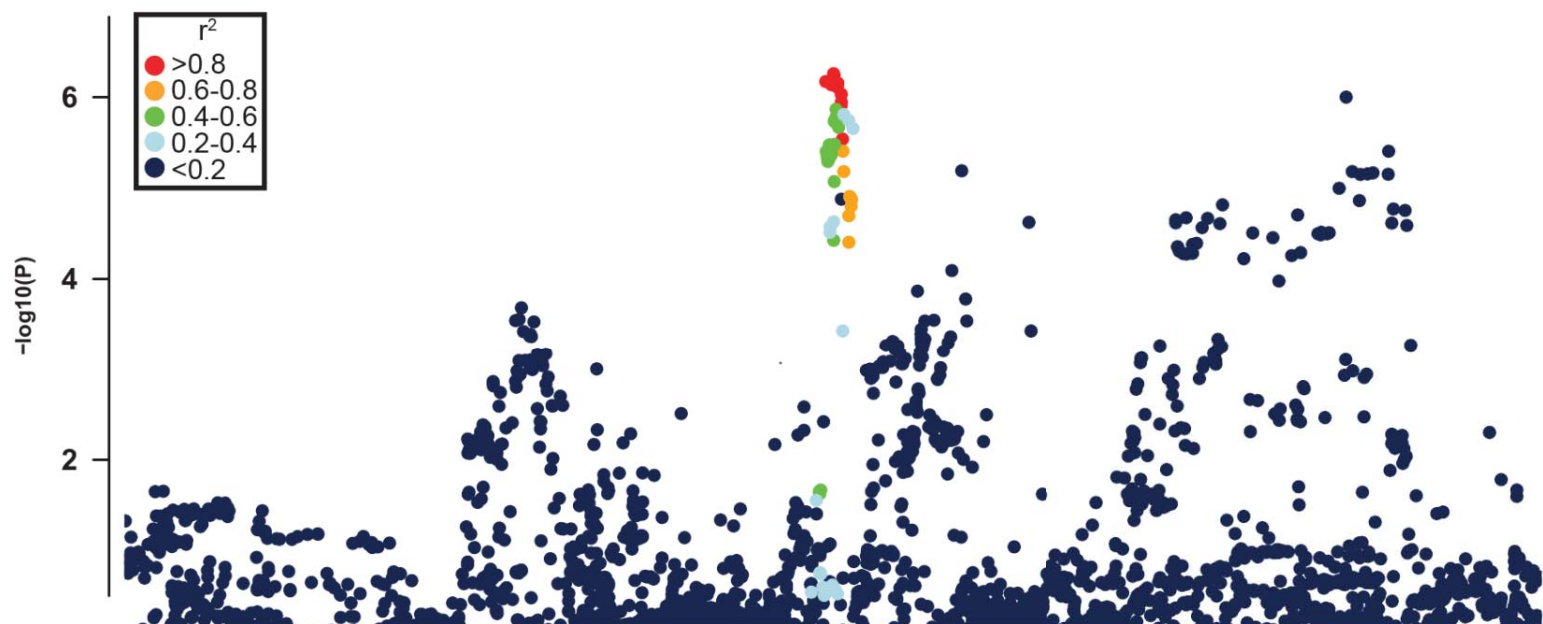
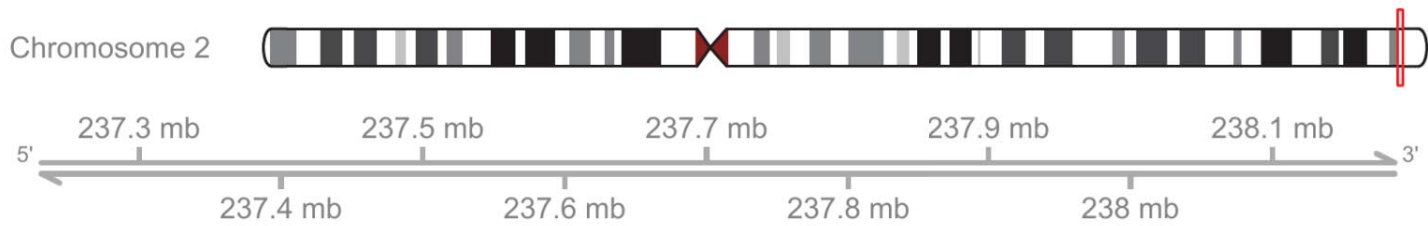
Uncorrected



Zscore of the effect allele: T

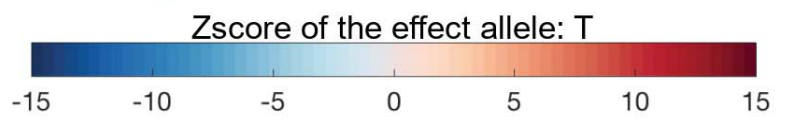
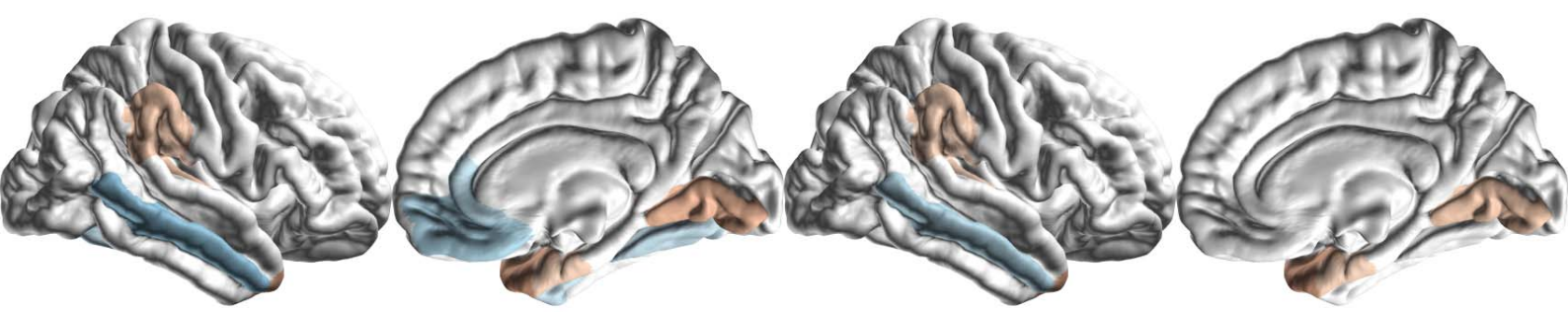


Middle Temporal SA: rs1344762

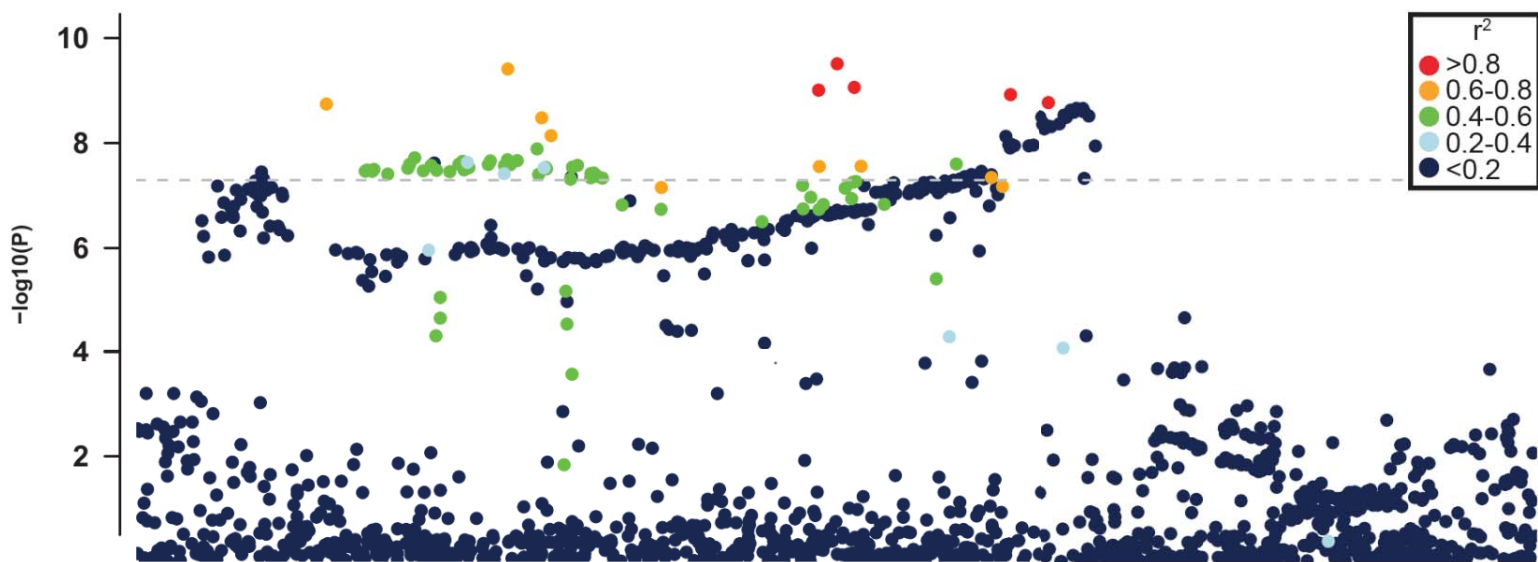
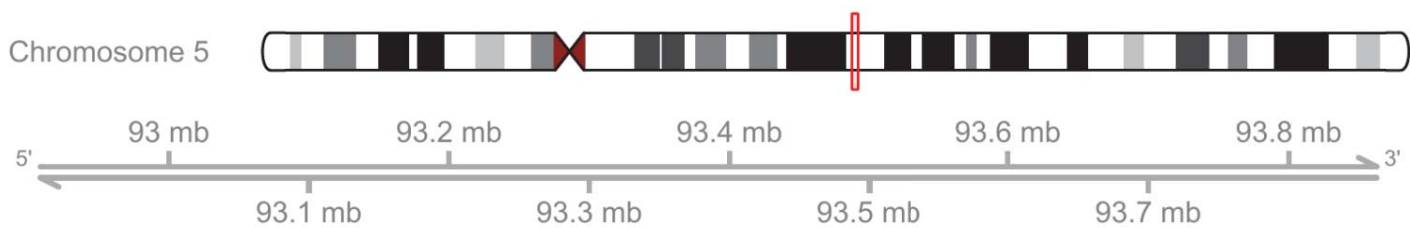


Corrected for total SA

Uncorrected



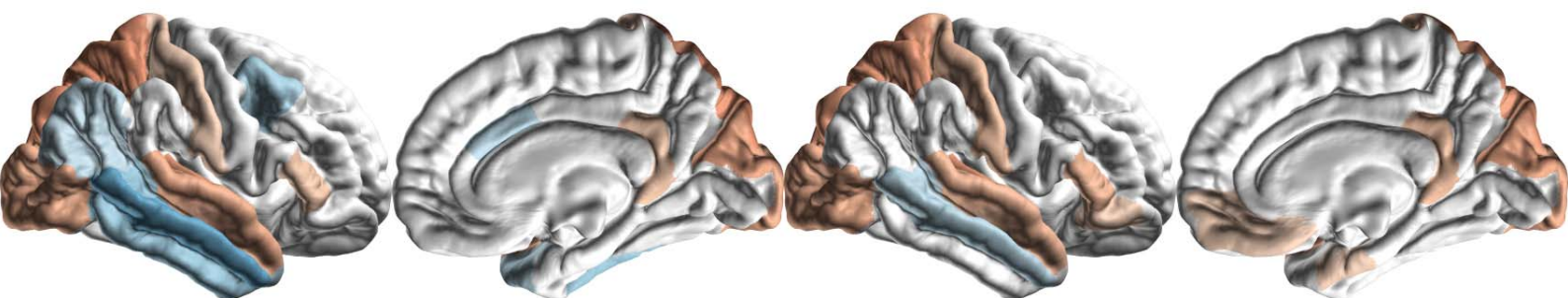
Middle Temporal SA: rs141834426



GZ>CP
CP>GZ

Corrected for total SA

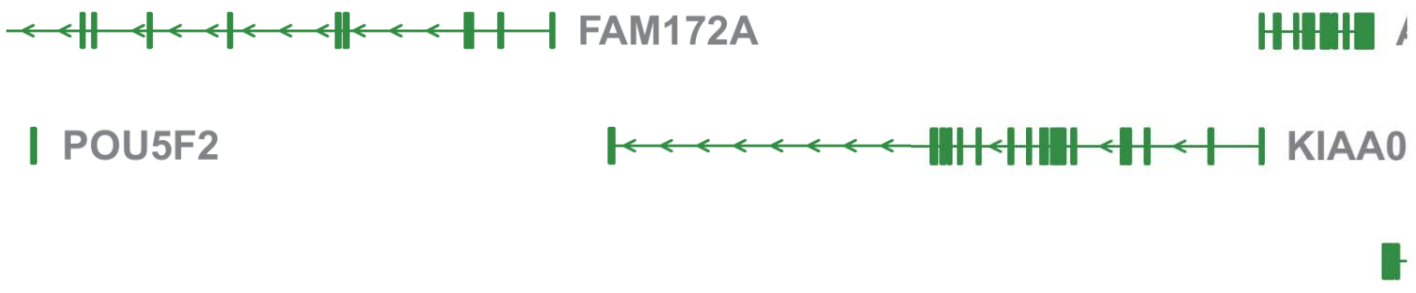
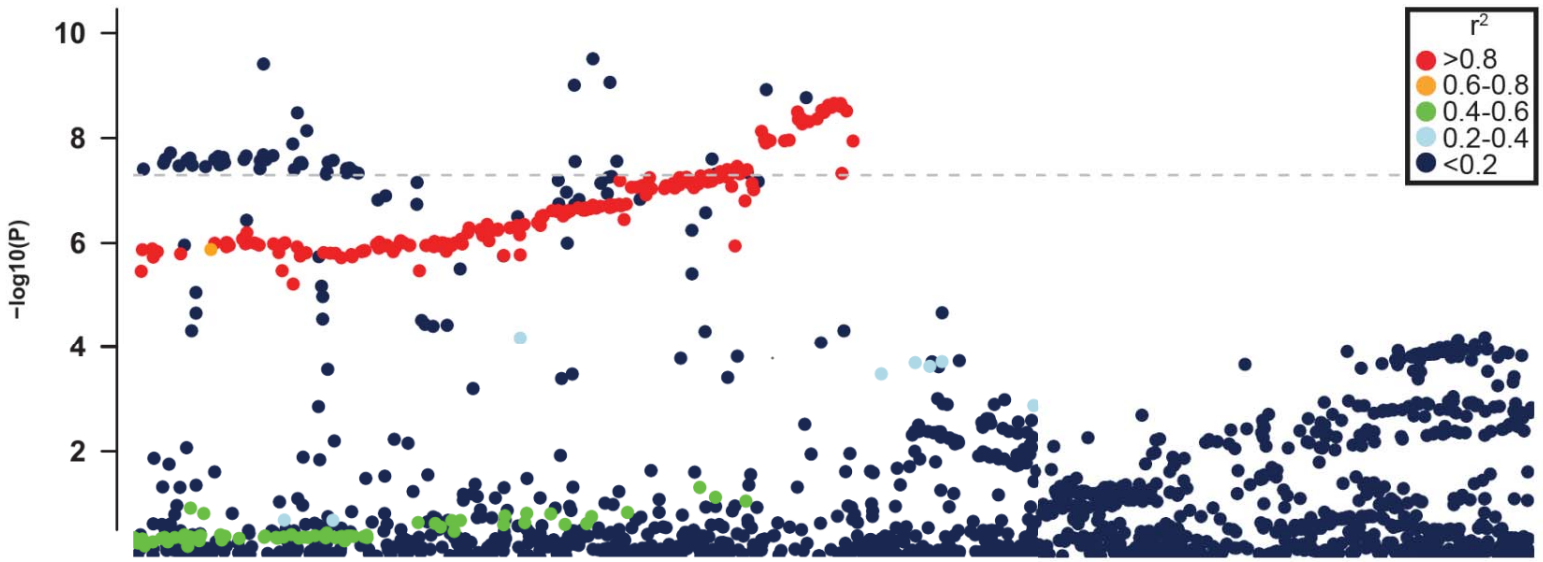
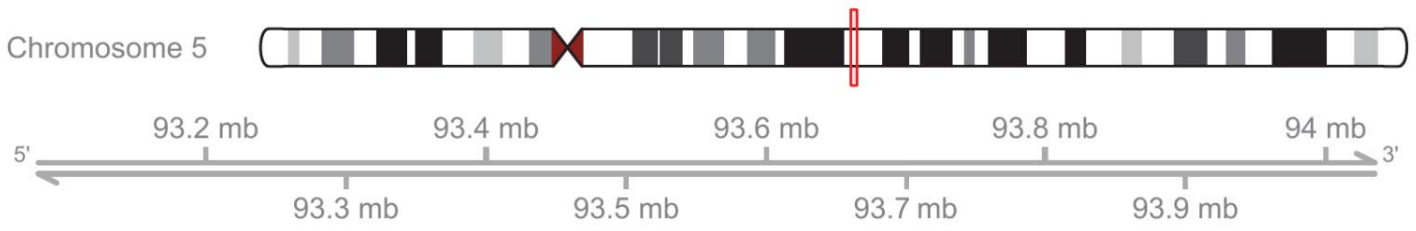
Uncorrected



Zscore of the effect allele: C

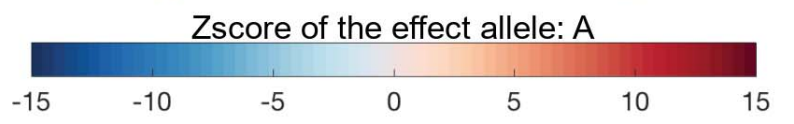
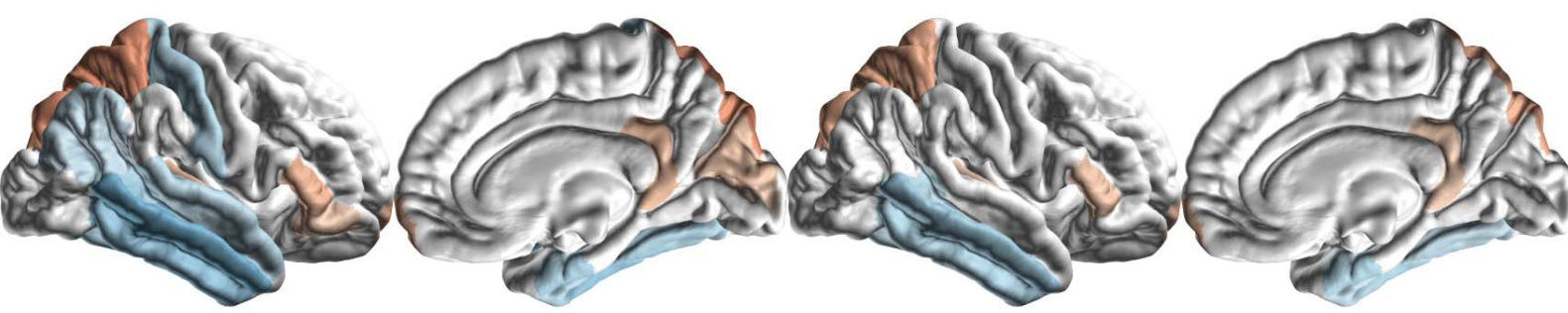


Middle Temporal SA: rs17376456



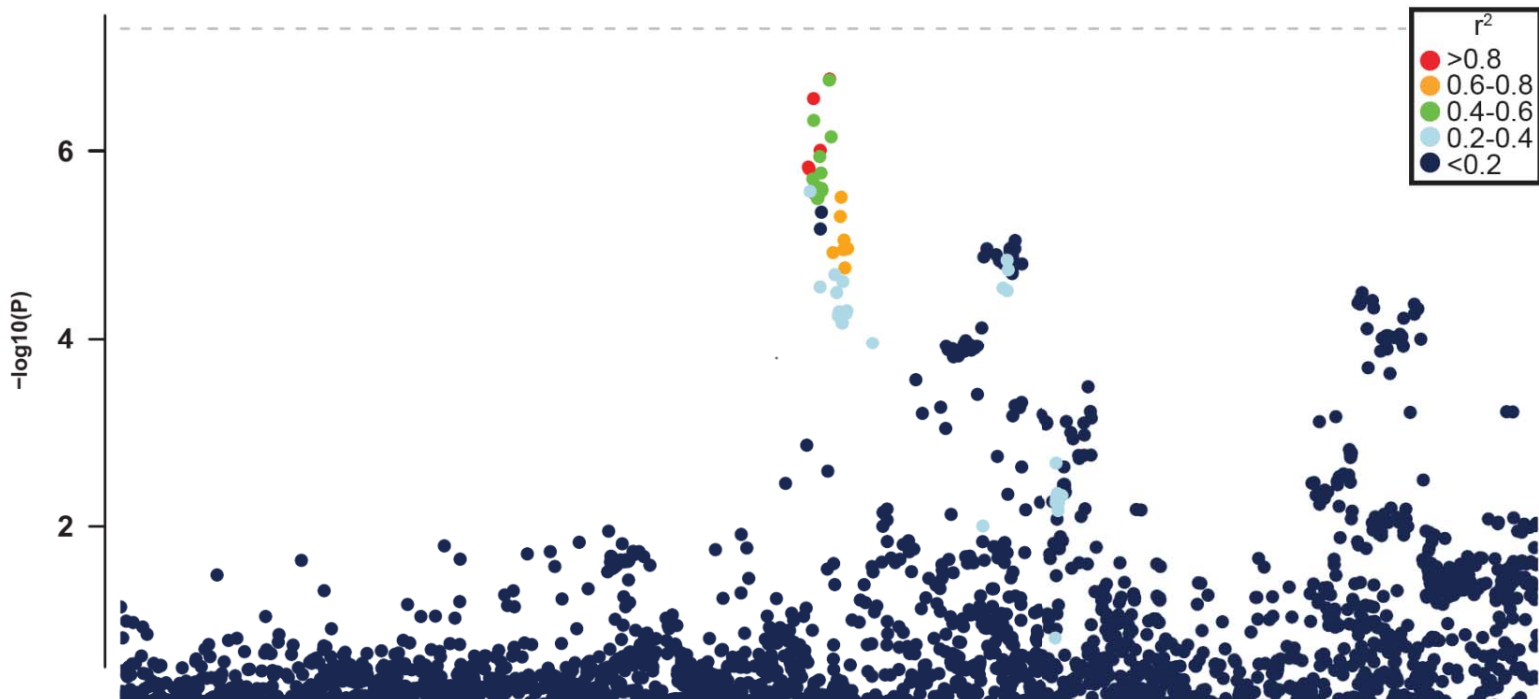
Corrected for total SA

Uncorrected



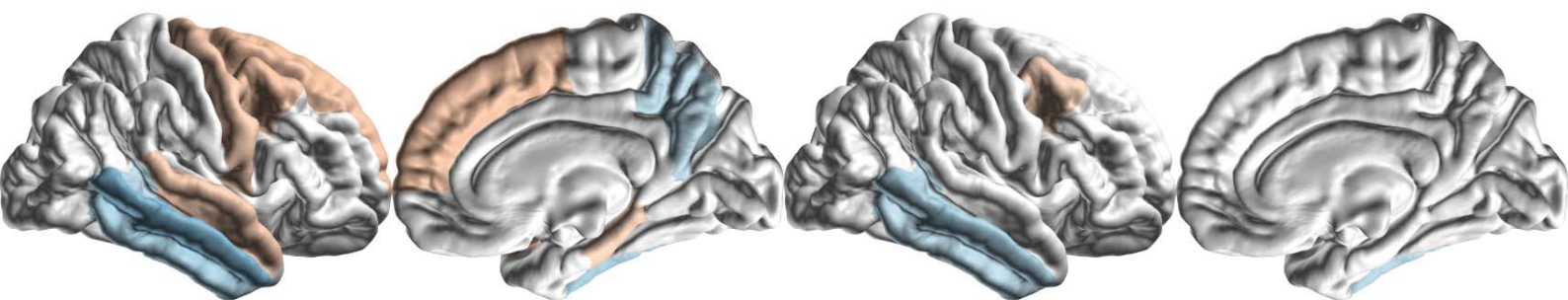
Middle Temporal SA: rs9328010

Chromosome 5

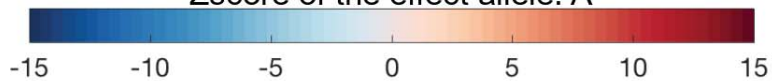


Corrected for total SA

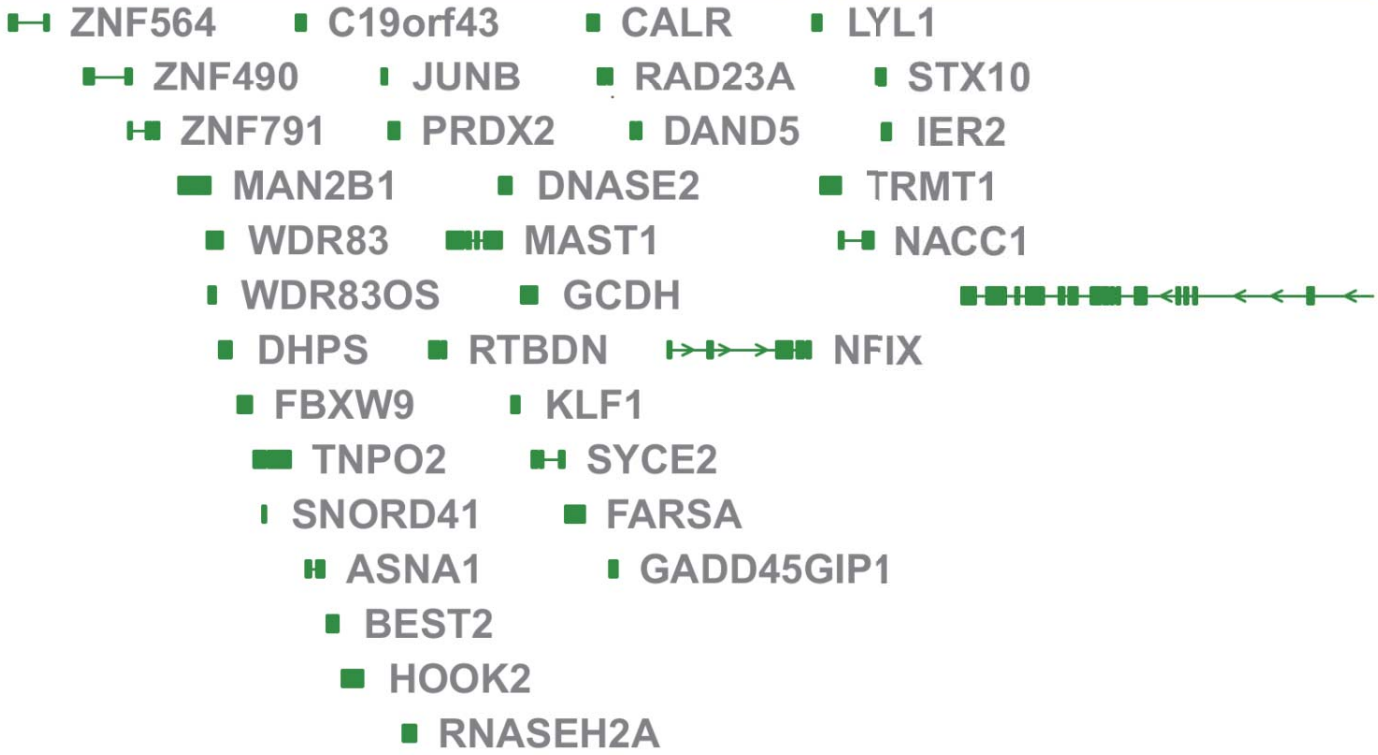
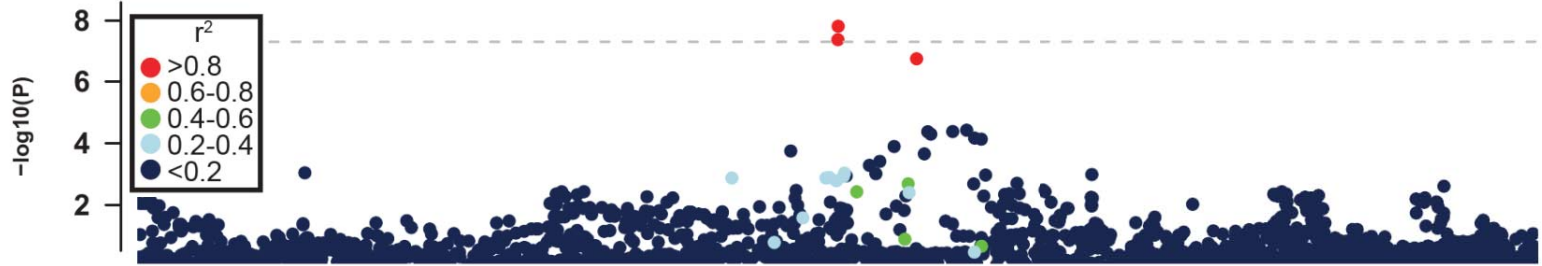
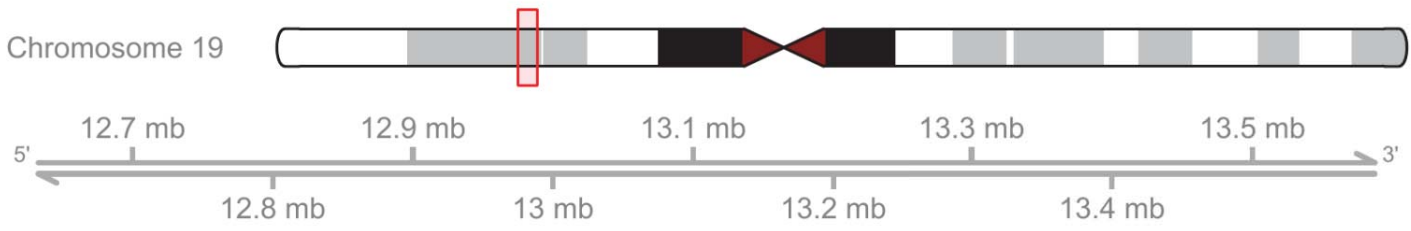
Uncorrected



Zscore of the effect allele: A



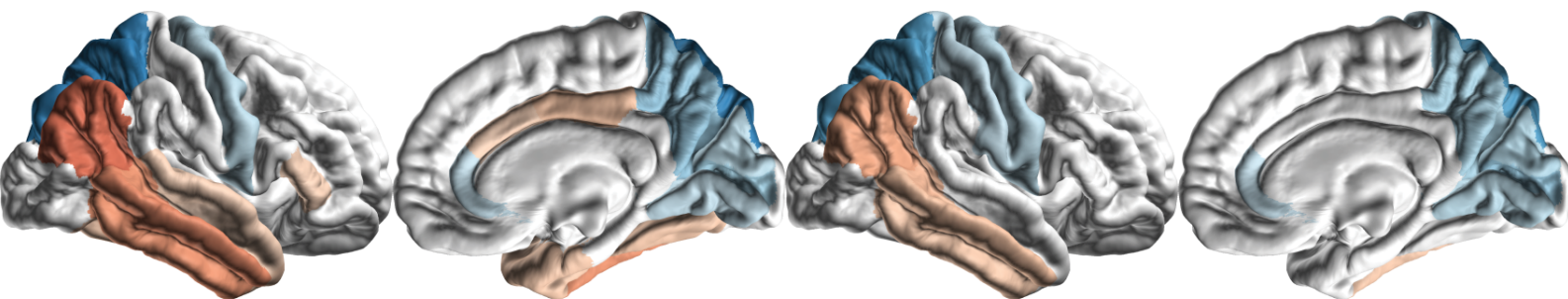
Middle Temporal SA: rs73006822



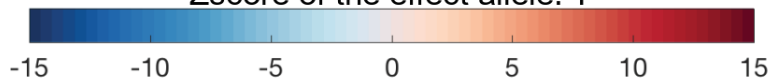
CP>GZZ>CP

Corrected for total SA

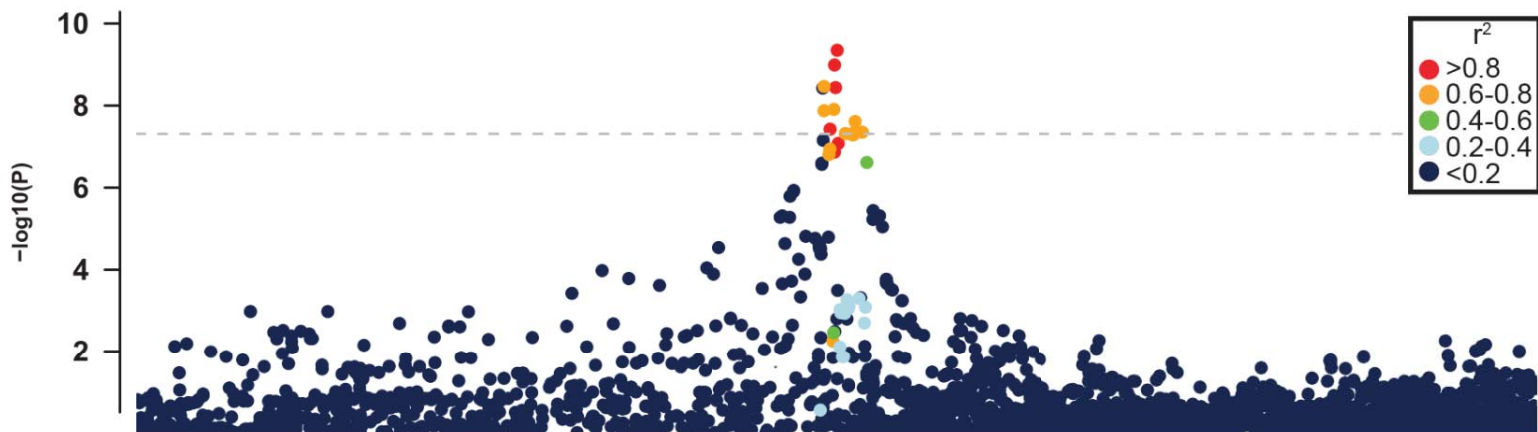
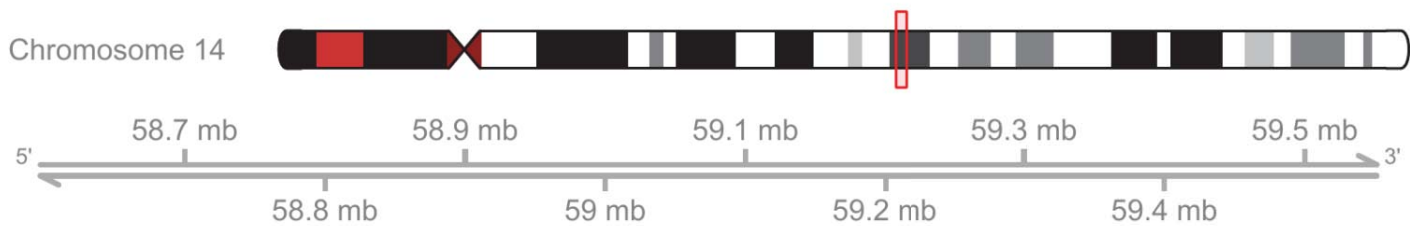
Uncorrected



Zscore of the effect allele: T



Middle Temporal TH: rs160459



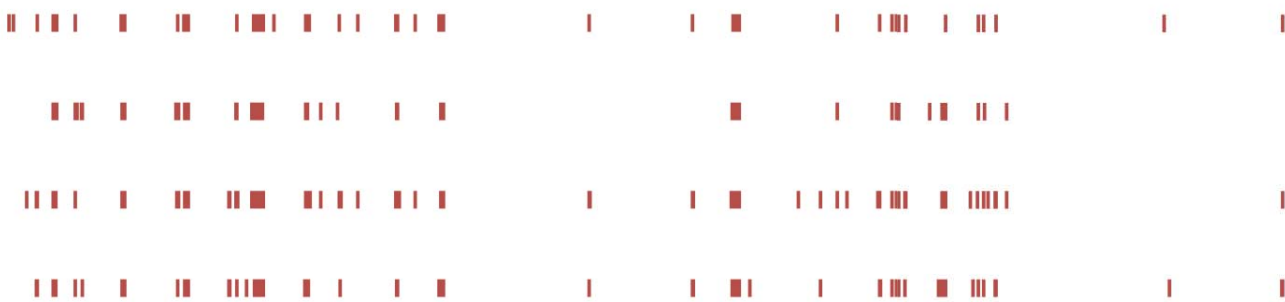
C14orf37 ARID4A DACT1

ACTR10 TOMM20L

PSMA3 TIMM9

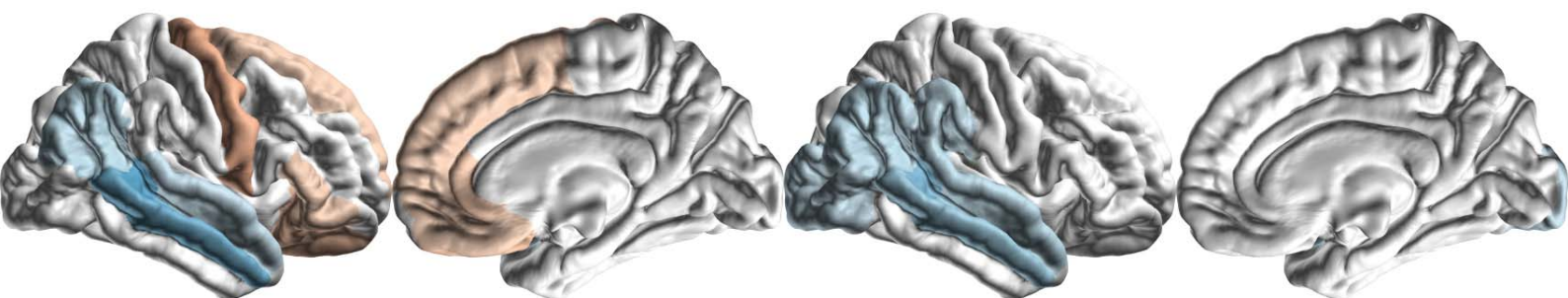
KIAA0586

Angular temporal dIPFC >ingulate

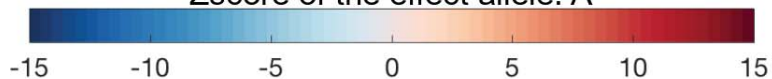


Corrected for average TH

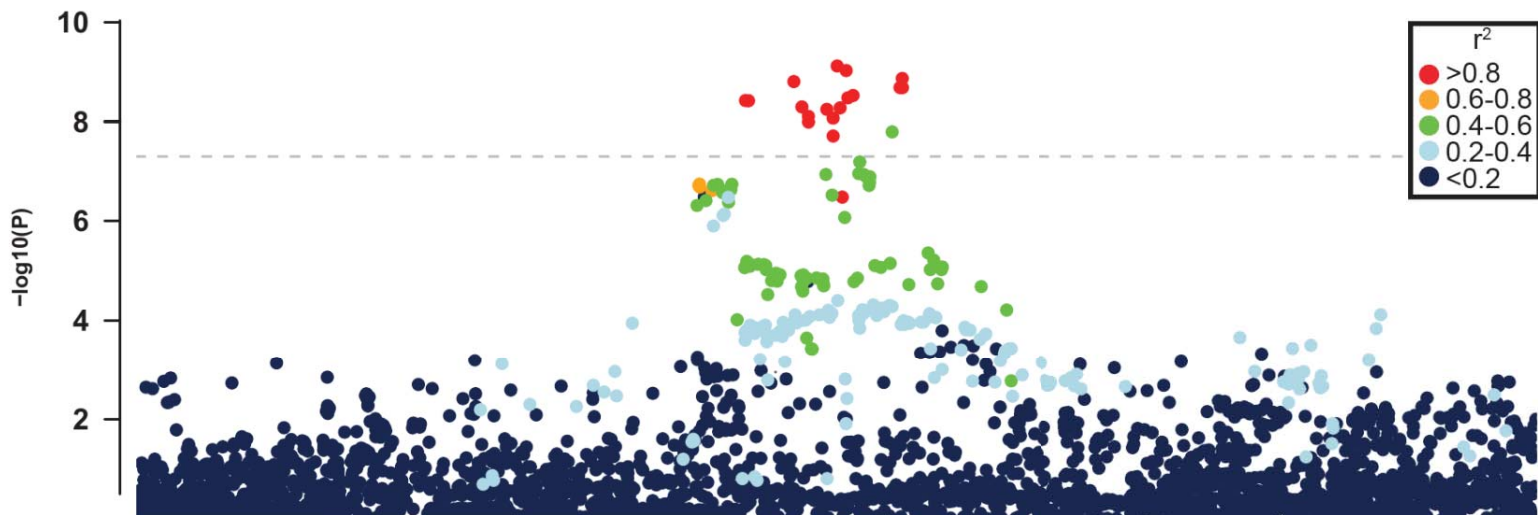
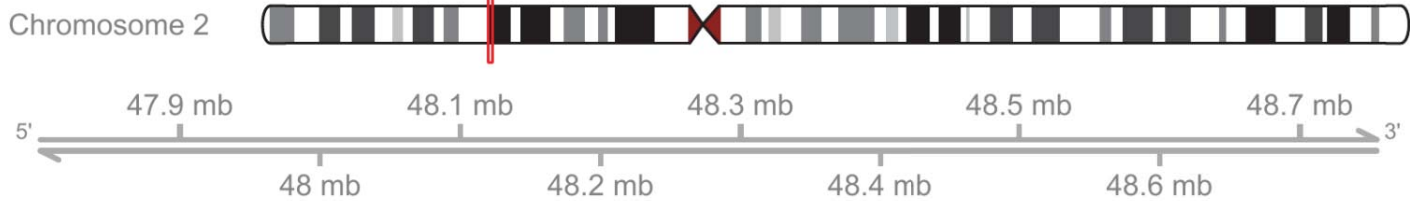
Uncorrected



Zscore of the effect allele: A

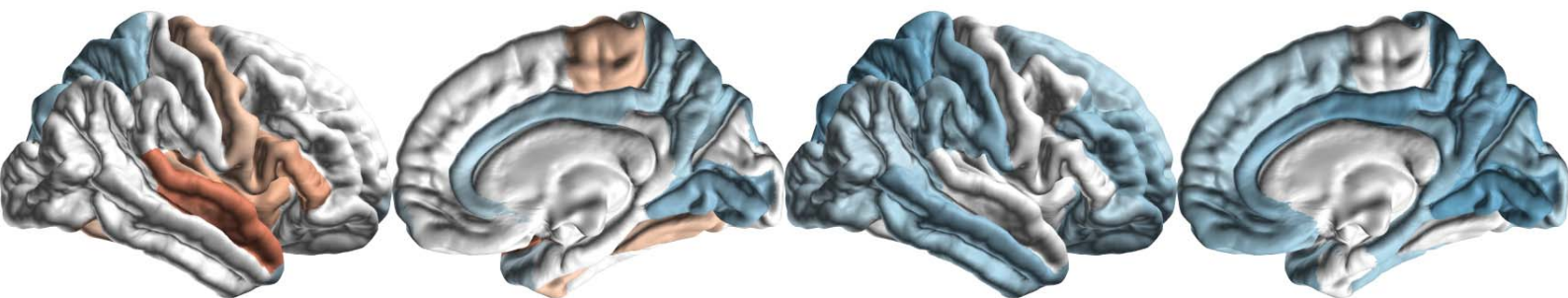


Superior Temporal SA: rs7601767

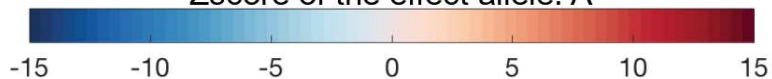


Corrected for total SA

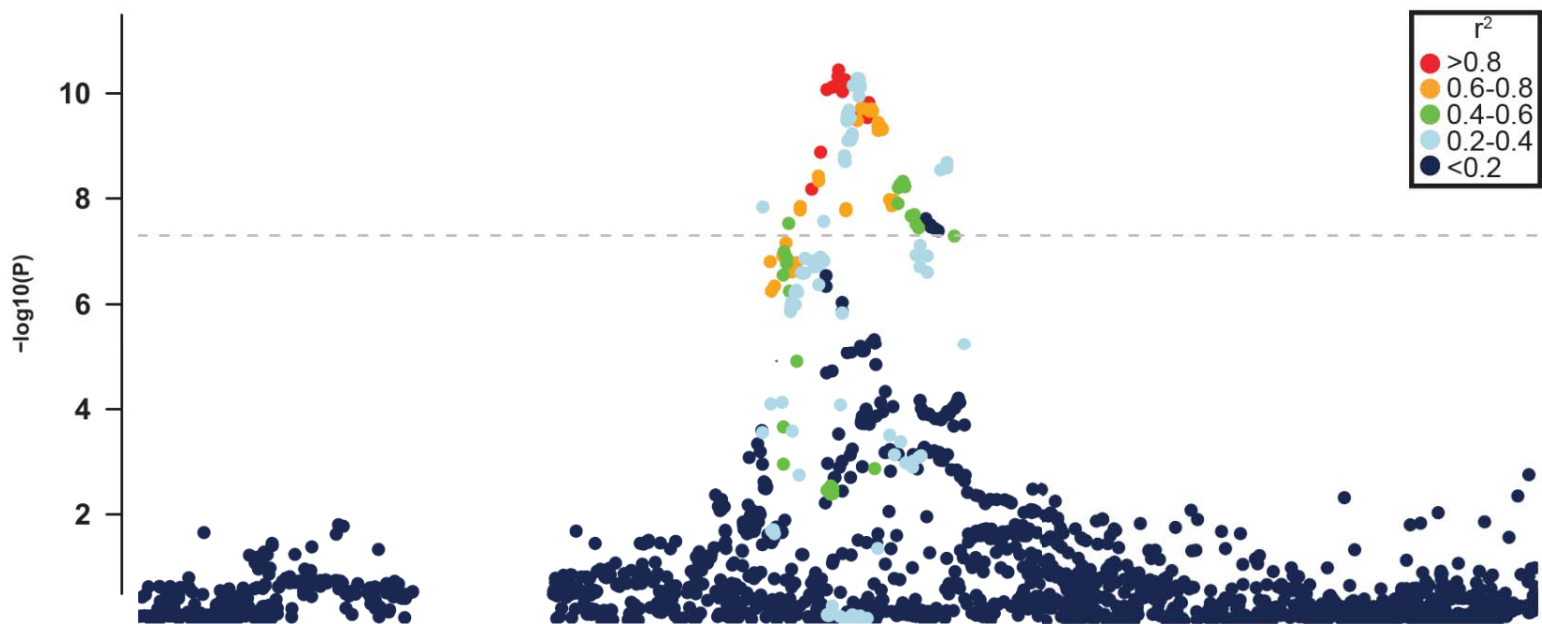
Uncorrected



Zscore of the effect allele: A



Superior Temporal SA: rs13011264



EPC2

KIF5C

LYPD6

LYPD6B

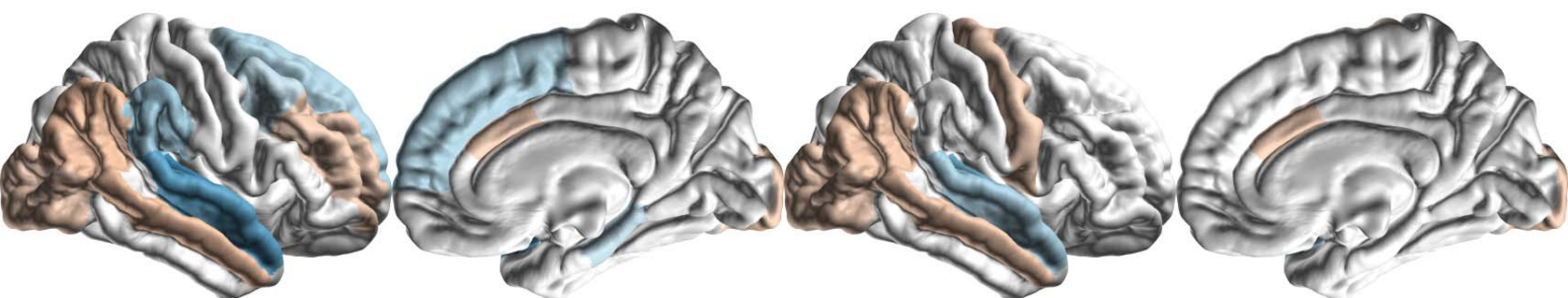
MI

GZ>CP

CP>GZ

Corrected for total SA

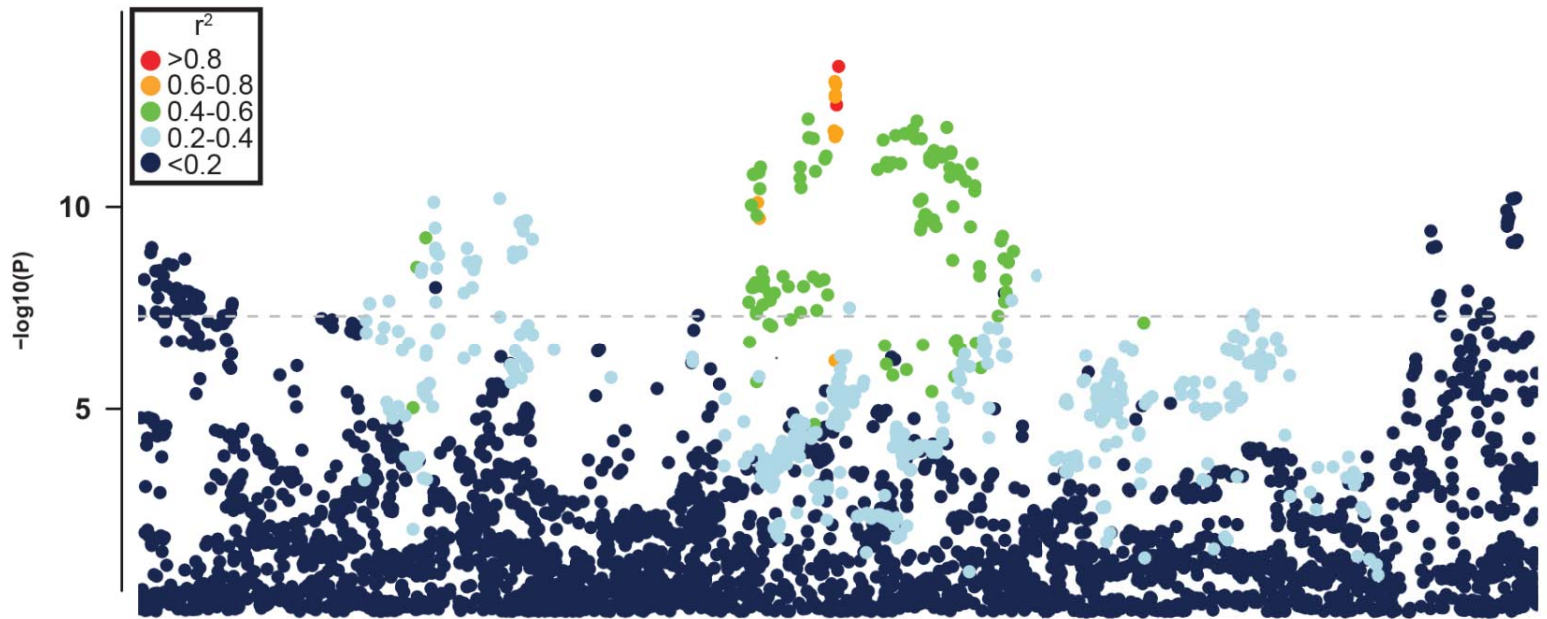
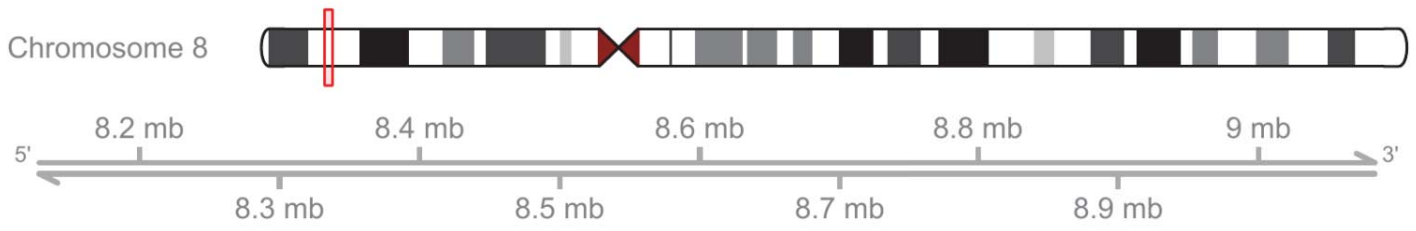
Uncorrected



Zscore of the effect allele: A

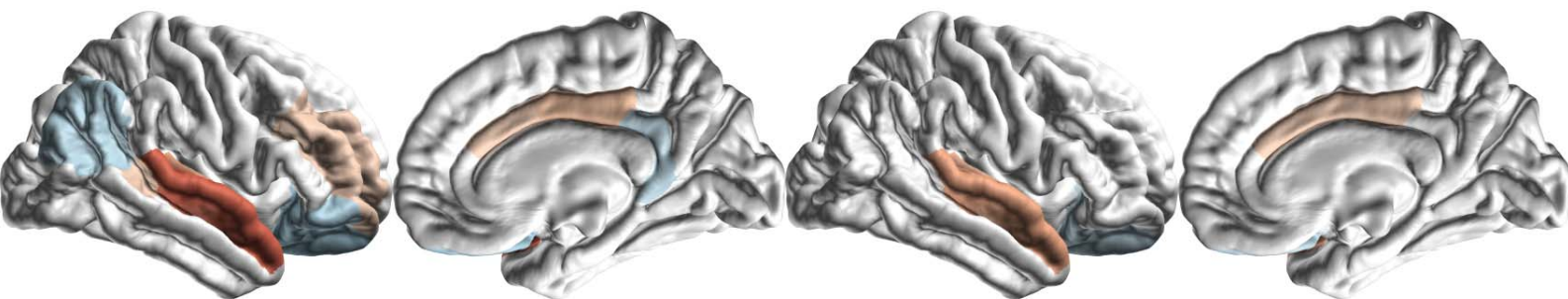


Superior Temporal SA: rs4841029

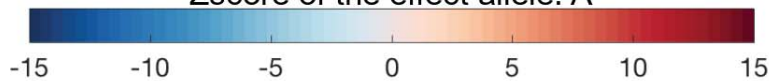


Corrected for total SA

Uncorrected

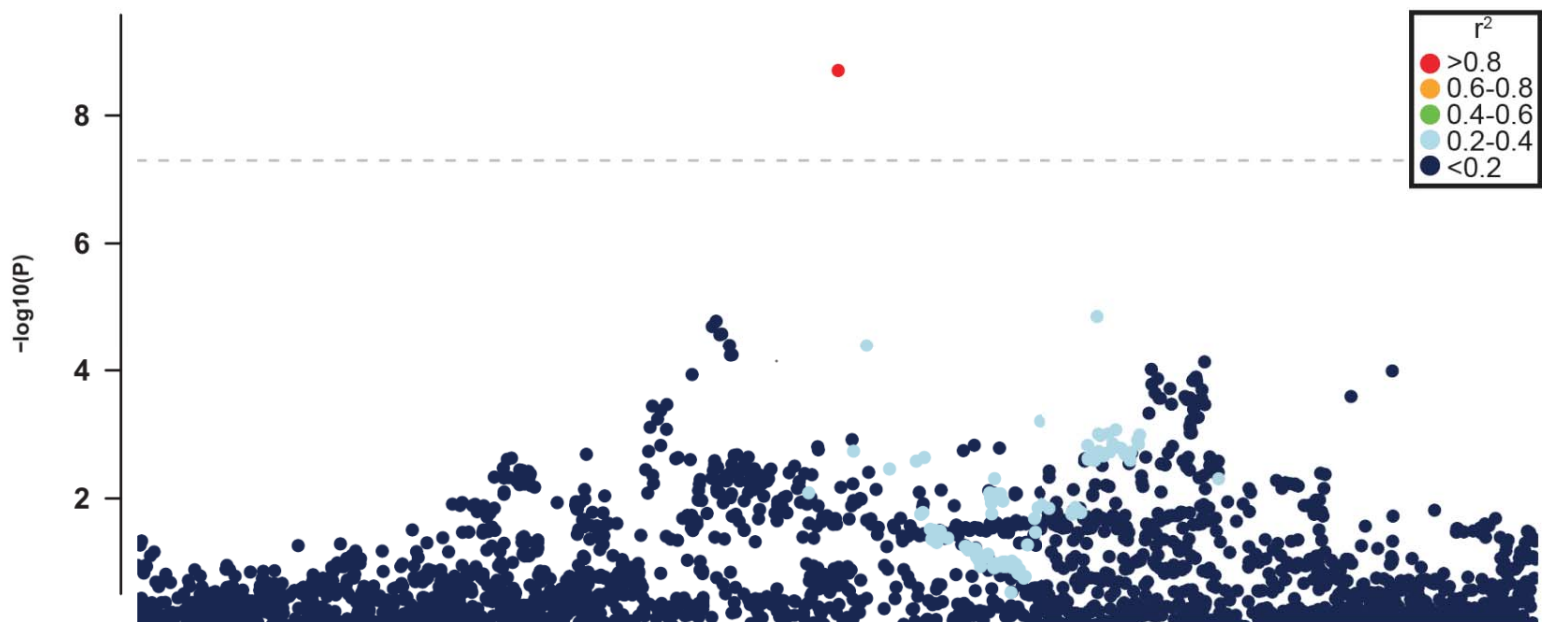
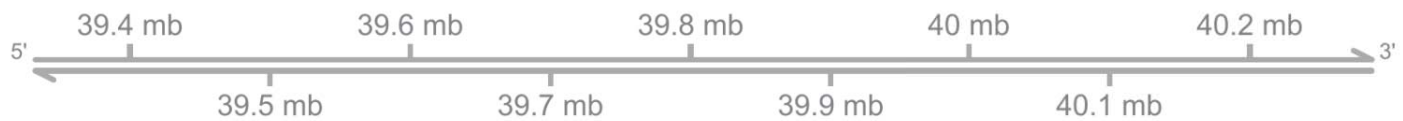


Zscore of the effect allele: A



Superior Temporal SA: rs115241741

Chromosome 15



C15orf54

THBS1

GPR17

FSIP1

GZ>CP

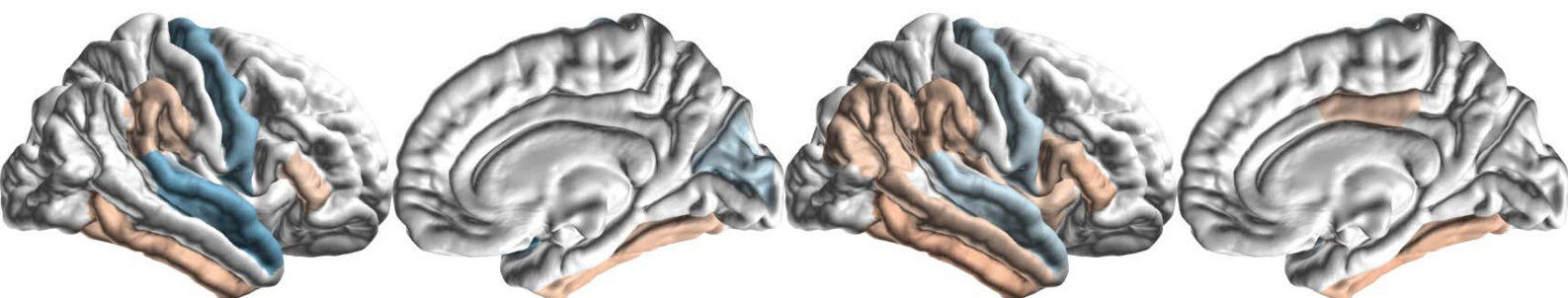


CP>GZ



Corrected for total SA

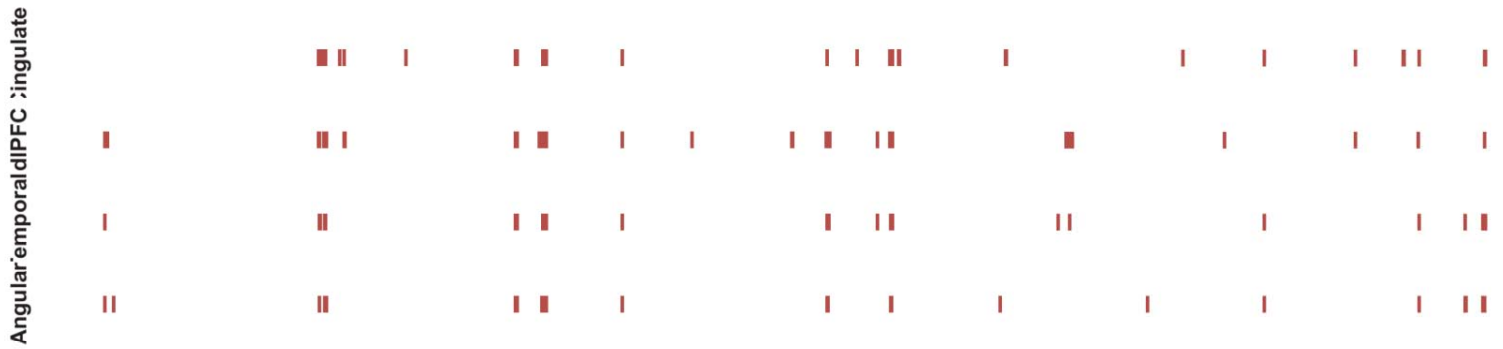
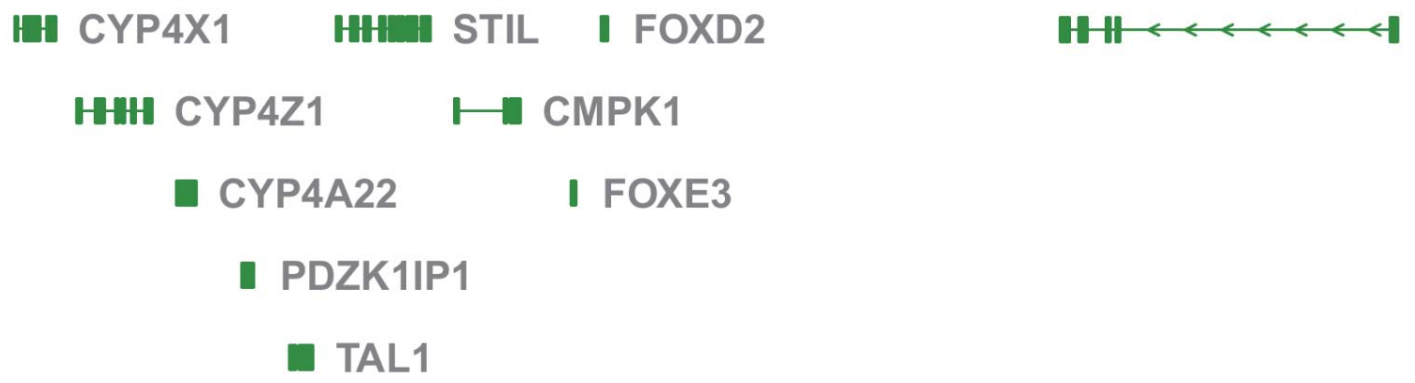
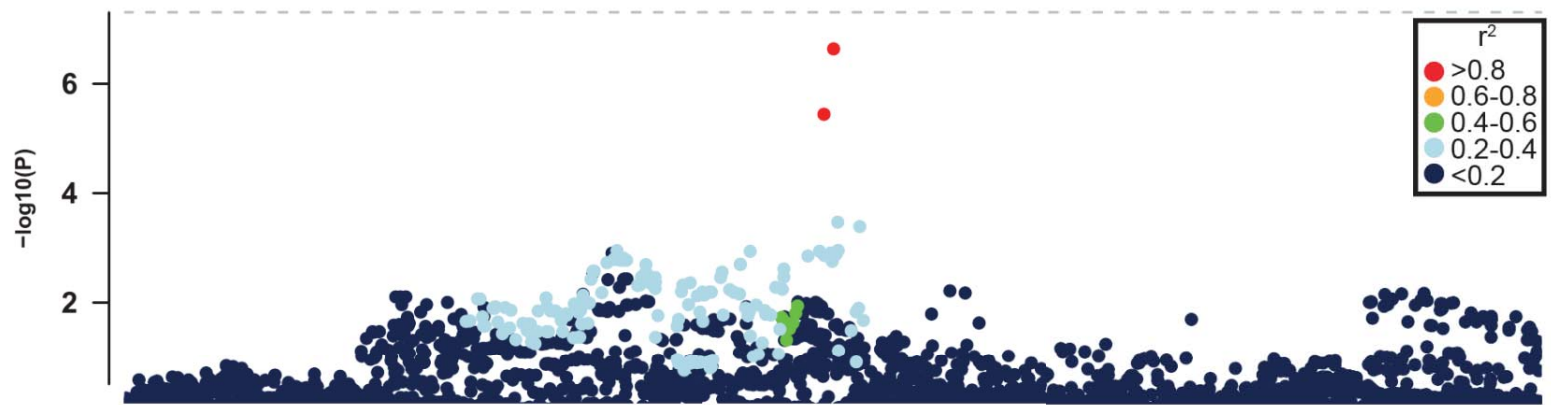
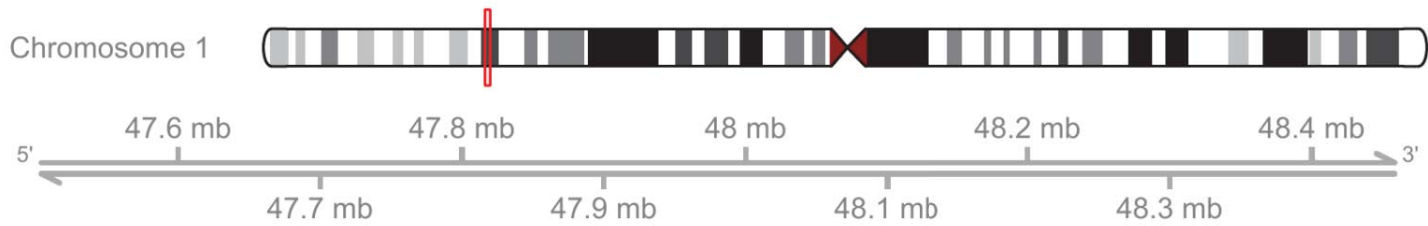
Uncorrected



Zscore of the effect allele: A

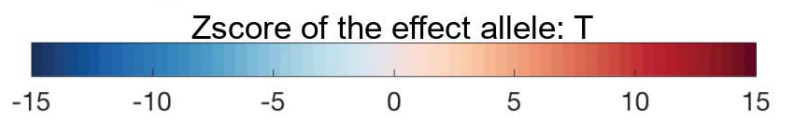
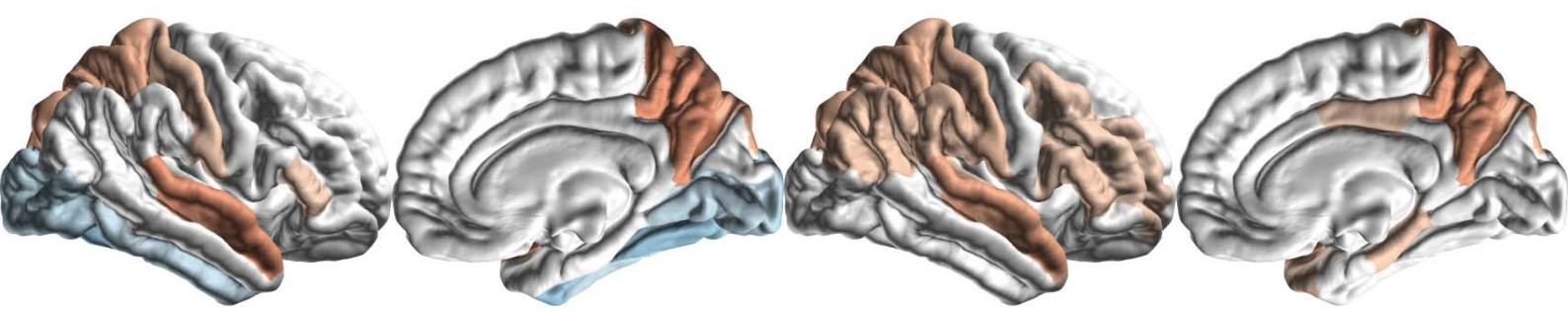


Superior Temporal TH: rs6658111

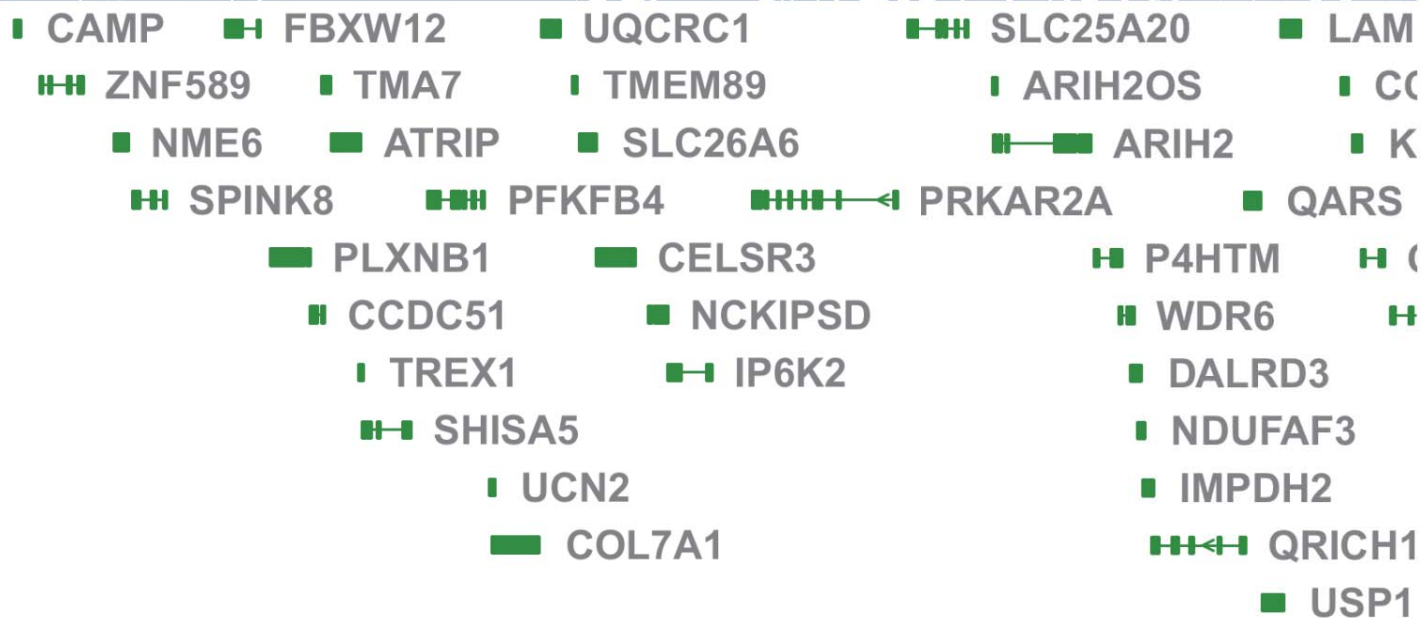
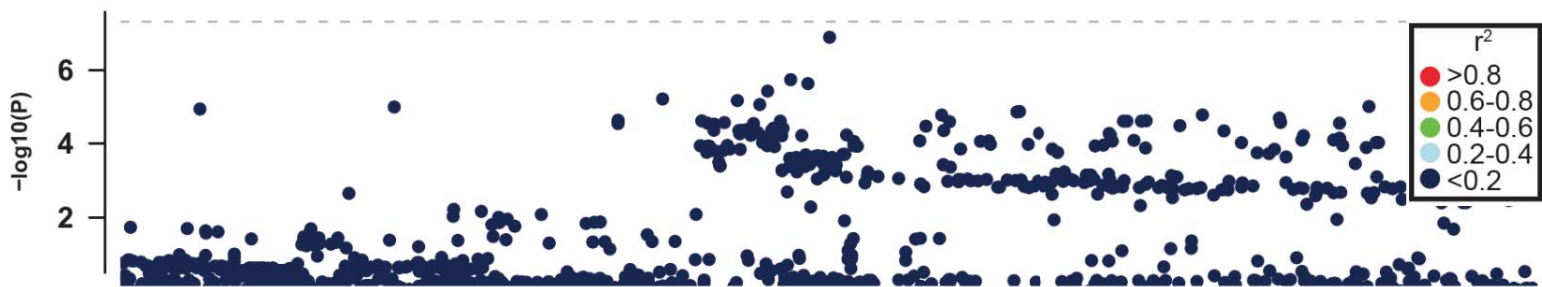
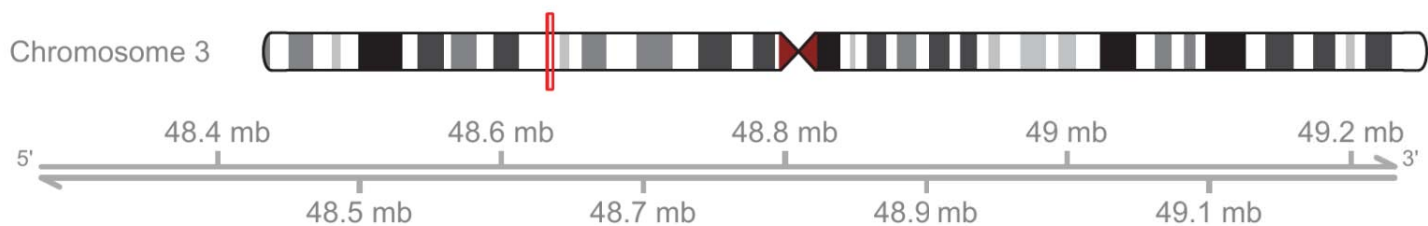


Corrected for average TH

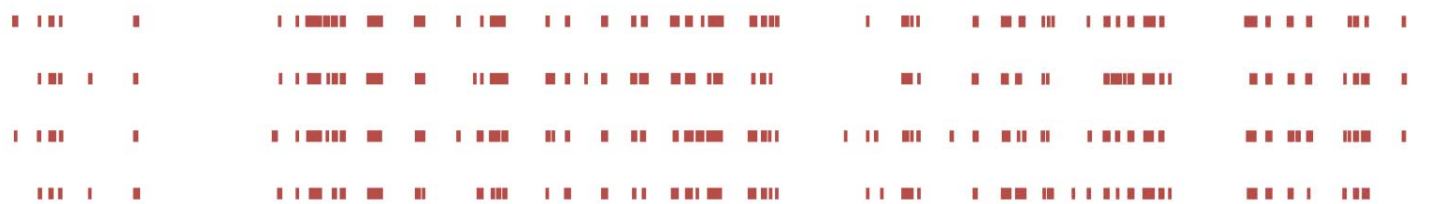
Uncorrected



Superior Temporal TH: rs9848203

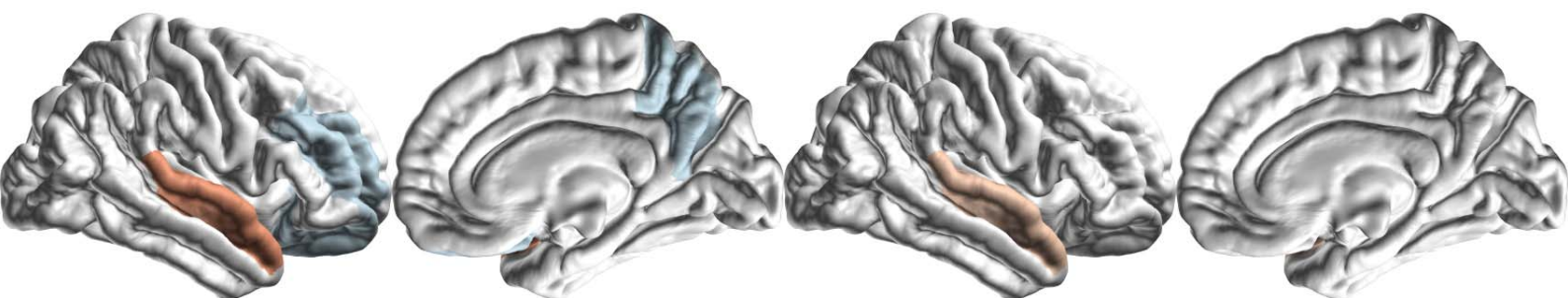


AngulamporziPFNgulate

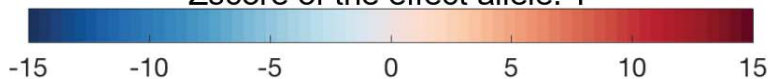


Corrected for average TH

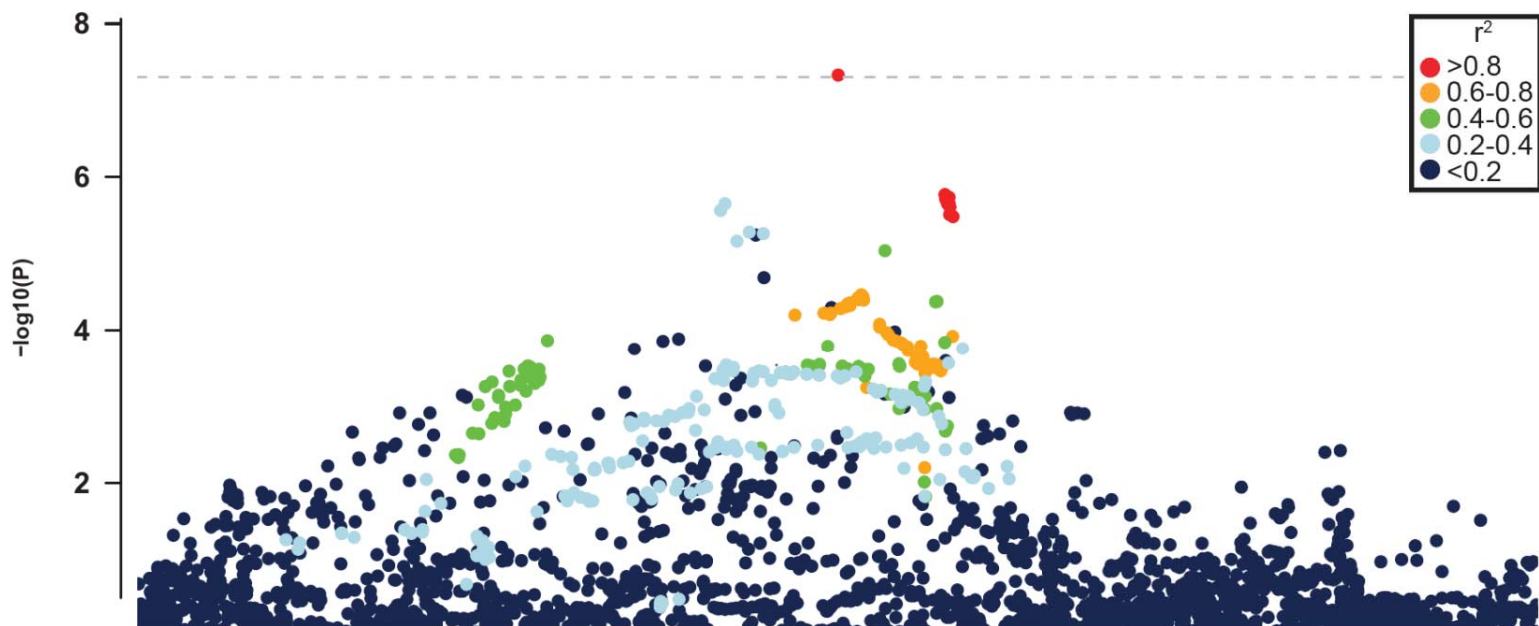
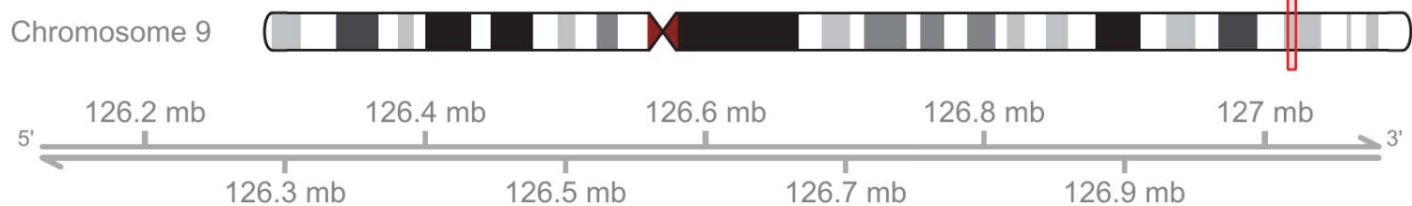
Uncorrected



Zscore of the effect allele: T



Banks of the Superior Temporal Sulcus SA: rs7862092



CRB2

LHX2

→

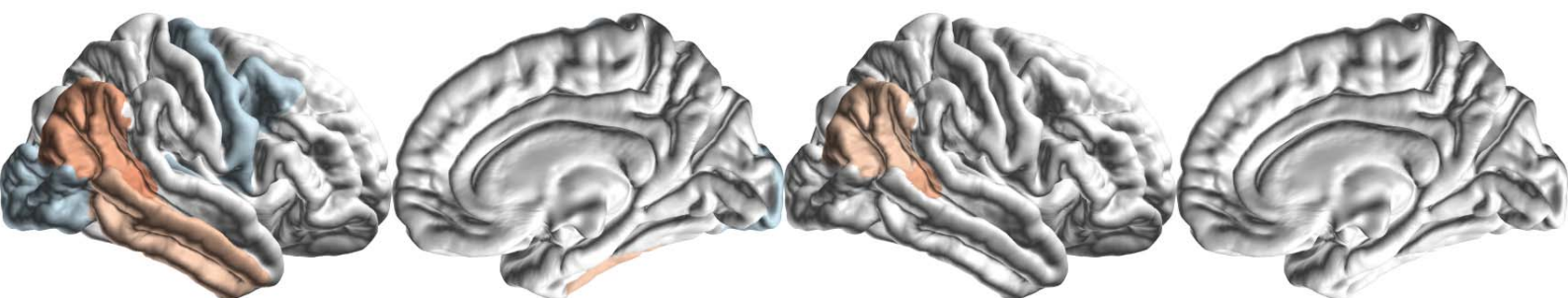
DENND1A

GZ>CP

CP>GZ

Corrected for total SA

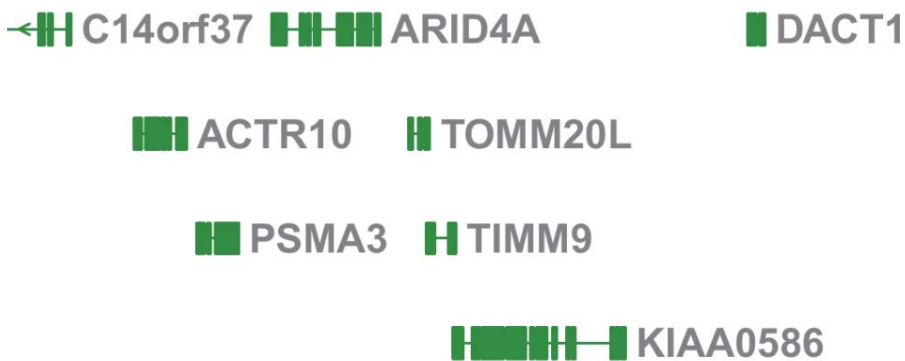
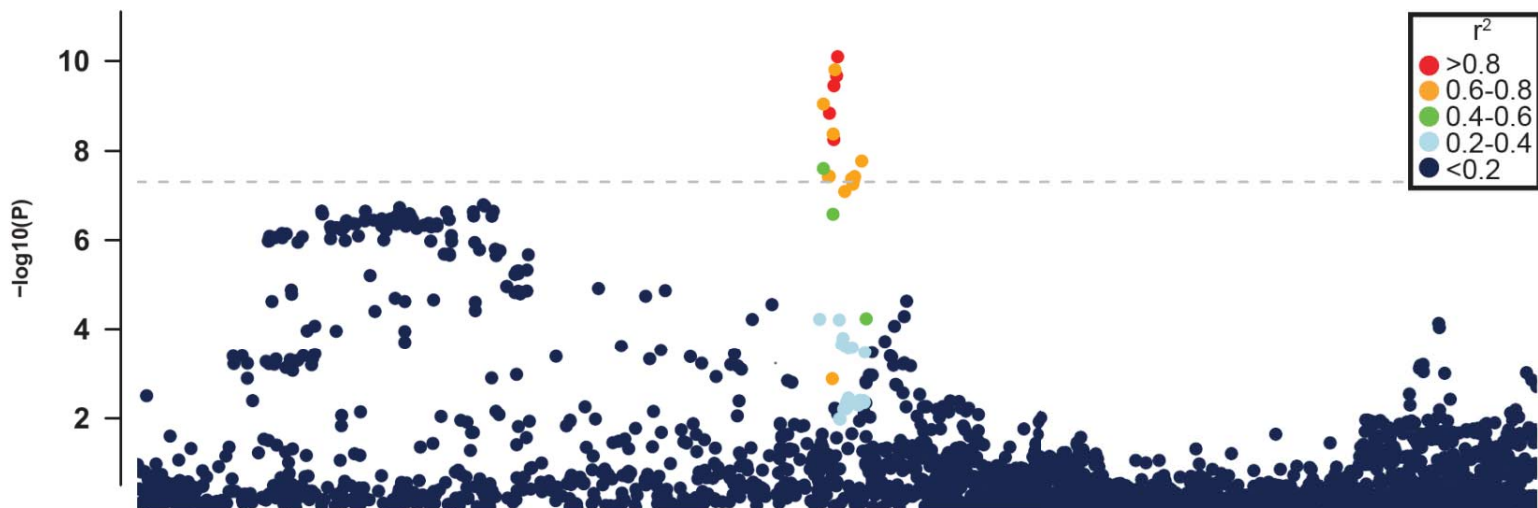
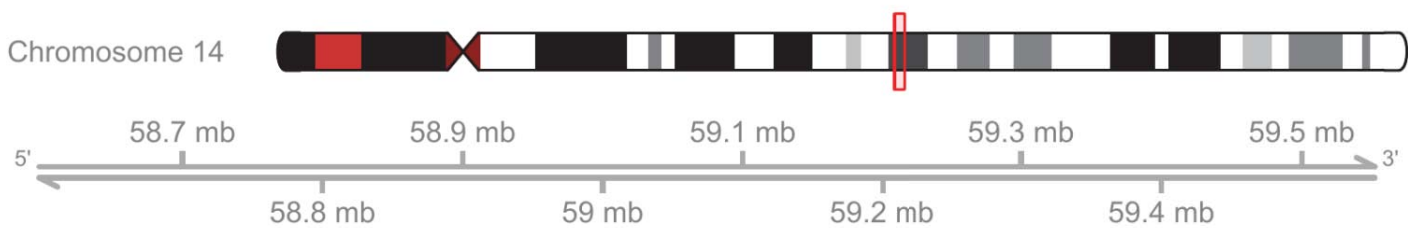
Uncorrected



Zscore of the effect allele: T

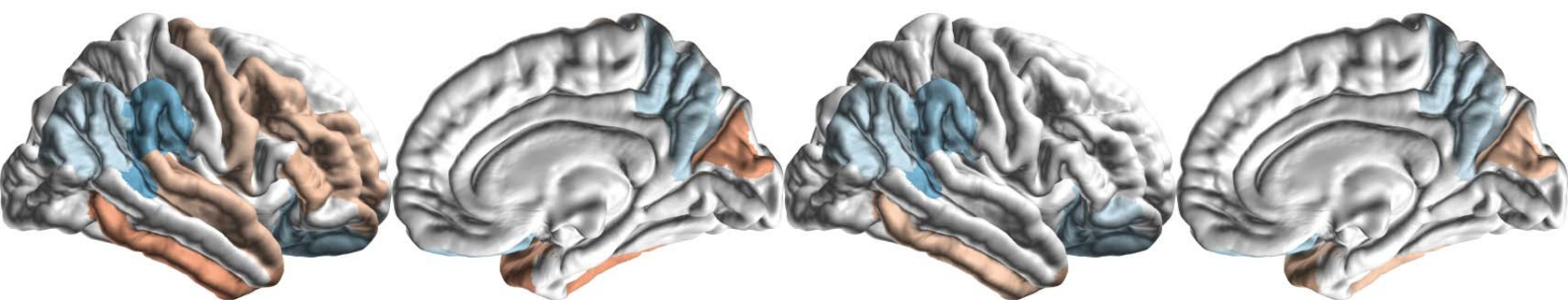


Banks of the Superior Temporal Sulcus SA: rs160458



Corrected for total SA

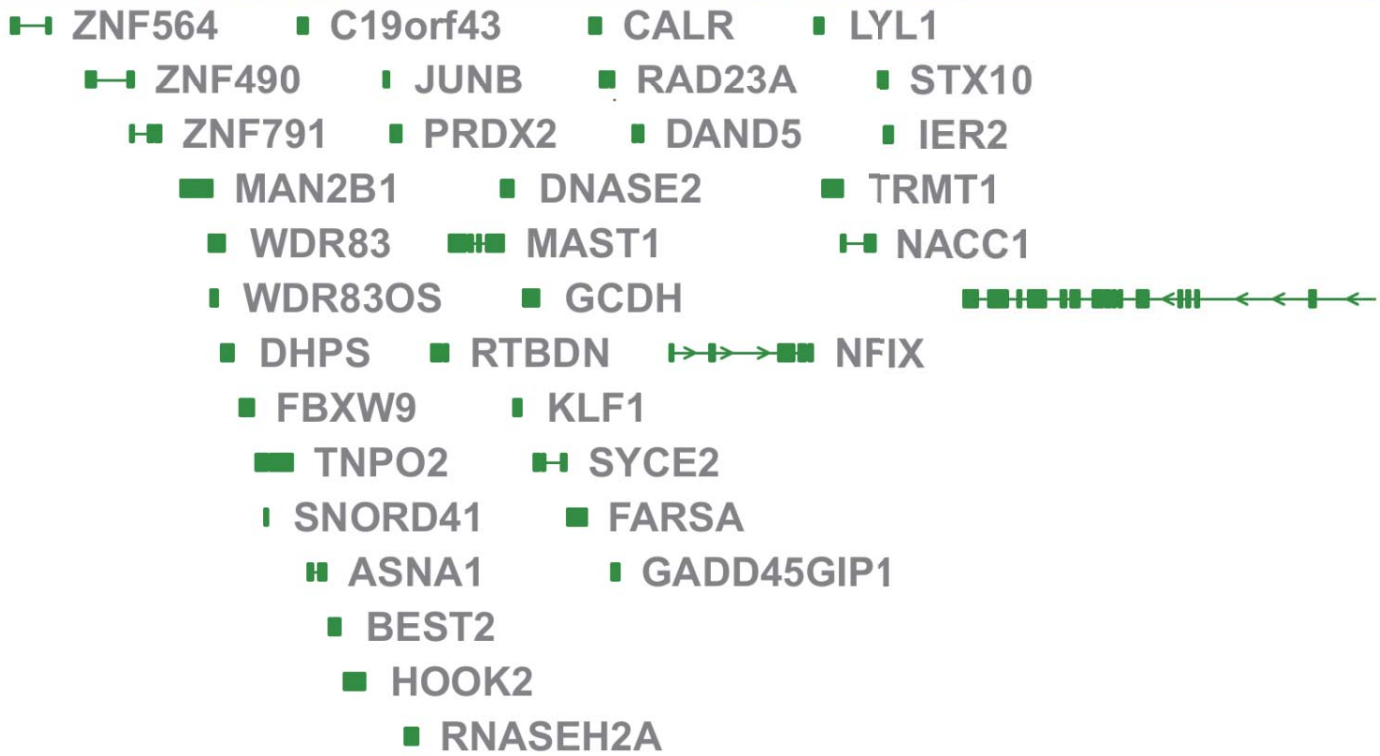
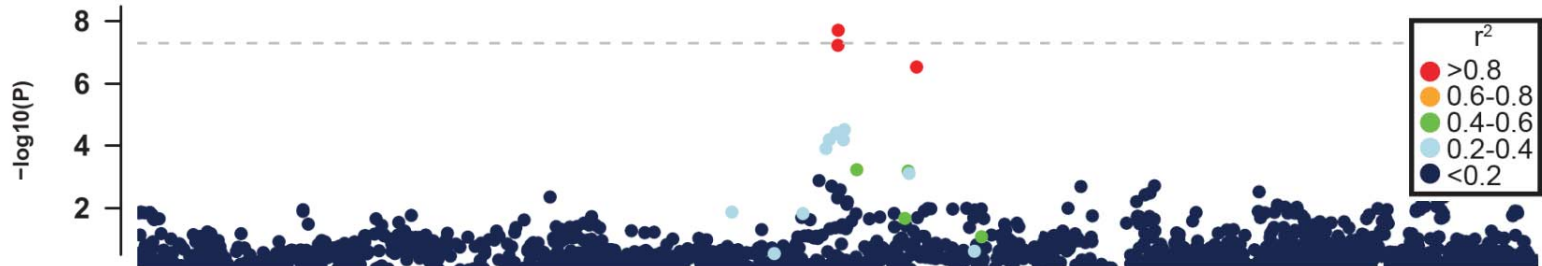
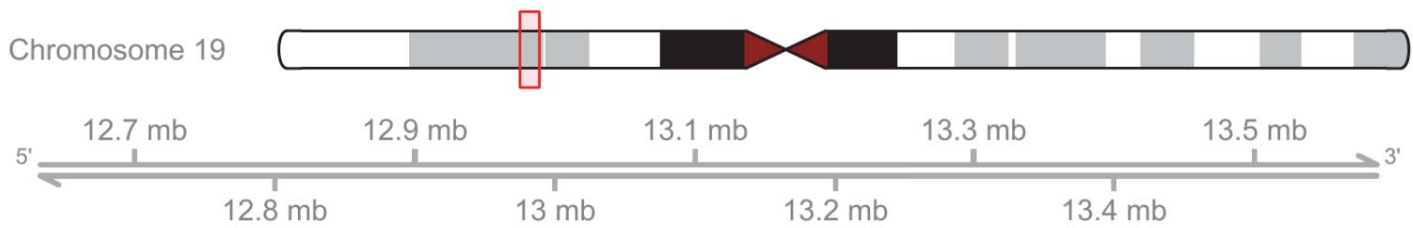
Uncorrected



Zscore of the effect allele: T



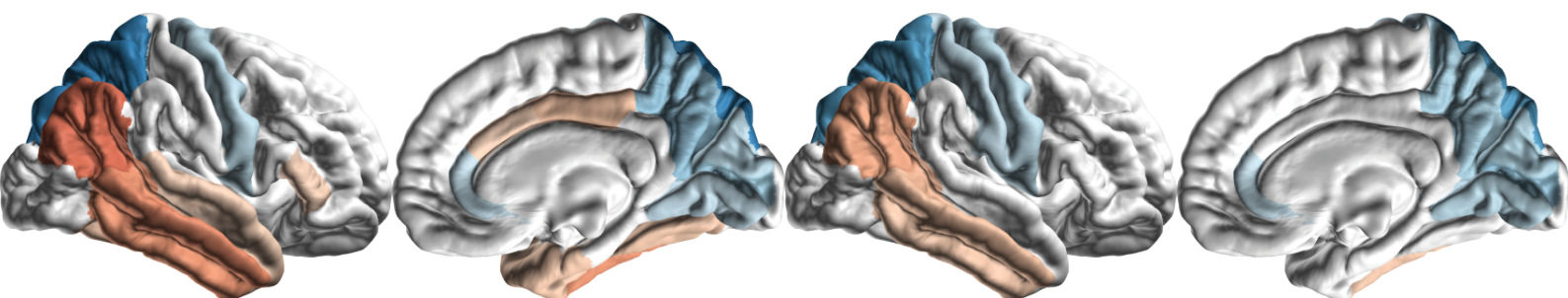
Banks of the Superior Temporal Sulcus SA: rs73006822



CP>GZZ>CP

Corrected for total SA

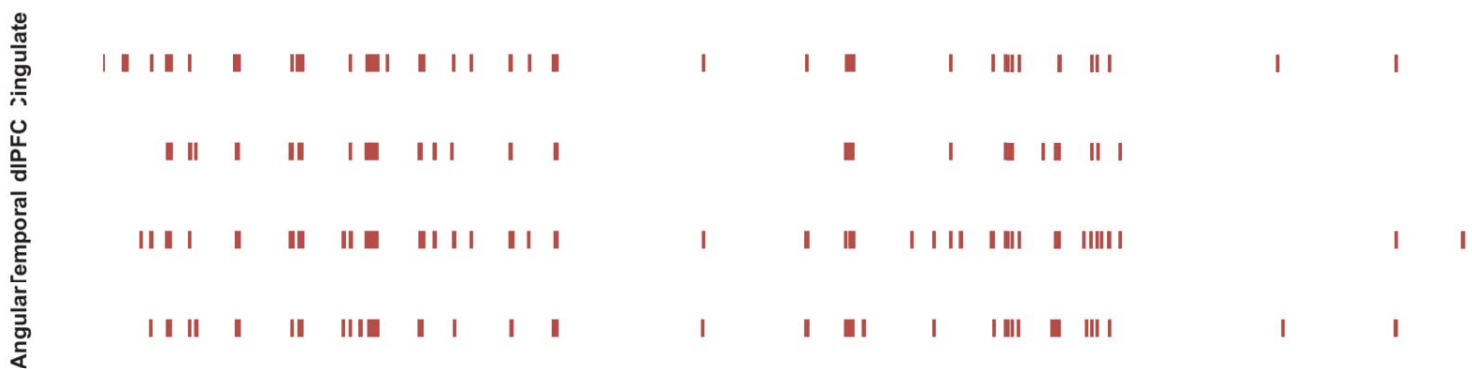
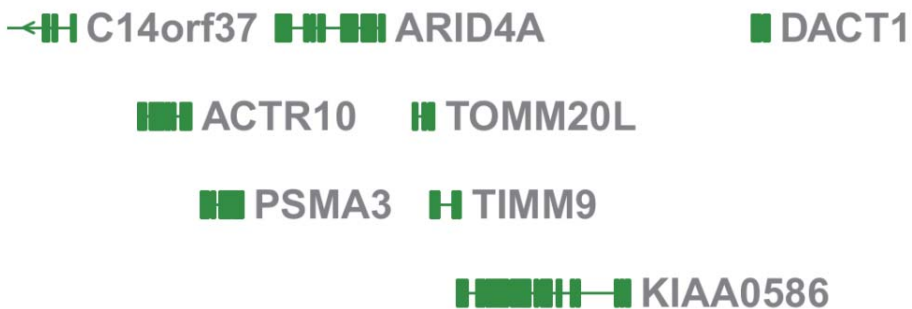
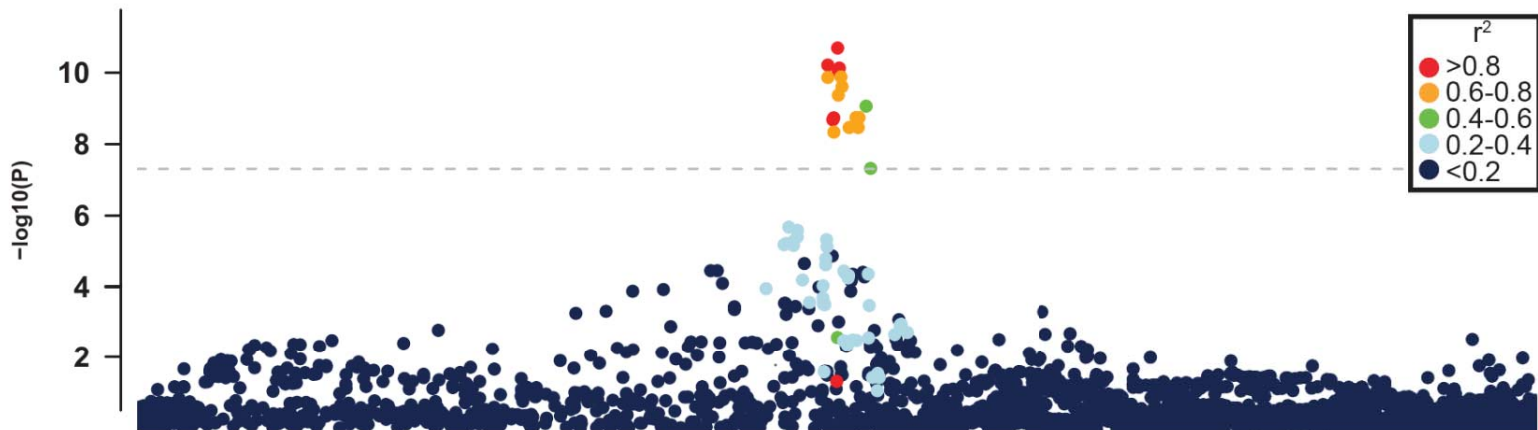
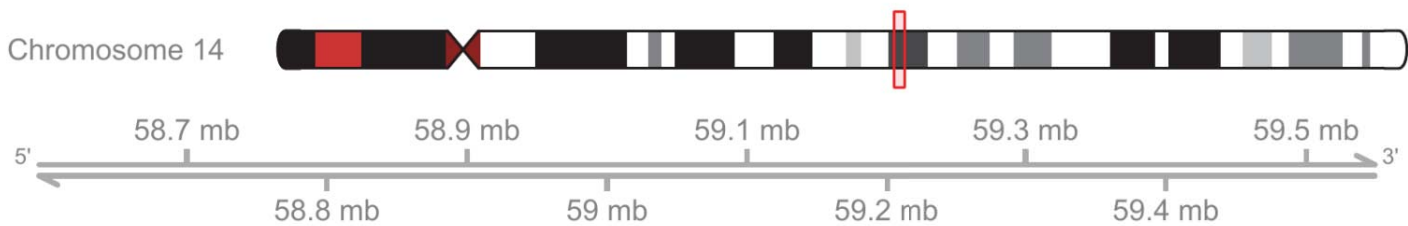
Uncorrected



Zscore of the effect allele: T

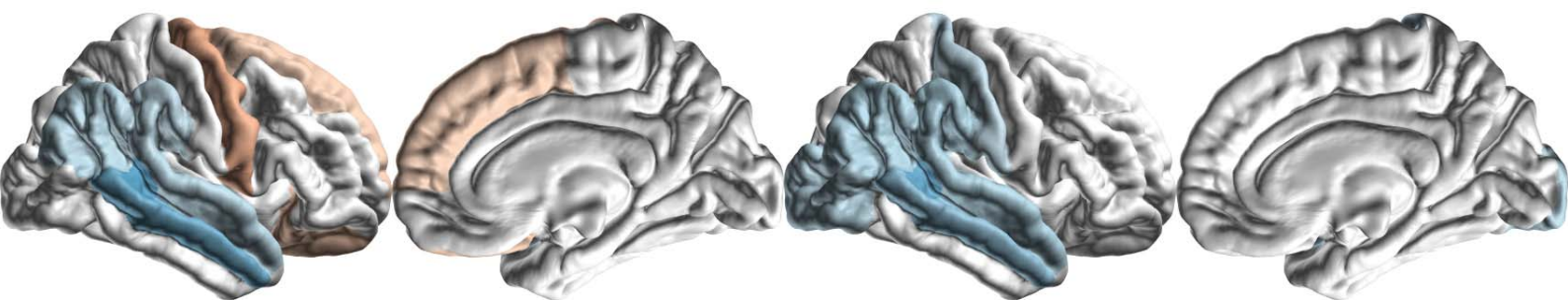


Banks of the Superior Temporal Sulcus TH: rs149142

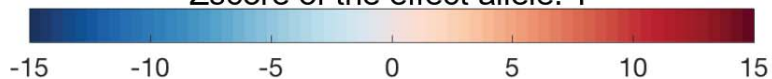


Corrected for average TH

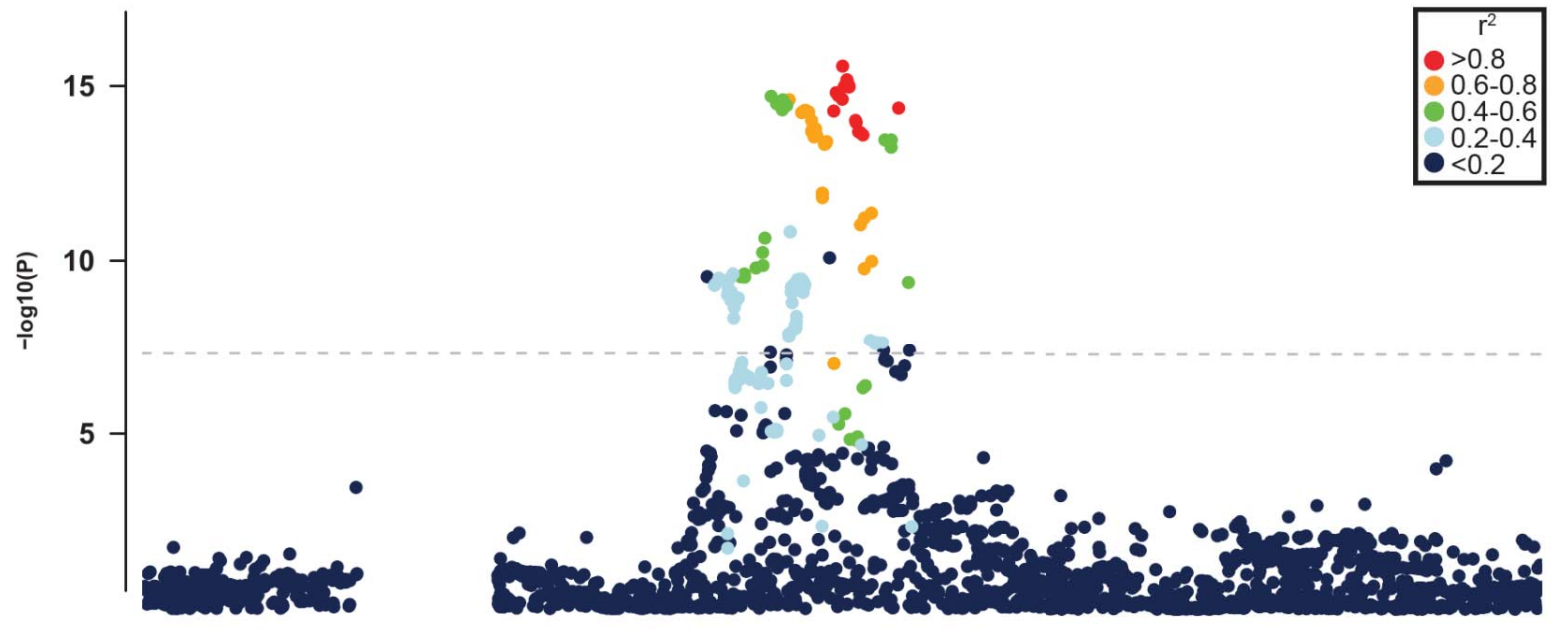
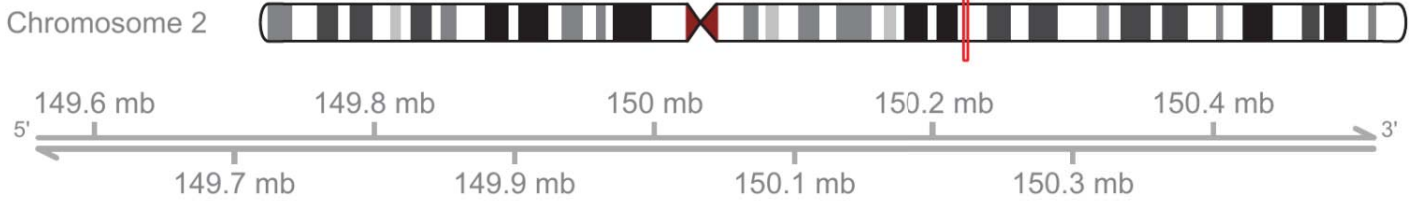
Uncorrected



Zscore of the effect allele: T

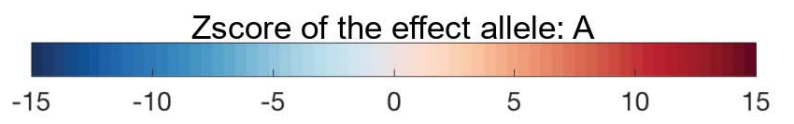
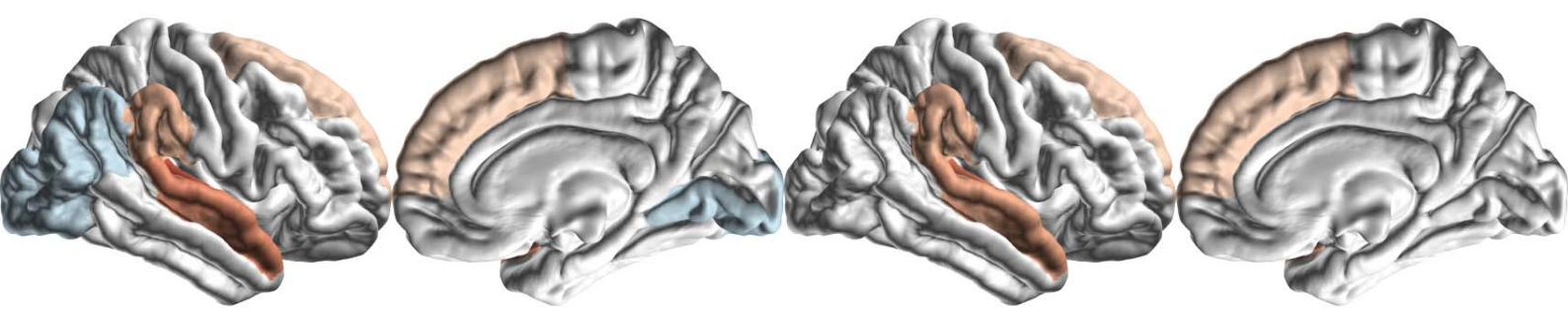


Transverse Temporal SA: rs11684511



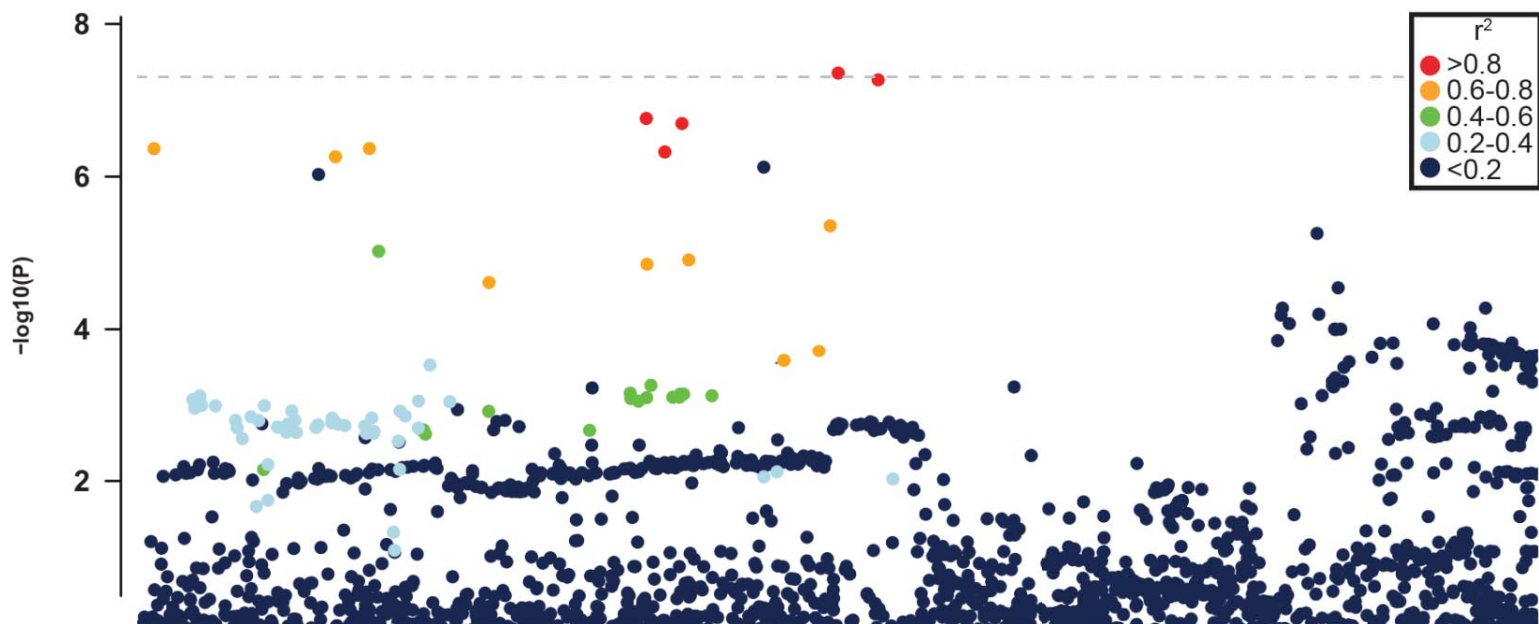
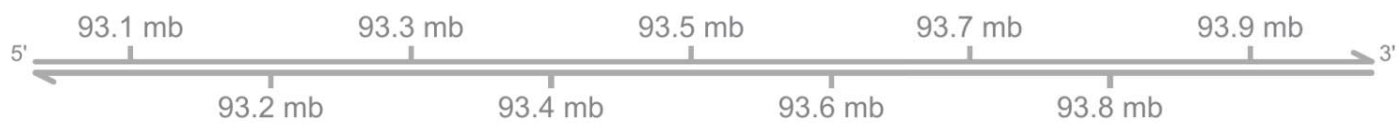
Corrected for total SA

Uncorrected



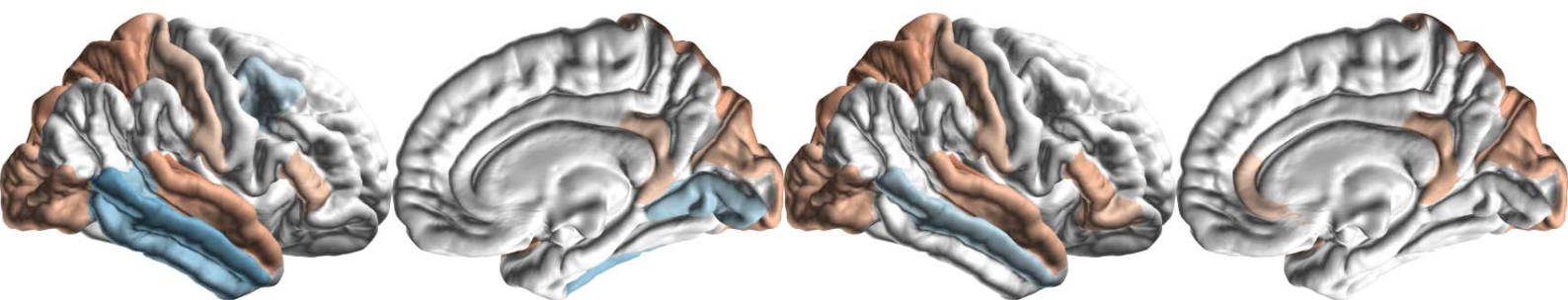
Transverse Temporal SA: rs115840240

Chromosome 5

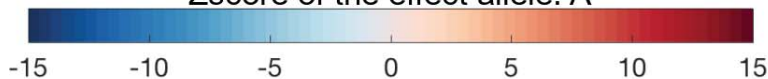


Corrected for total SA

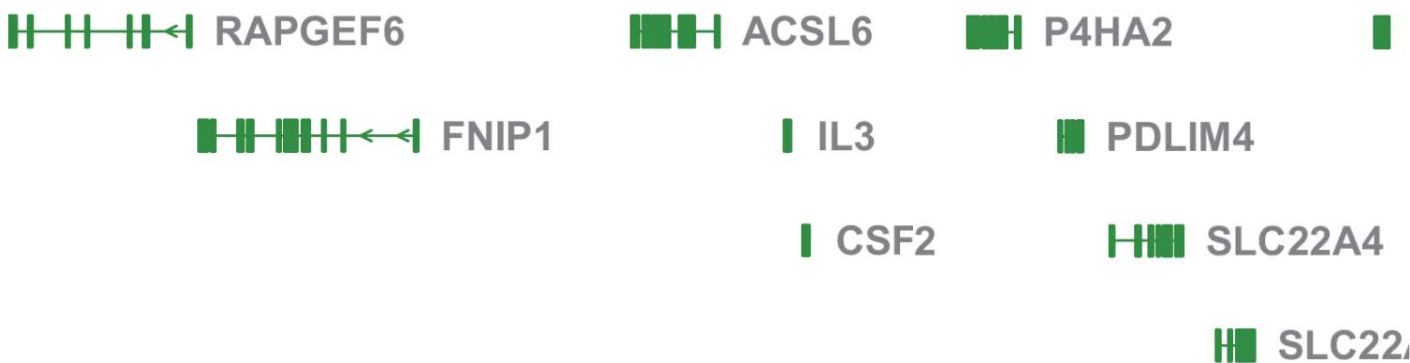
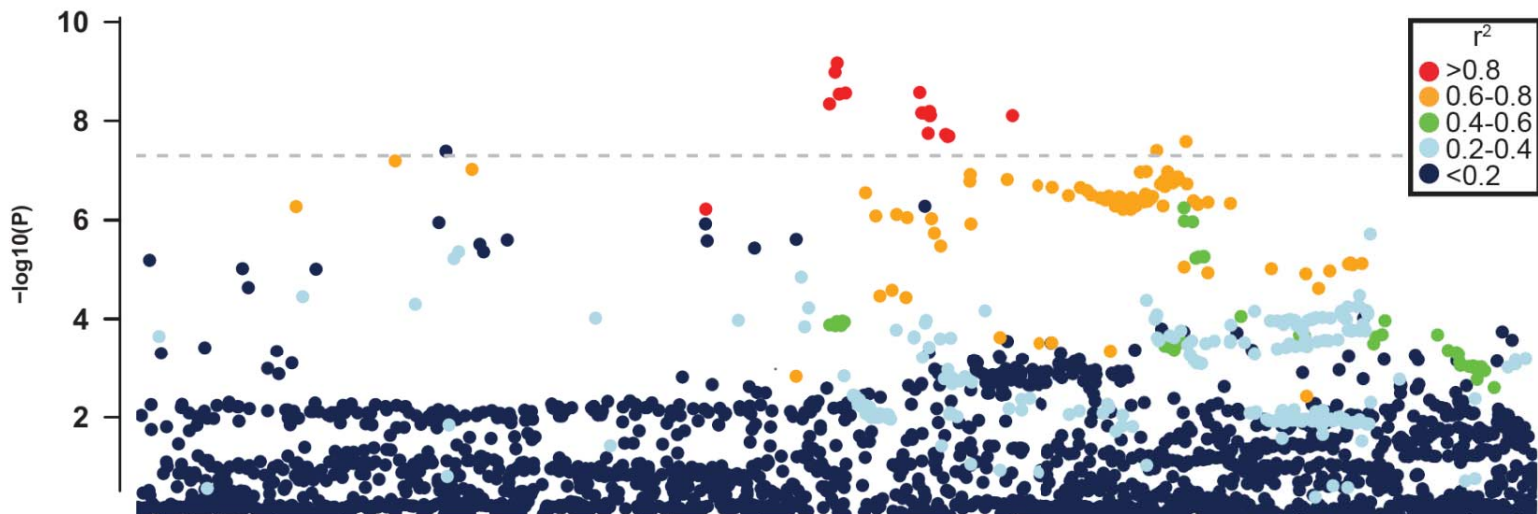
Uncorrected



Zscore of the effect allele: A

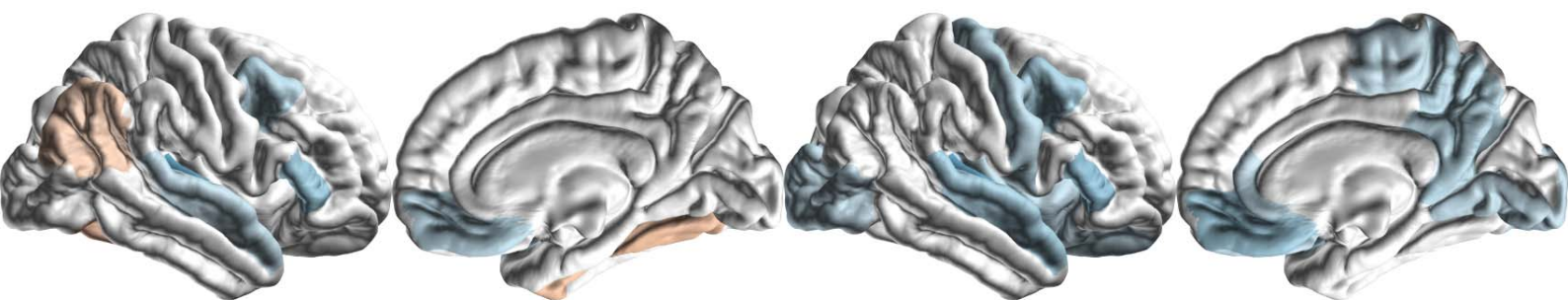


Transverse Temporal SA: rs7714191

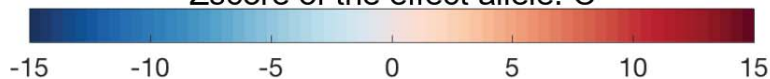


Corrected for total SA

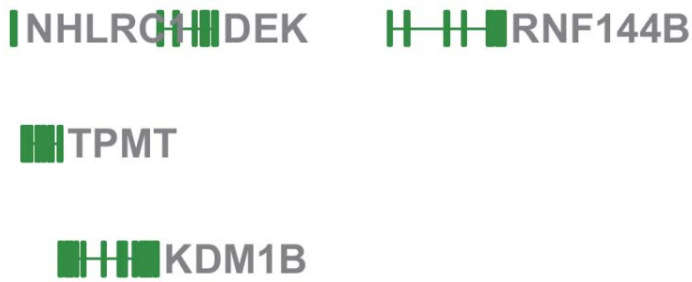
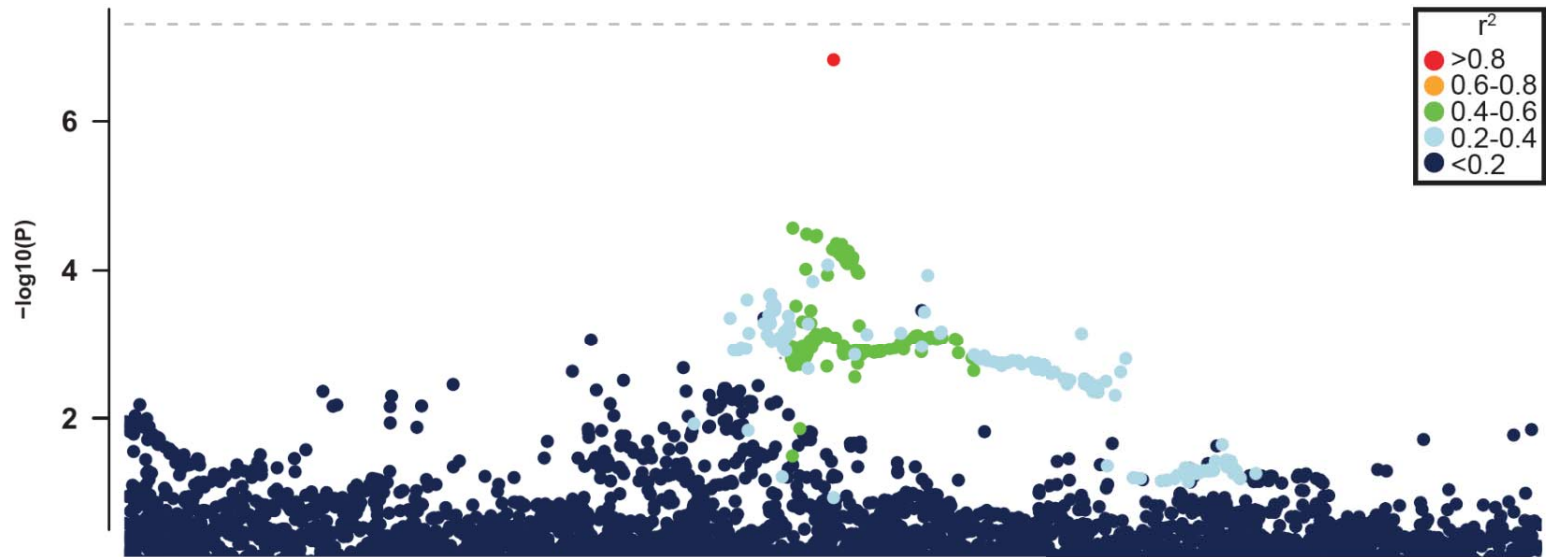
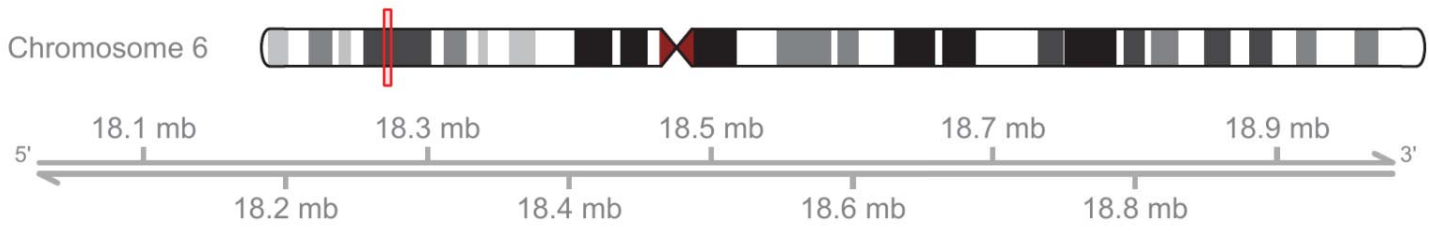
Uncorrected



Zscore of the effect allele: C

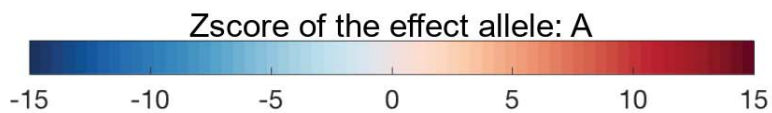
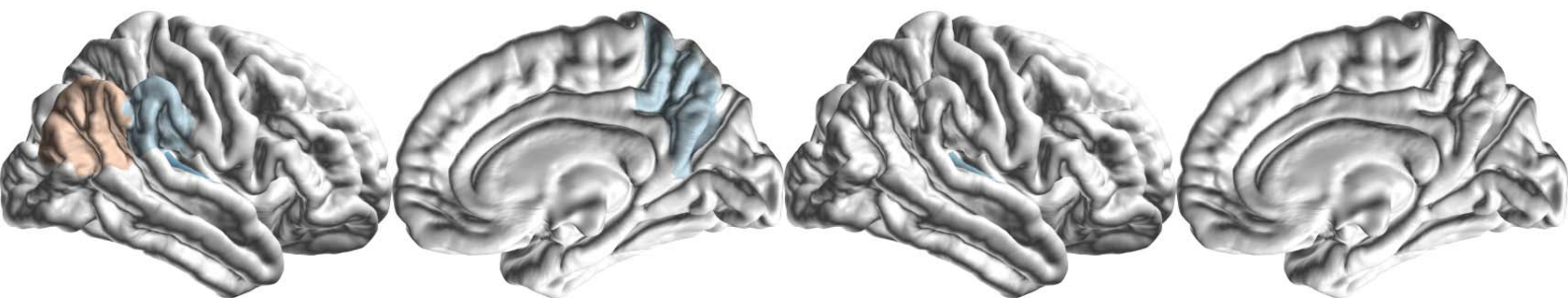


Transverse Temporal SA: rs72834200

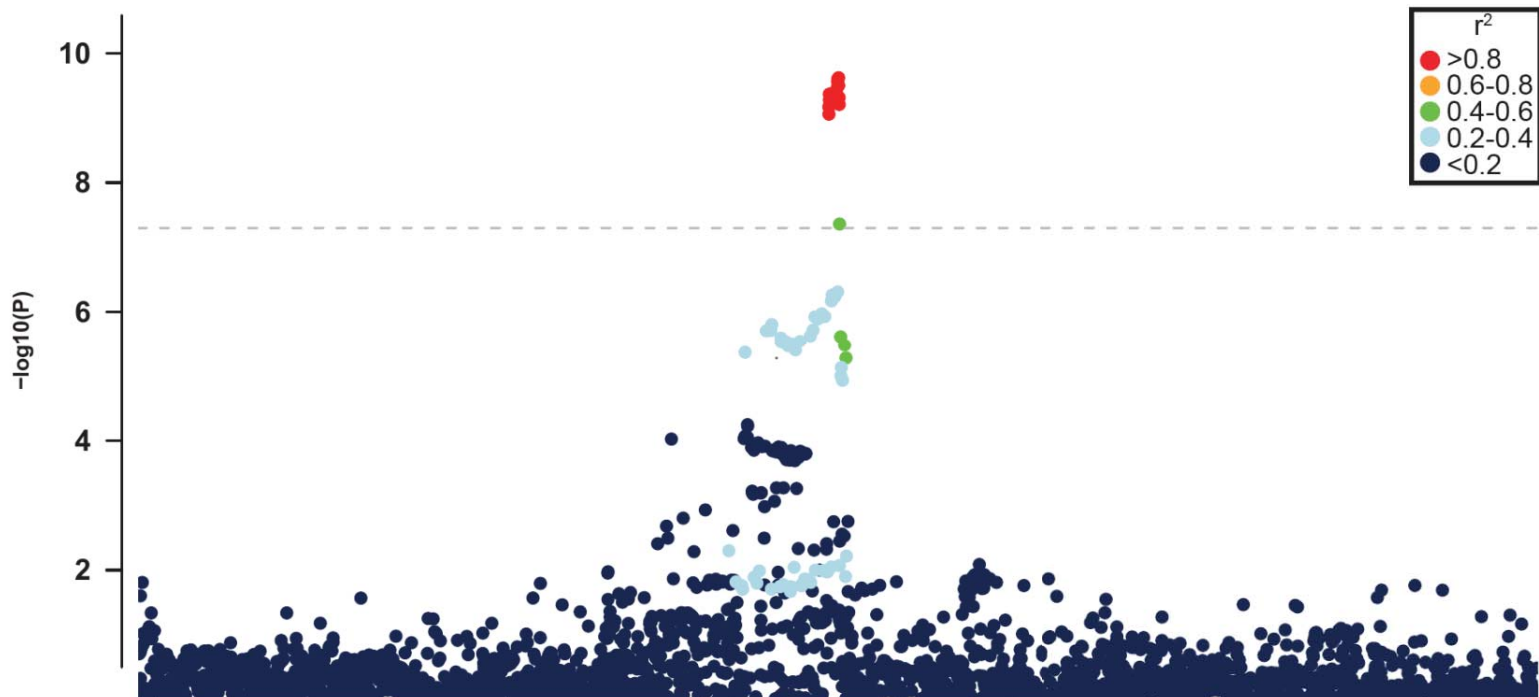
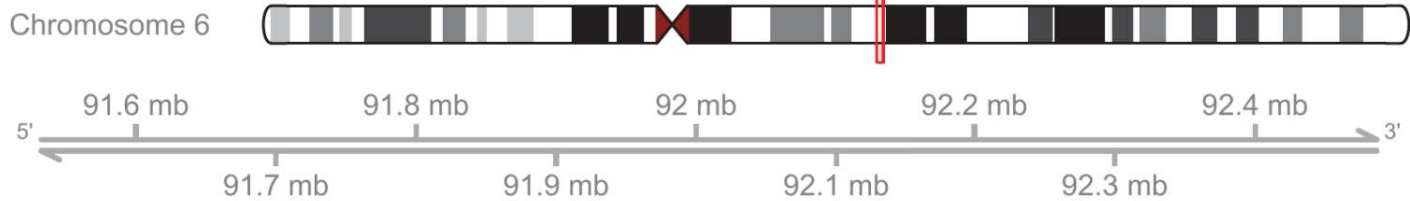


Corrected for total SA

Uncorrected



Transverse Temporal SA: rs4706391

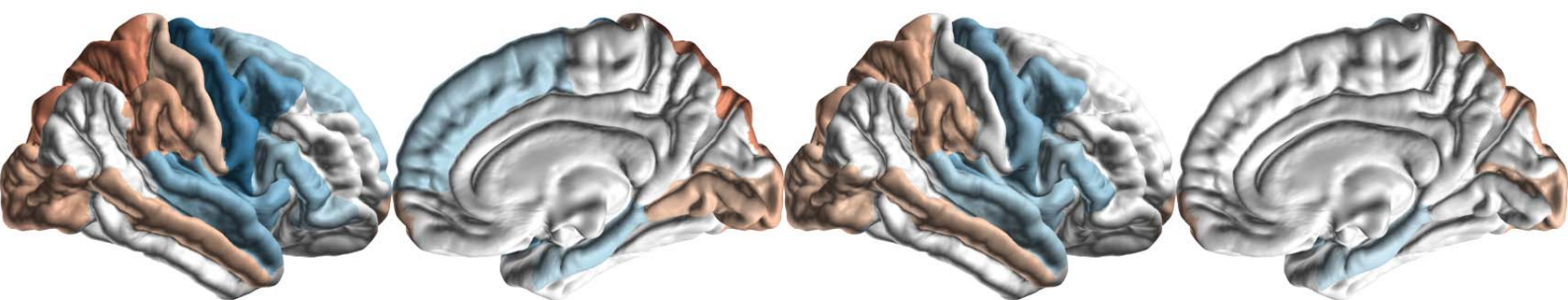


GZ>CP

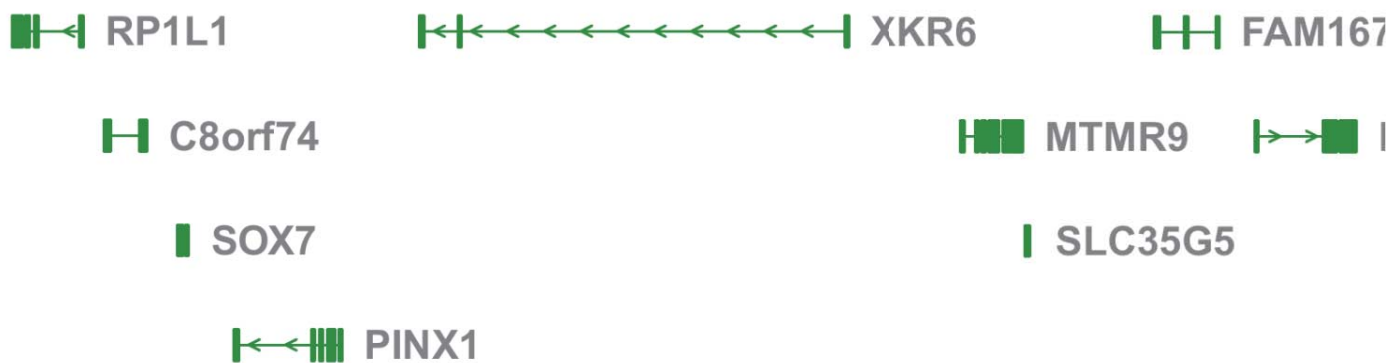
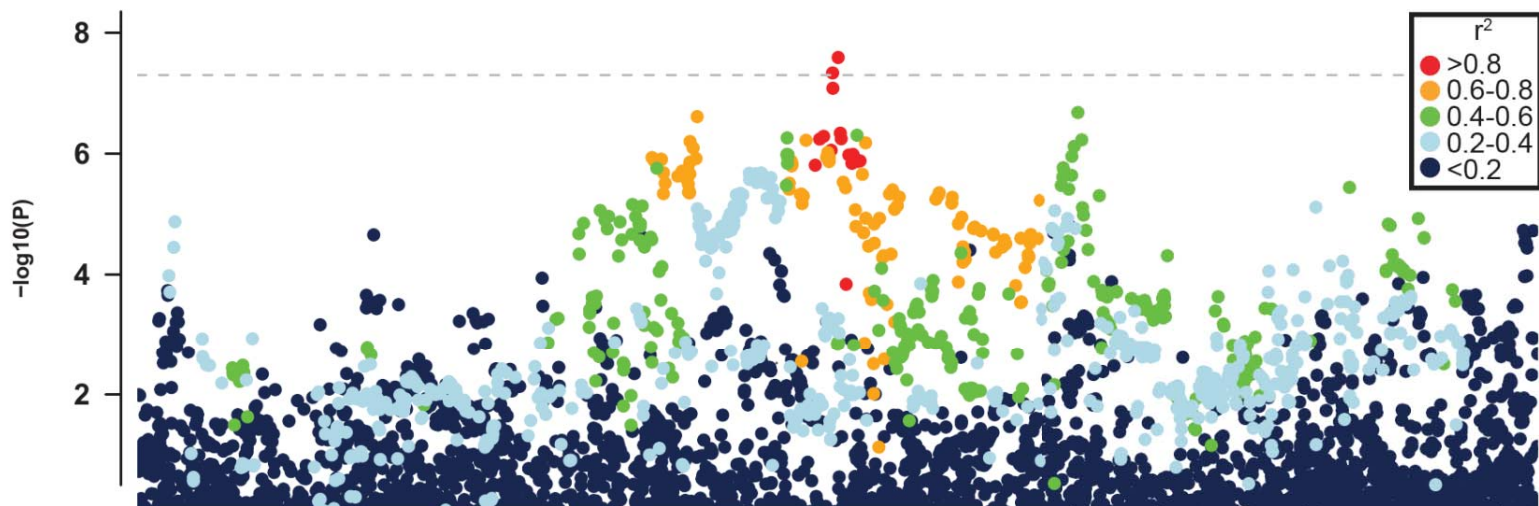
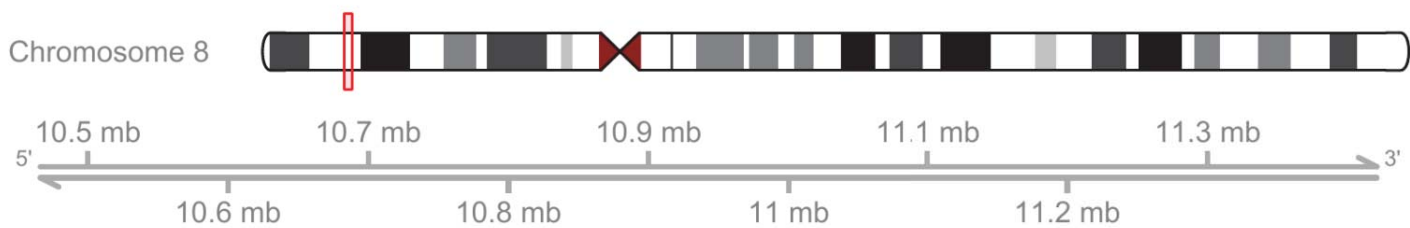
CP>GZ

Corrected for total SA

Uncorrected

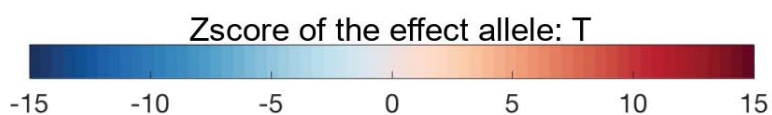
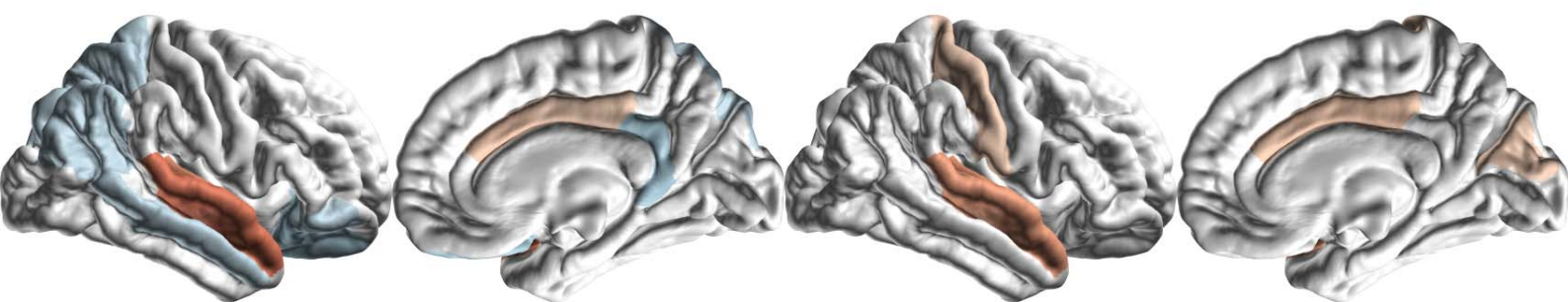


Transverse Temporal SA: rs2409691

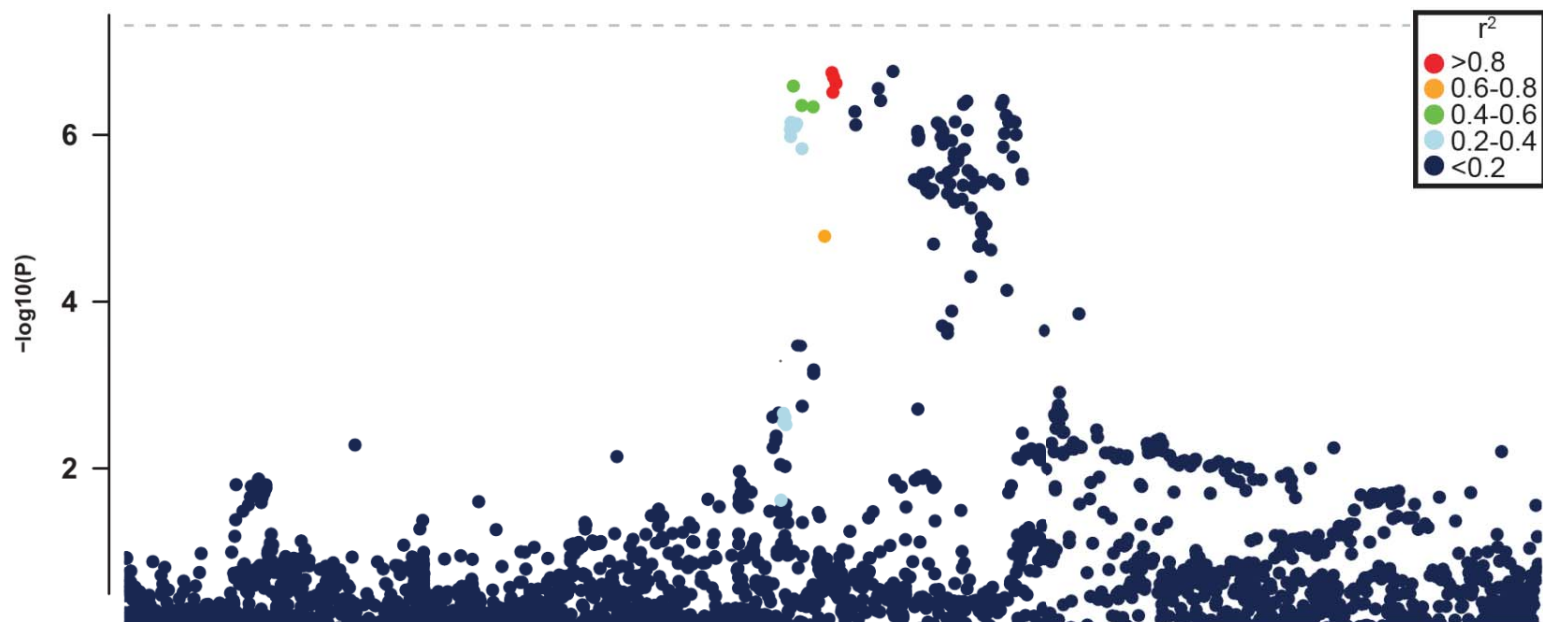


Corrected for total SA

Uncorrected



Transverse Temporal SA: rs4923822



C15orf54

THBS1

FSIP1

GZ>CP

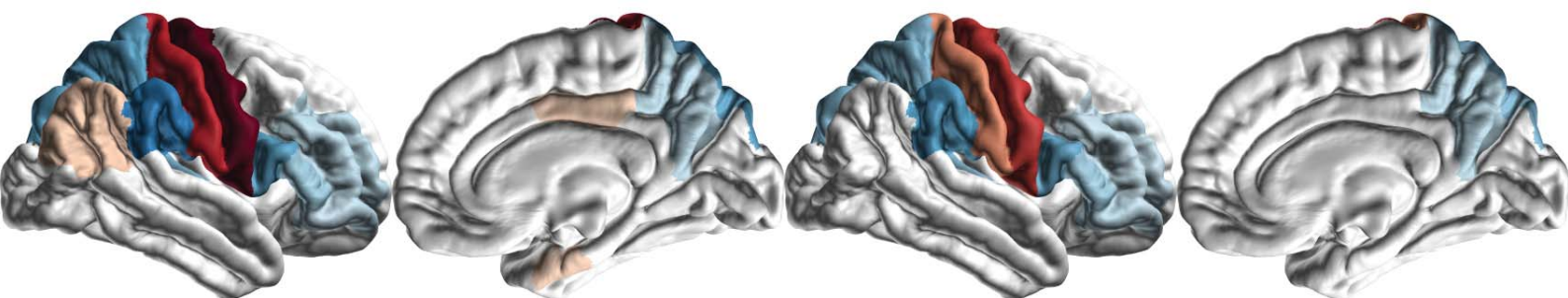


CP>GZ



Corrected for total SA

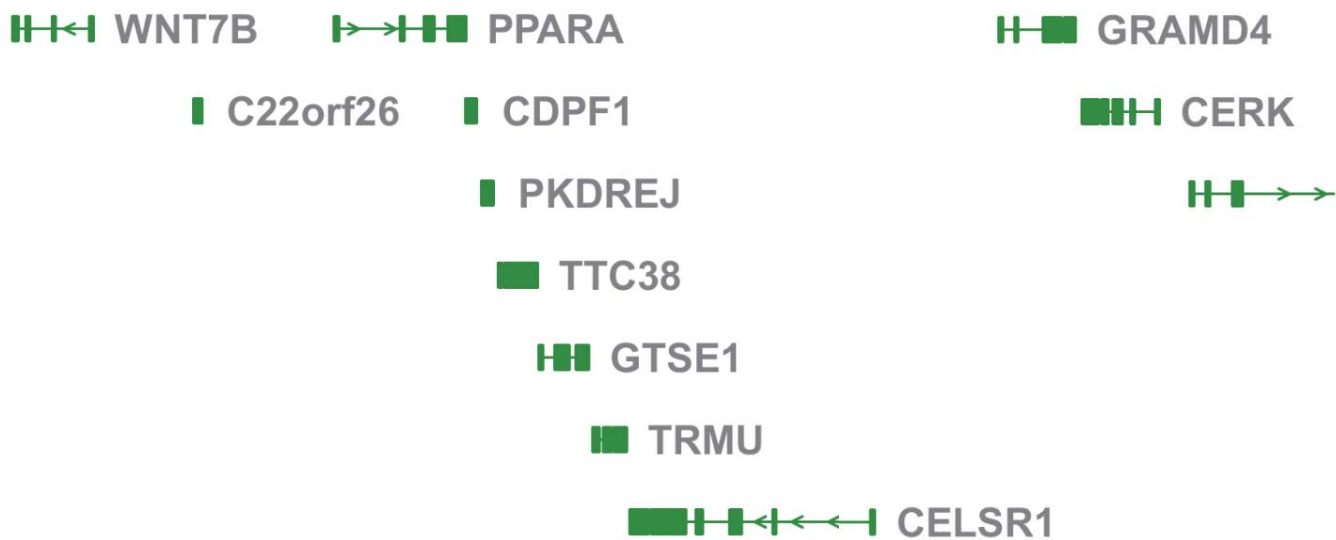
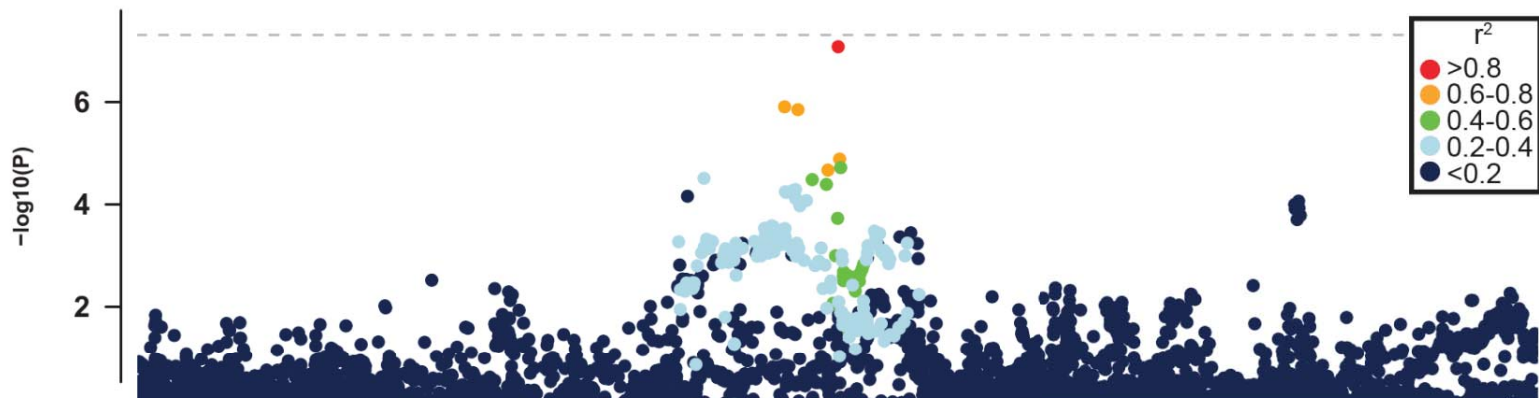
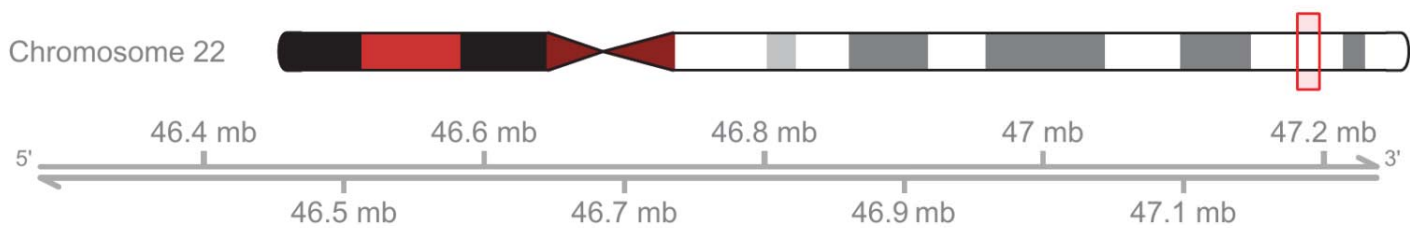
Uncorrected



Zscore of the effect allele: A



Transverse Temporal SA: rs9615351

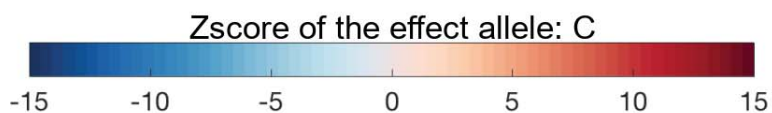
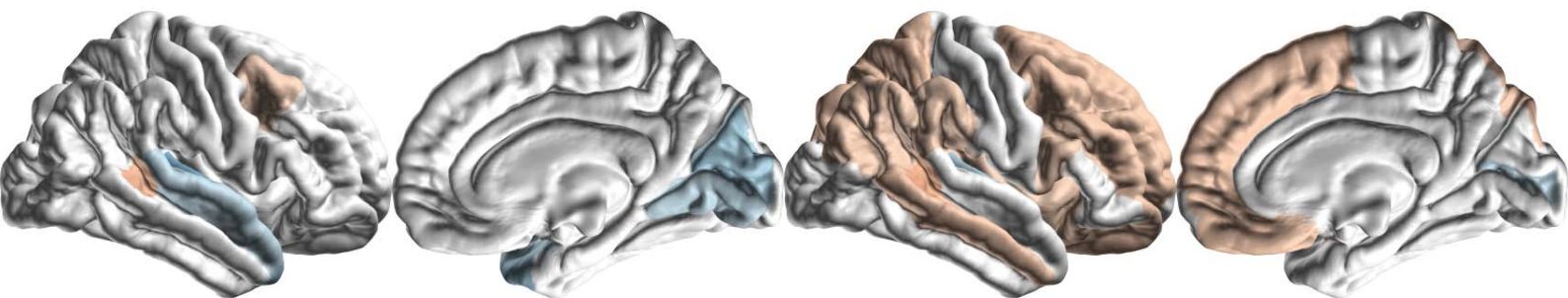


CP>GZ GZ>CP

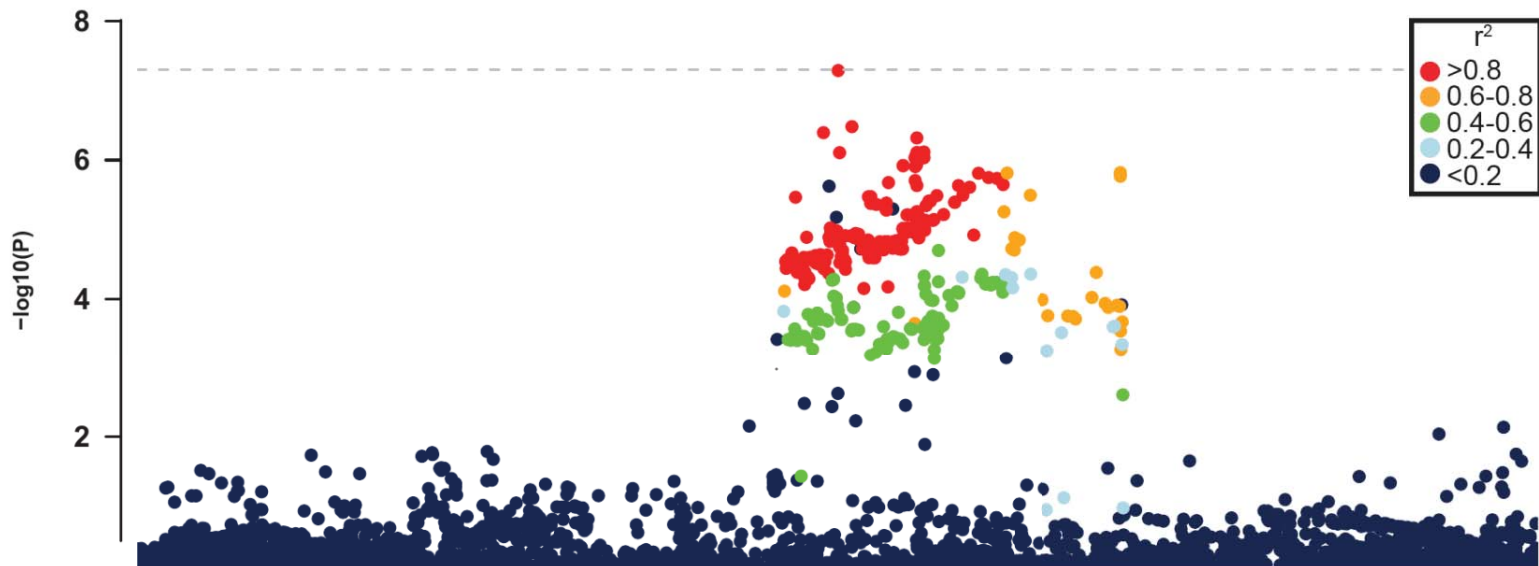
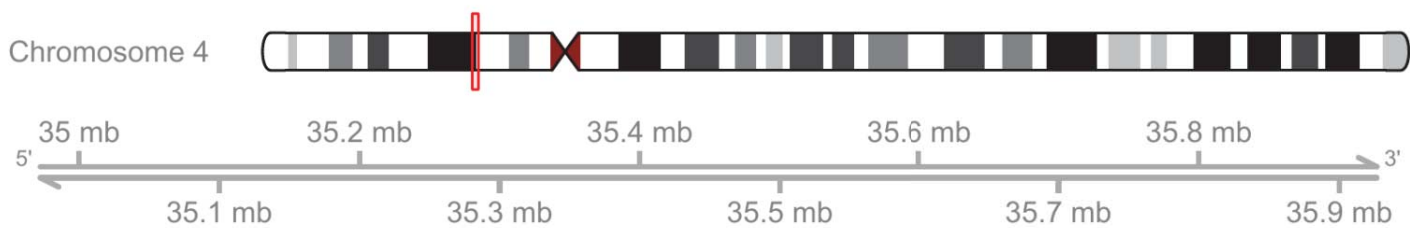


Corrected for total SA

Uncorrected

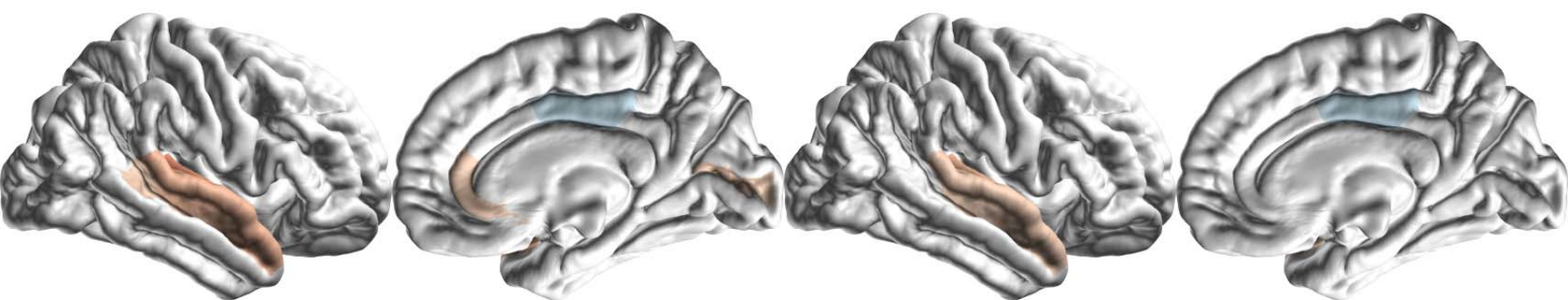


Transverse Temporal TH: rs57125540



Corrected for average TH

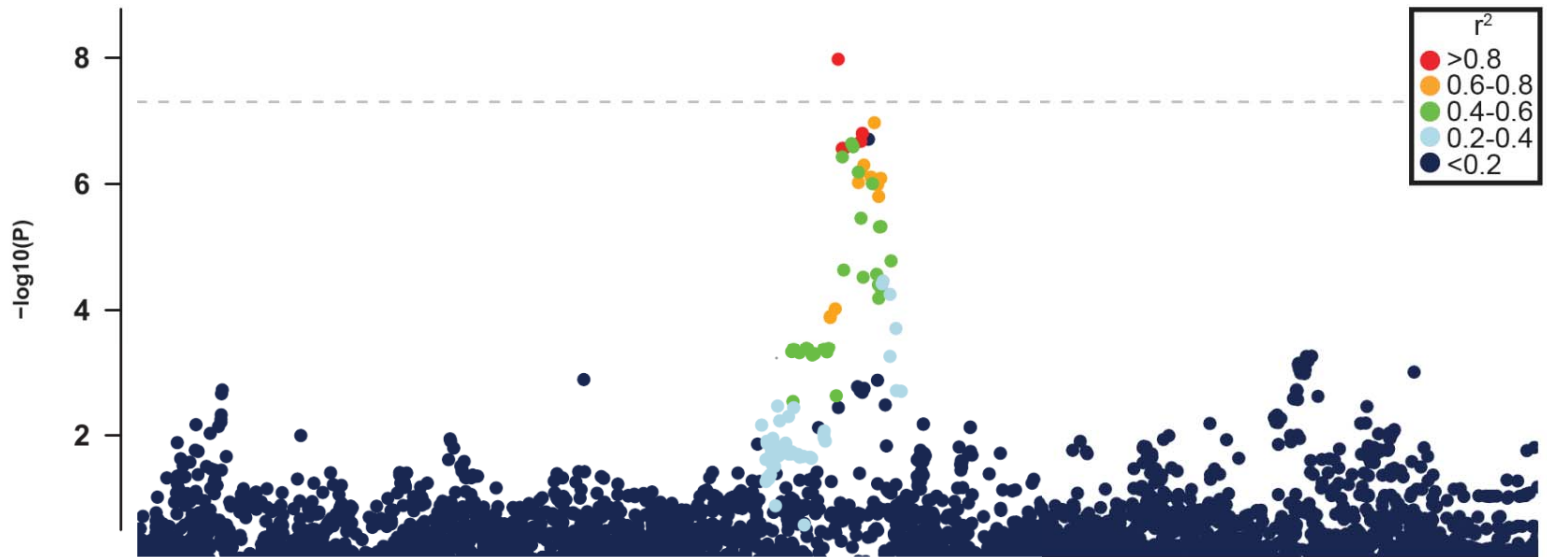
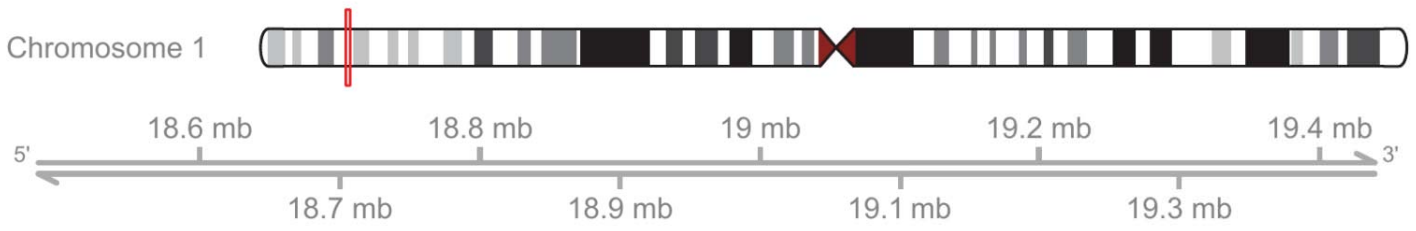
Uncorrected



Zscore of the effect allele: A

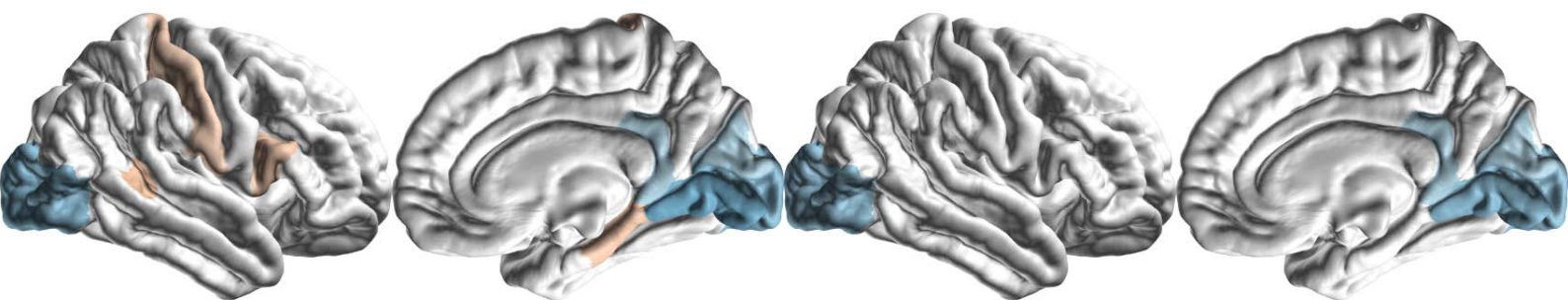


Lingual SA: rs1934057



Corrected for total SA

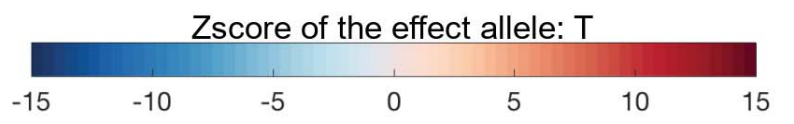
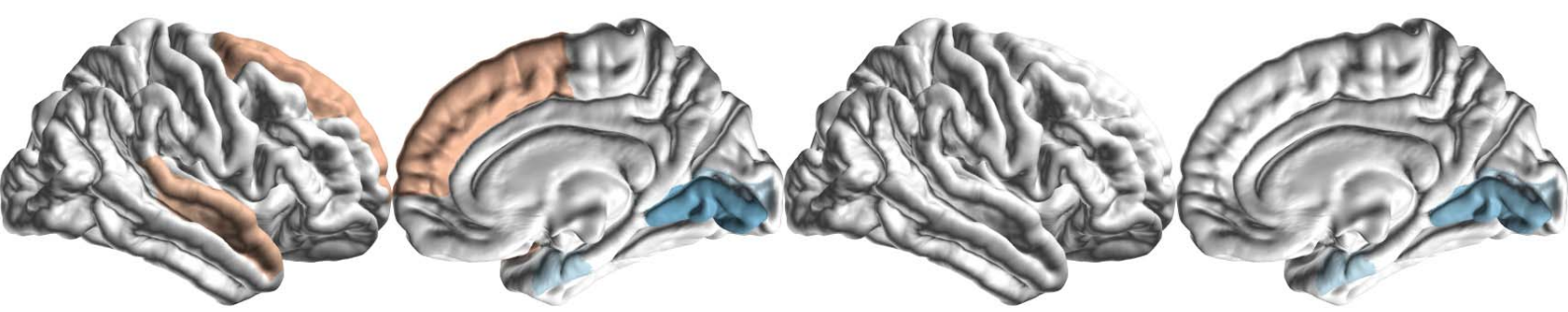
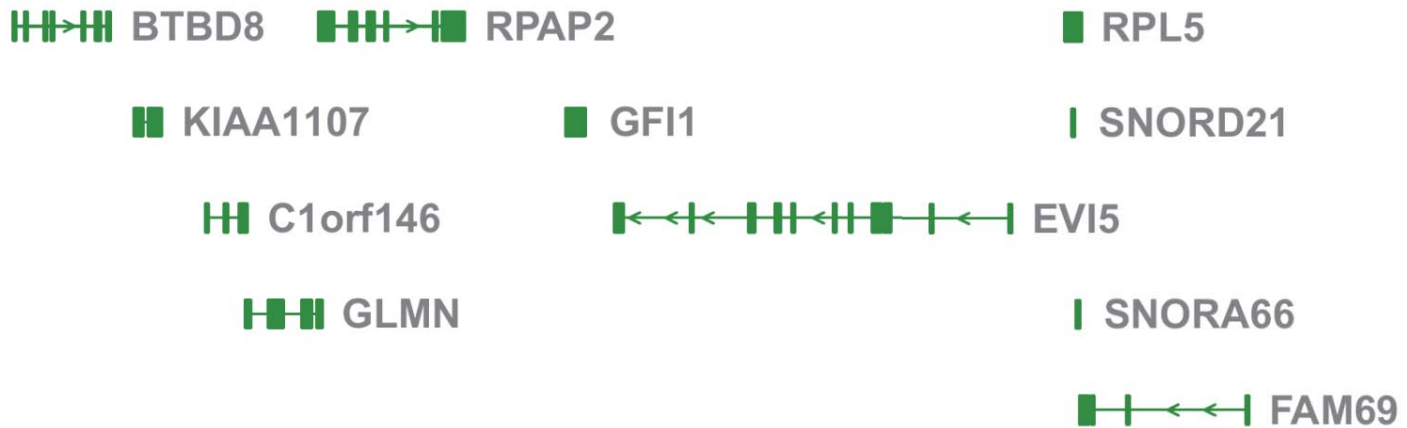
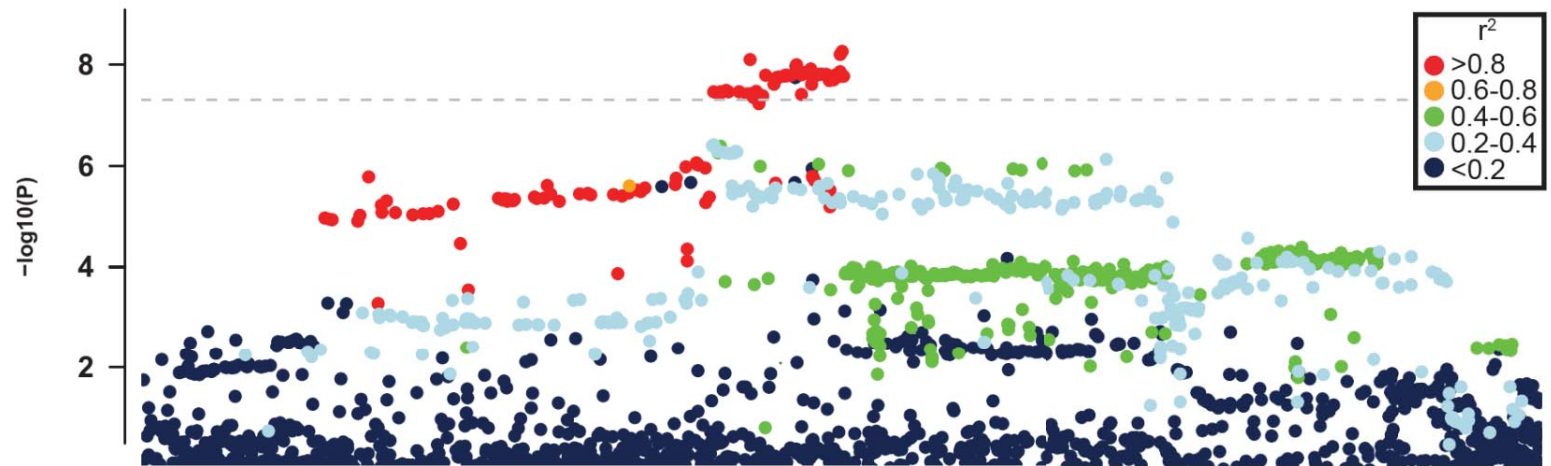
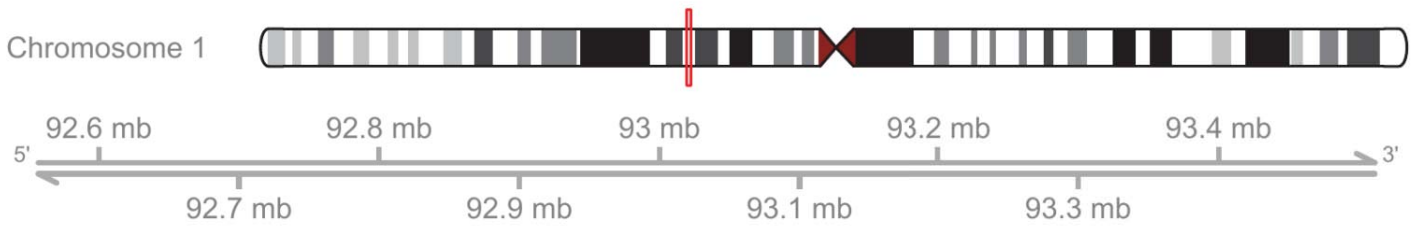
Uncorrected



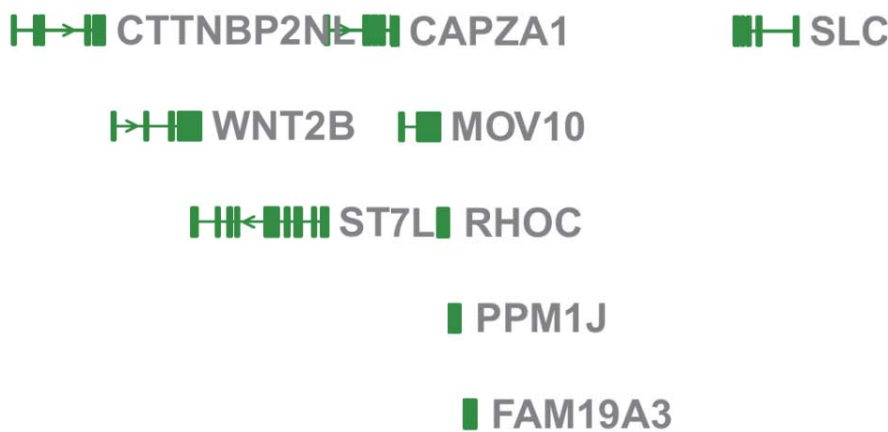
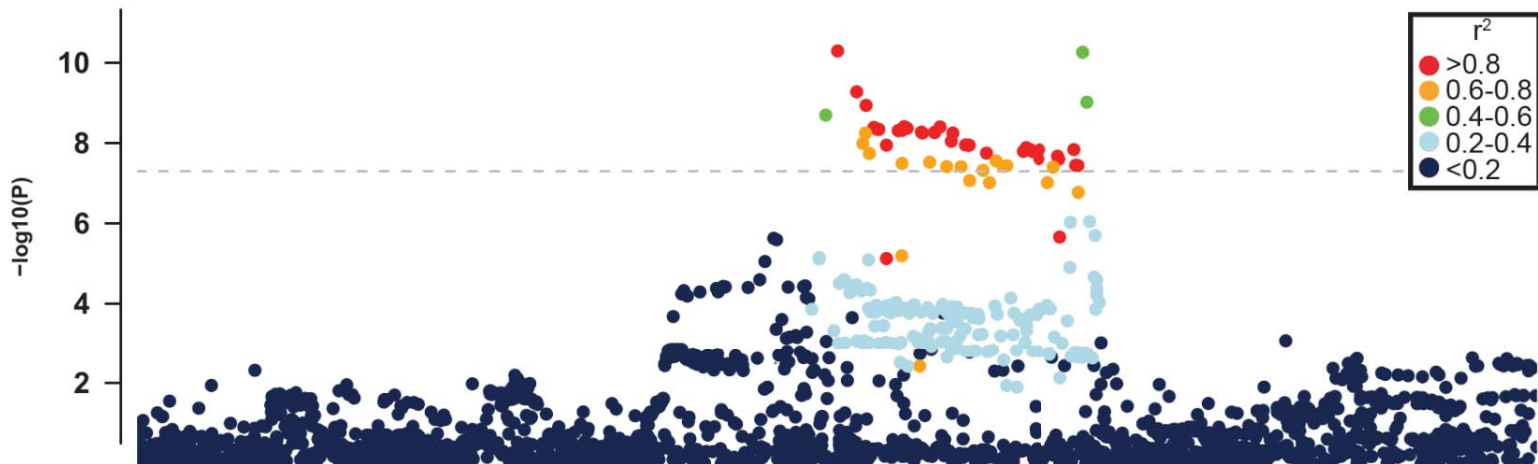
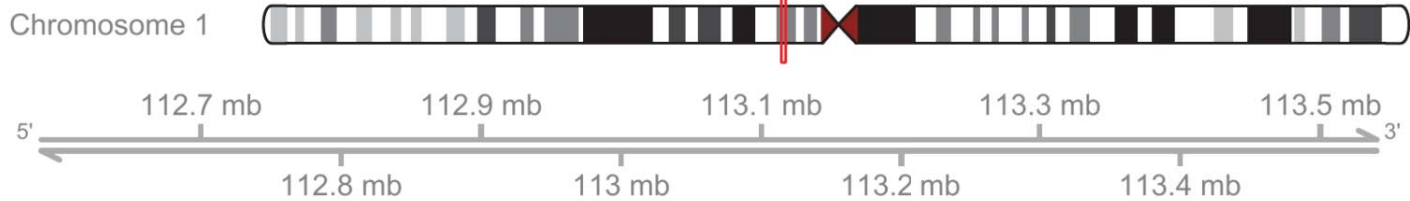
Zscore of the effect allele: T



Lingual SA: rs1556562



Lingual SA: rs910697

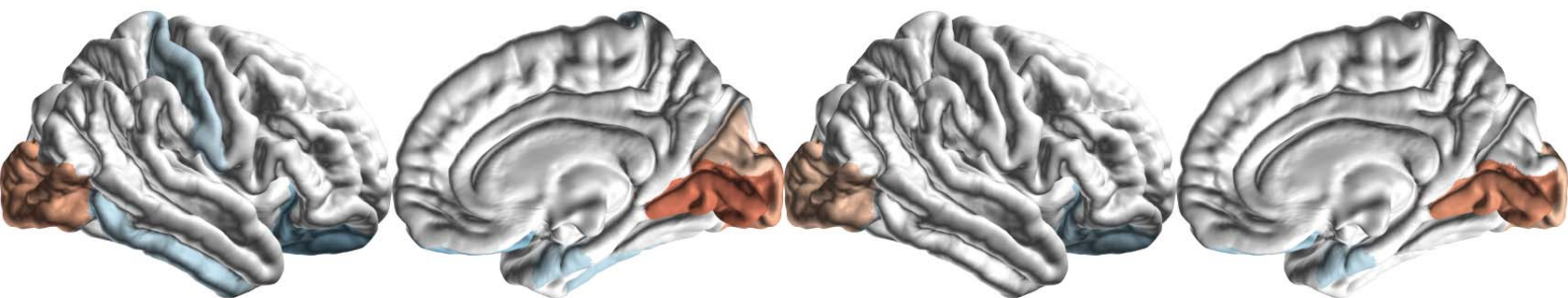


CP>GZ GZ>CP



Corrected for total SA

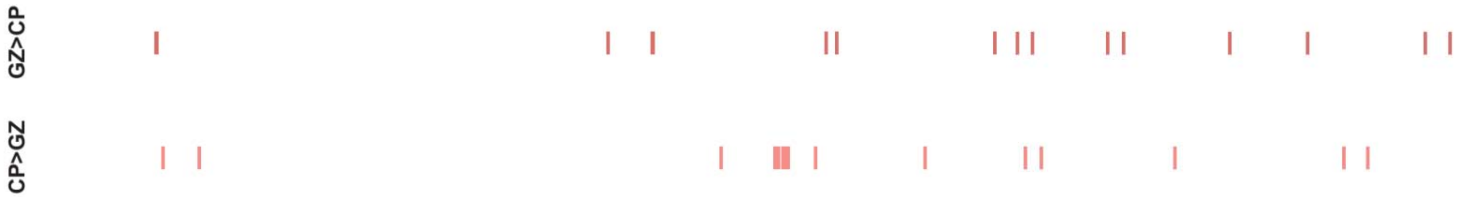
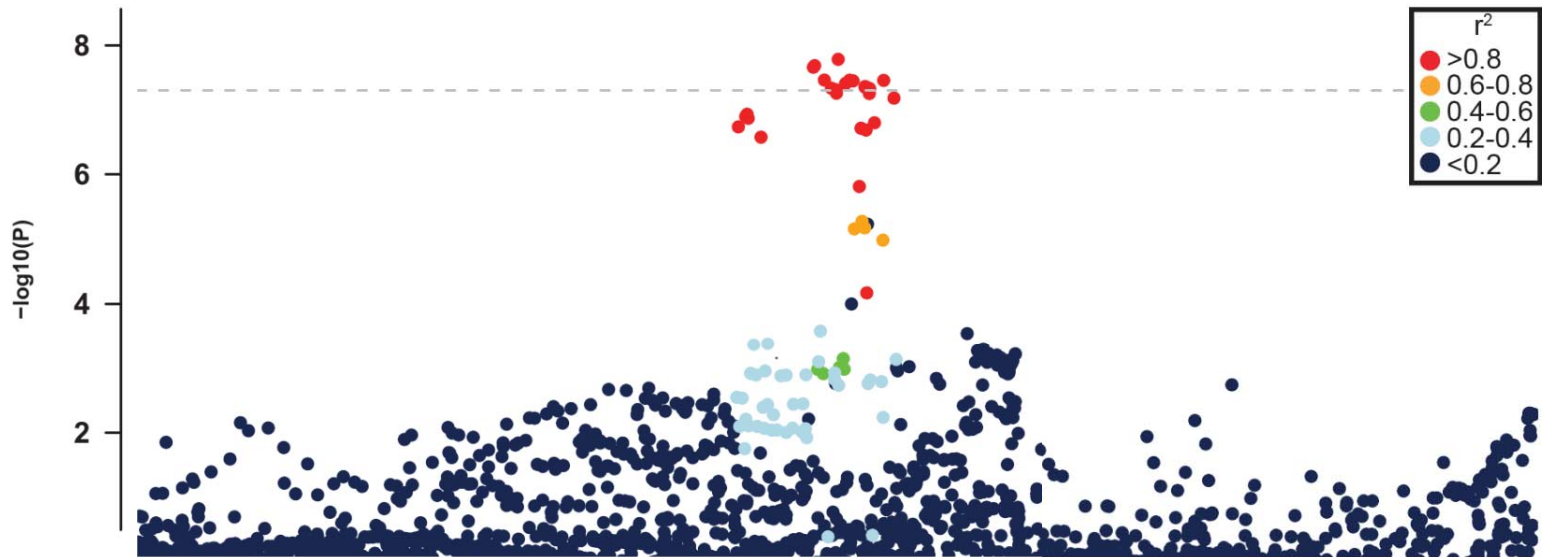
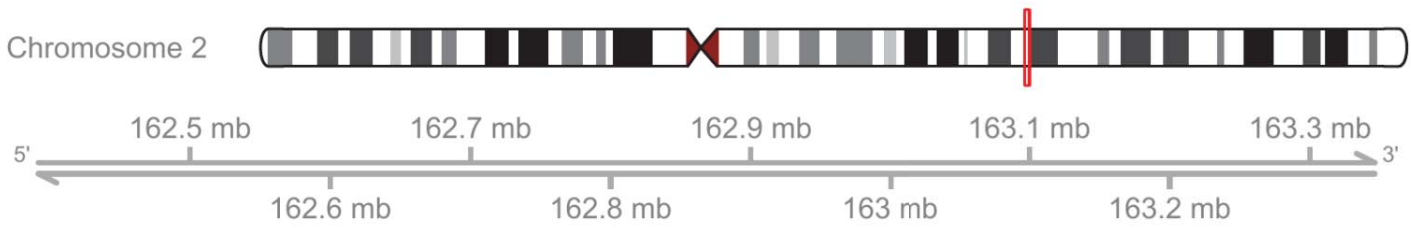
Uncorrected



Zscore of the effect allele: A

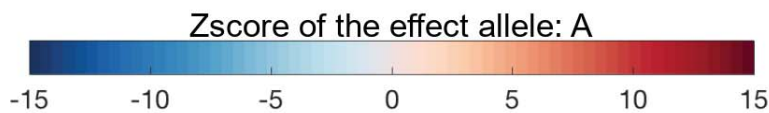
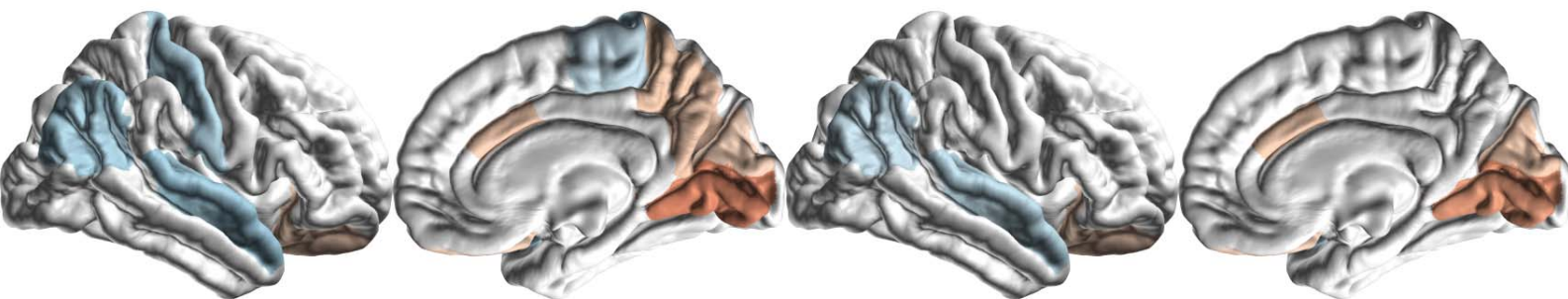


Lingual SA: rs1014444



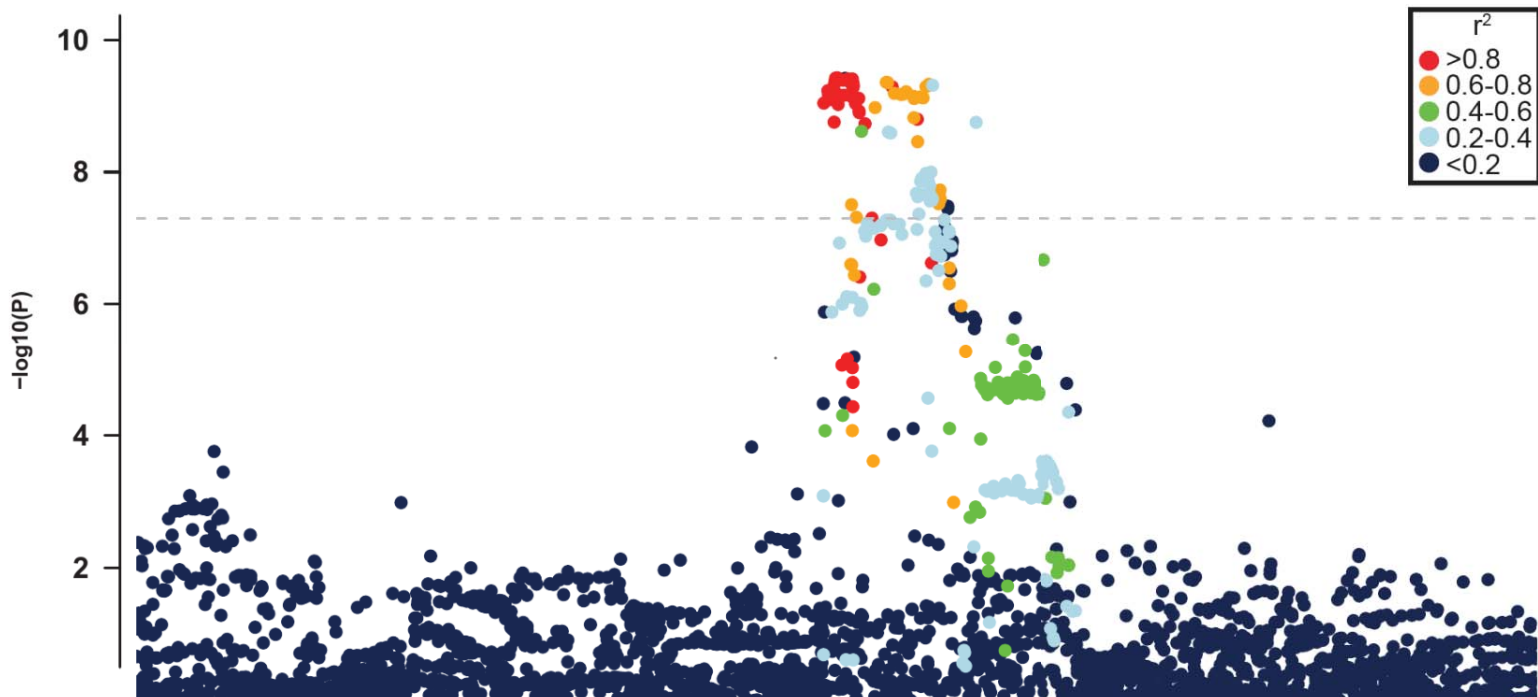
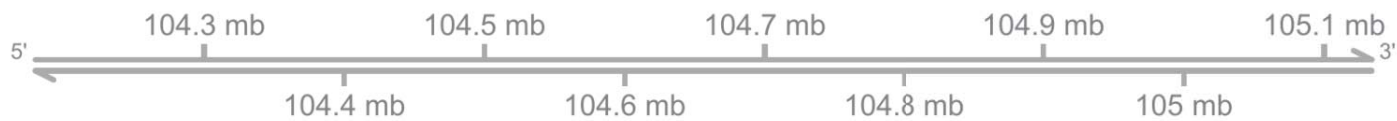
Corrected for total SA

Uncorrected



Lingual SA: rs2118004

Chromosome 3



GZ>CP

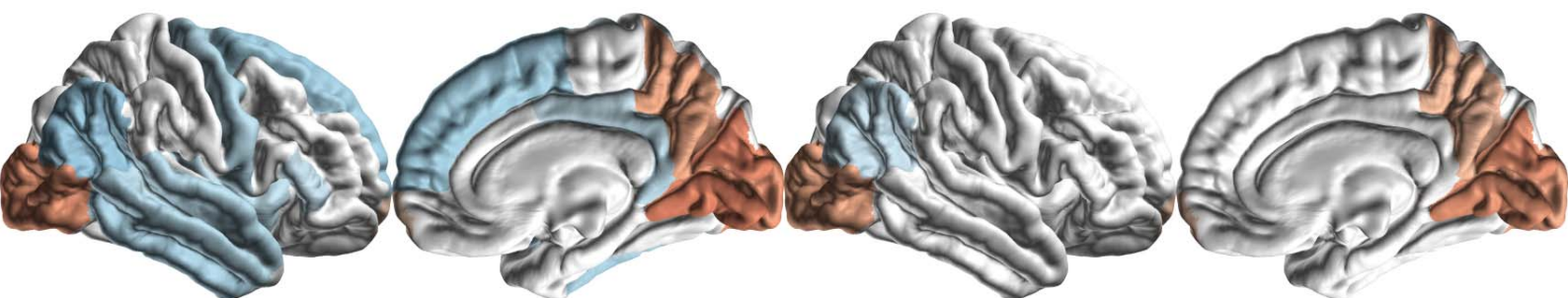


CP>GZ

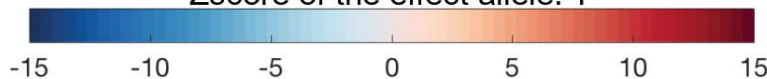


Corrected for total SA

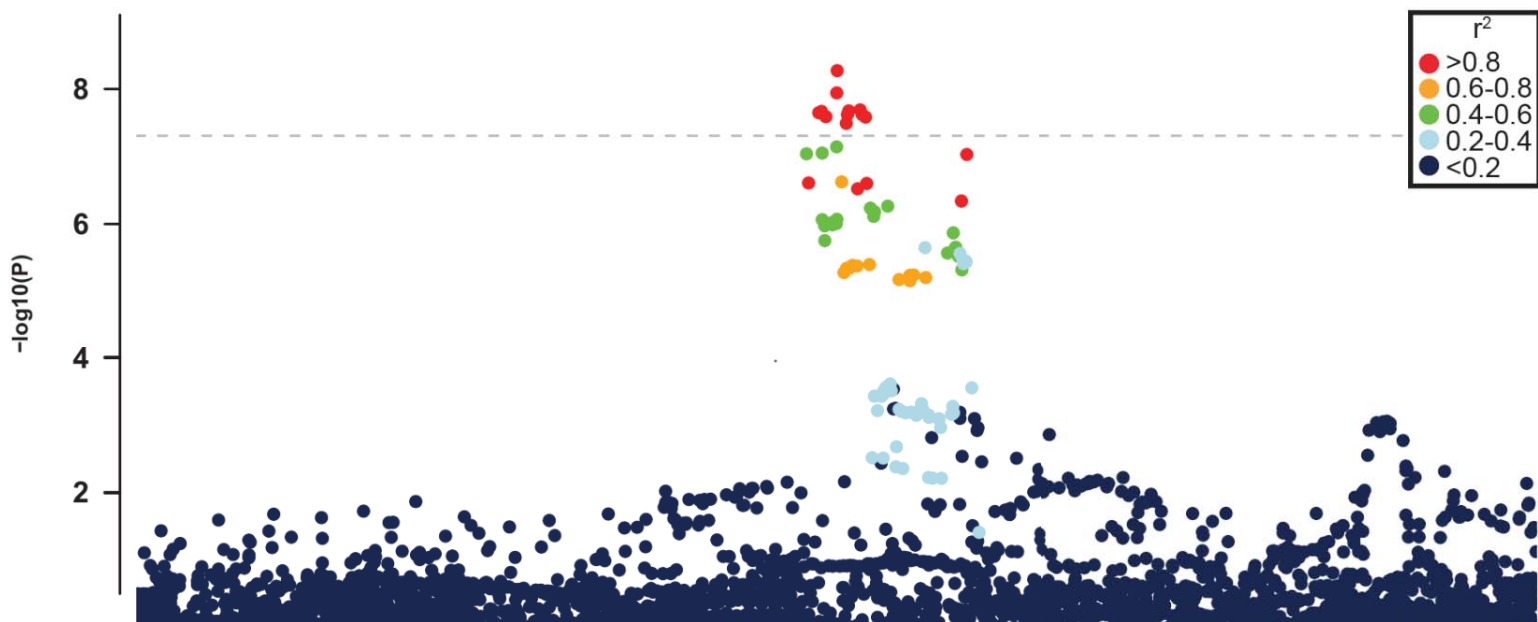
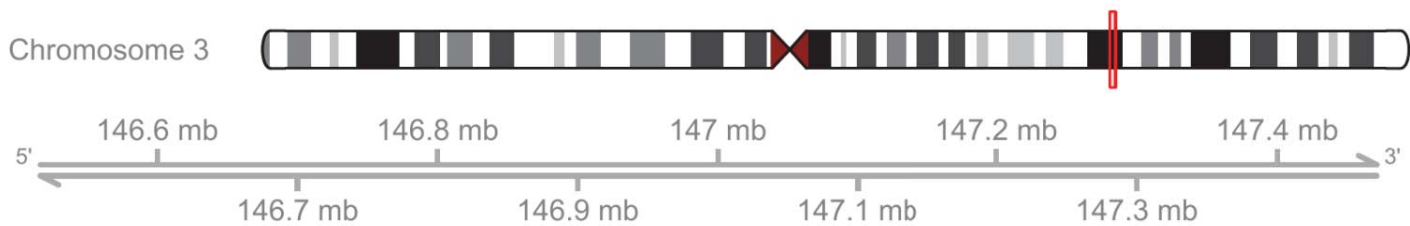
Uncorrected



Zscore of the effect allele: T



Lingual SA: rs61508189



ZIC4

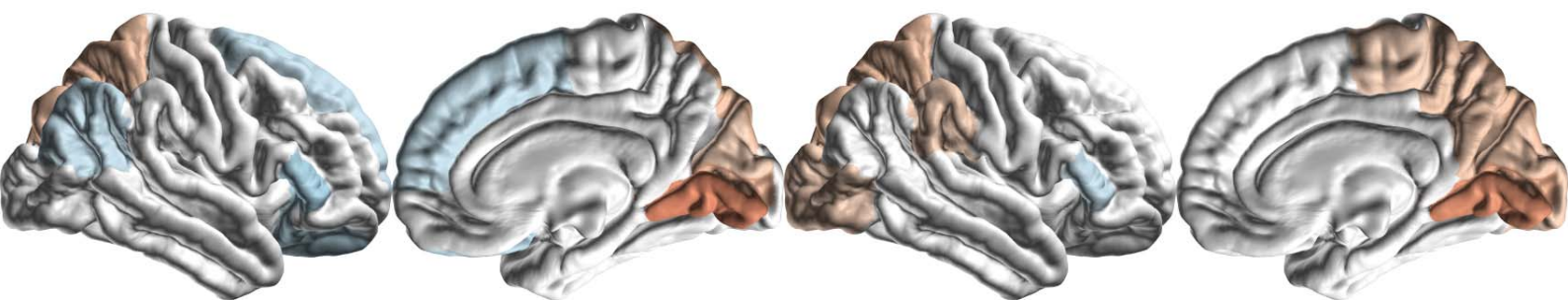
ZIC1

GZ>CP

CP>GZ

Corrected for total SA

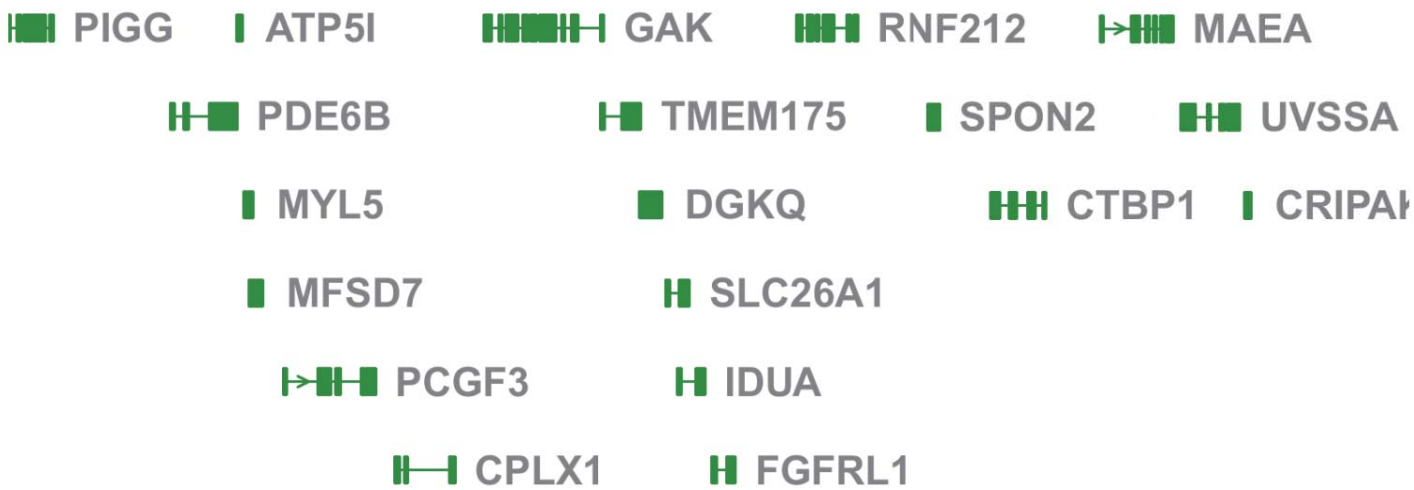
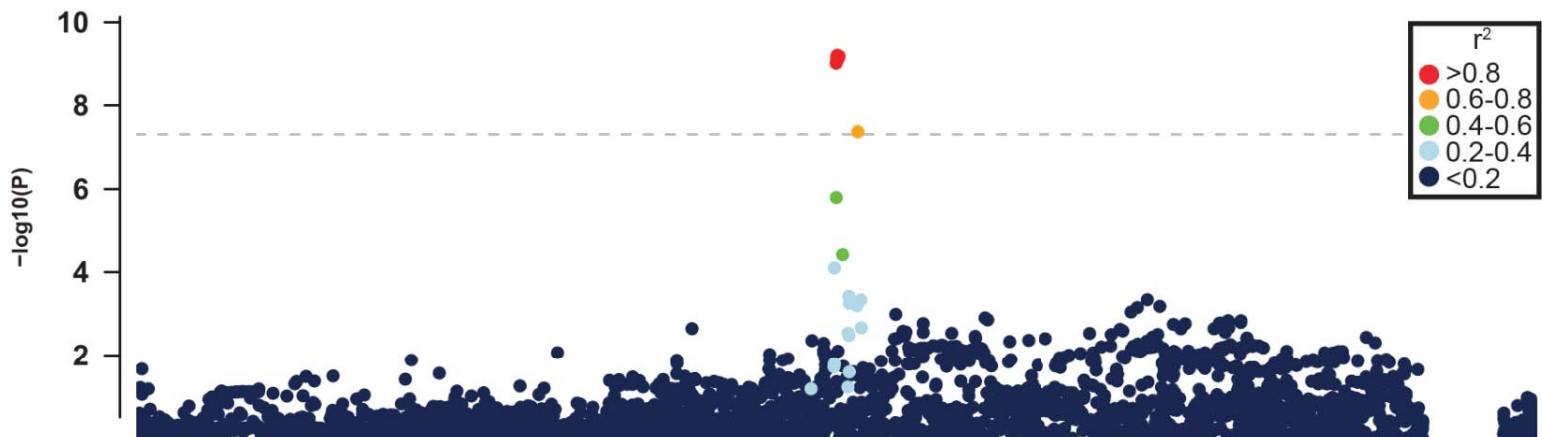
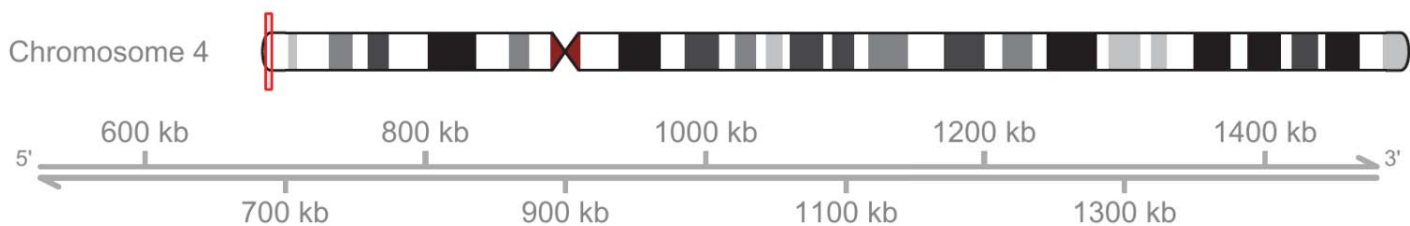
Uncorrected



Zscore of the effect allele: A



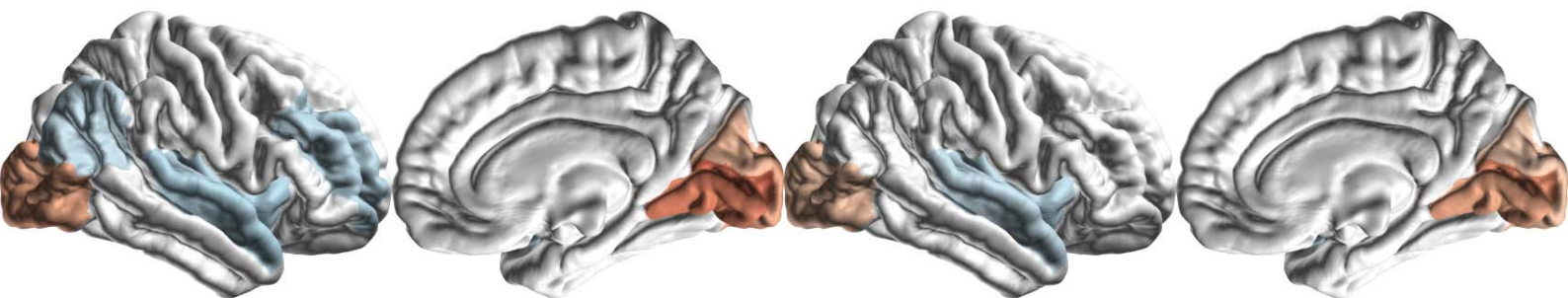
Lingual SA: rs6812278



CP>GZ GZ>CP

Corrected for total SA

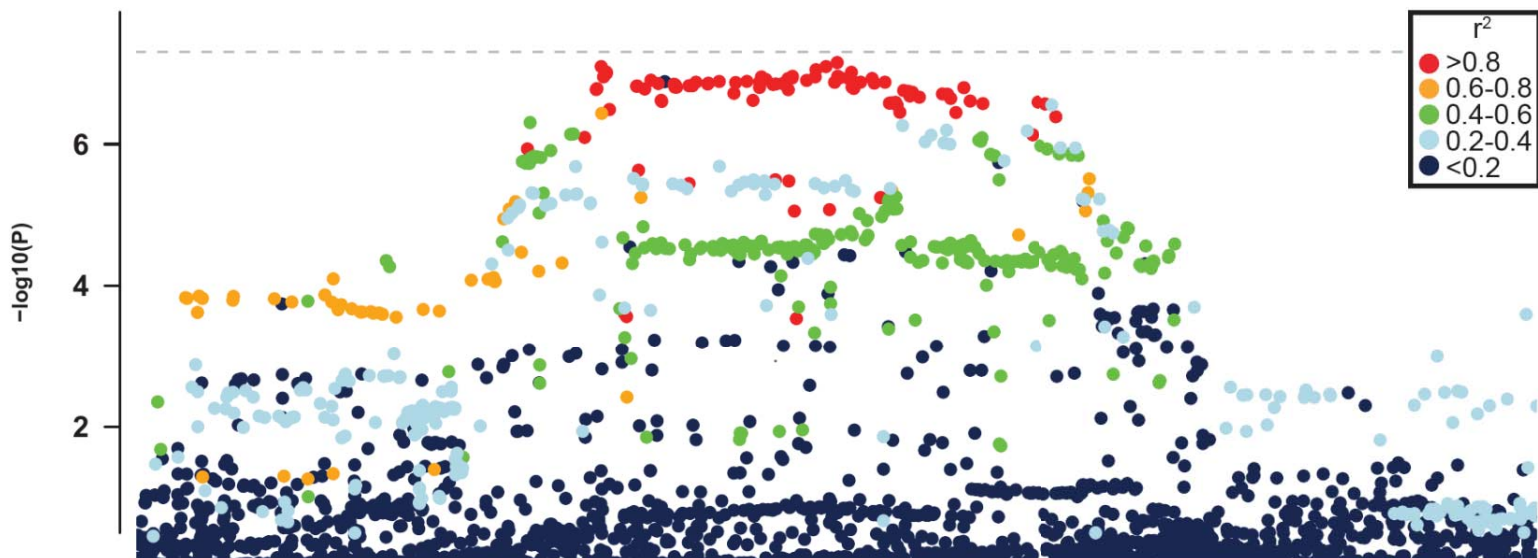
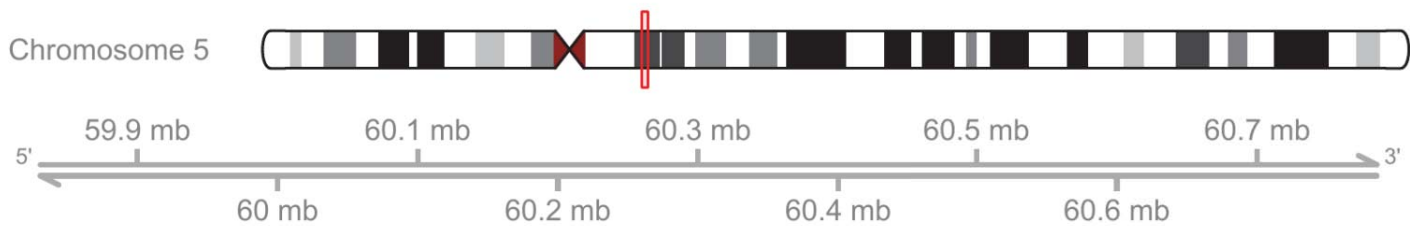
Uncorrected



Zscore of the effect allele: C

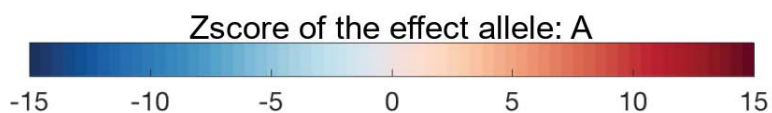
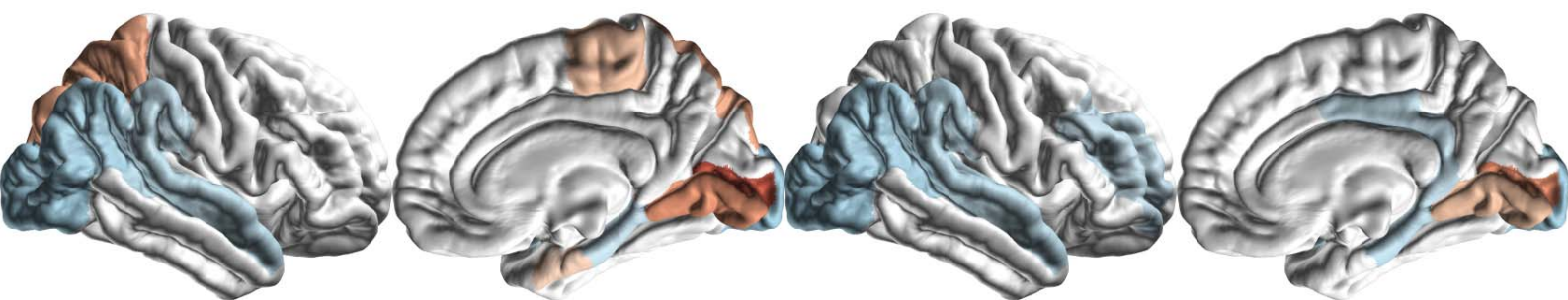


Lingual SA: rs17419290

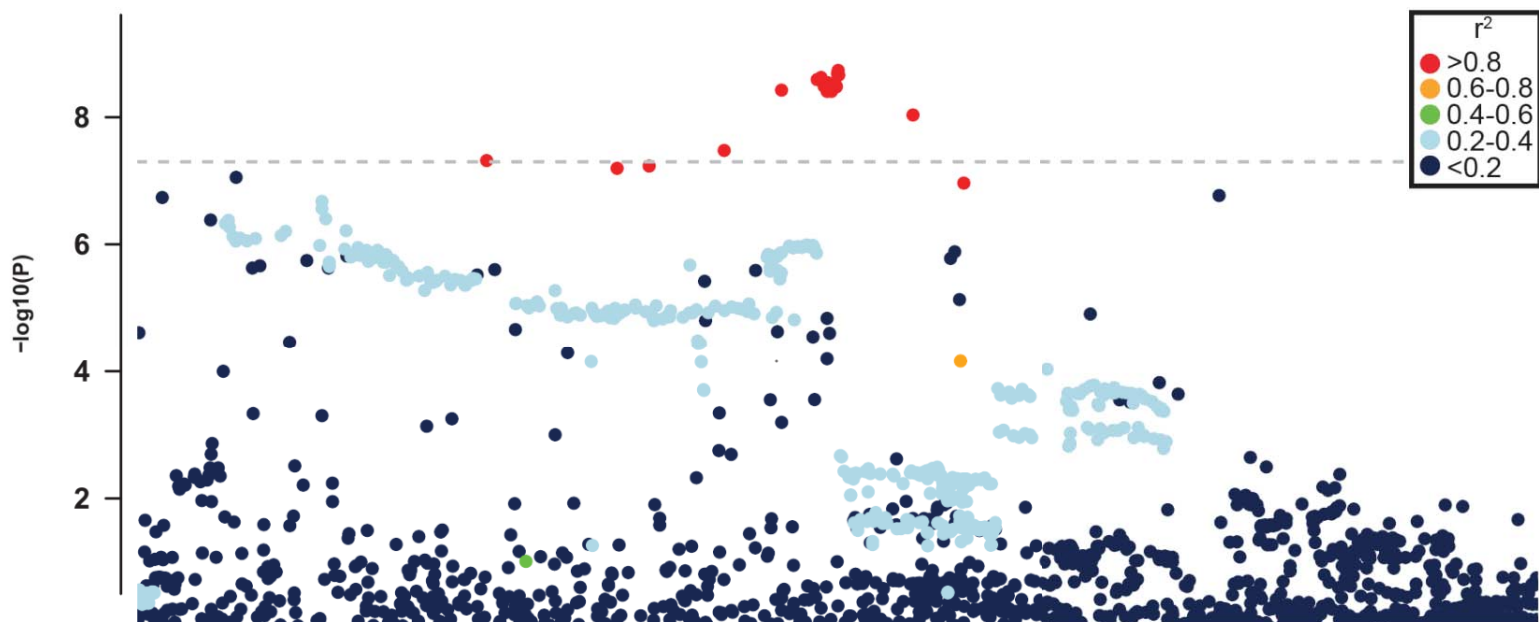
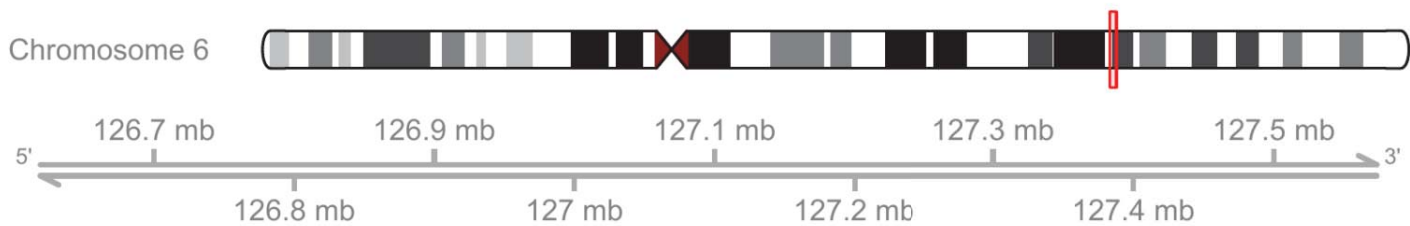


Corrected for total SA

Uncorrected



Lingual SA: rs9401907



CENPW

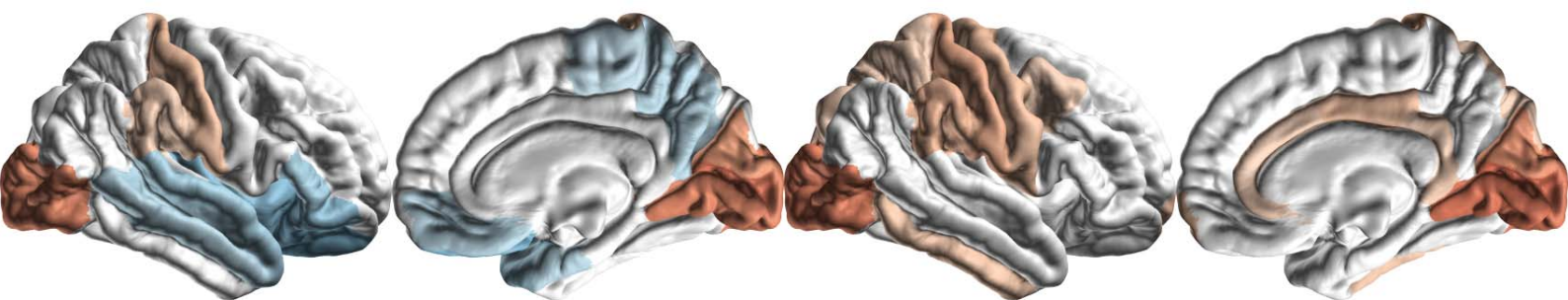
RSP

GZ>CP

CP>GZ

Corrected for total SA

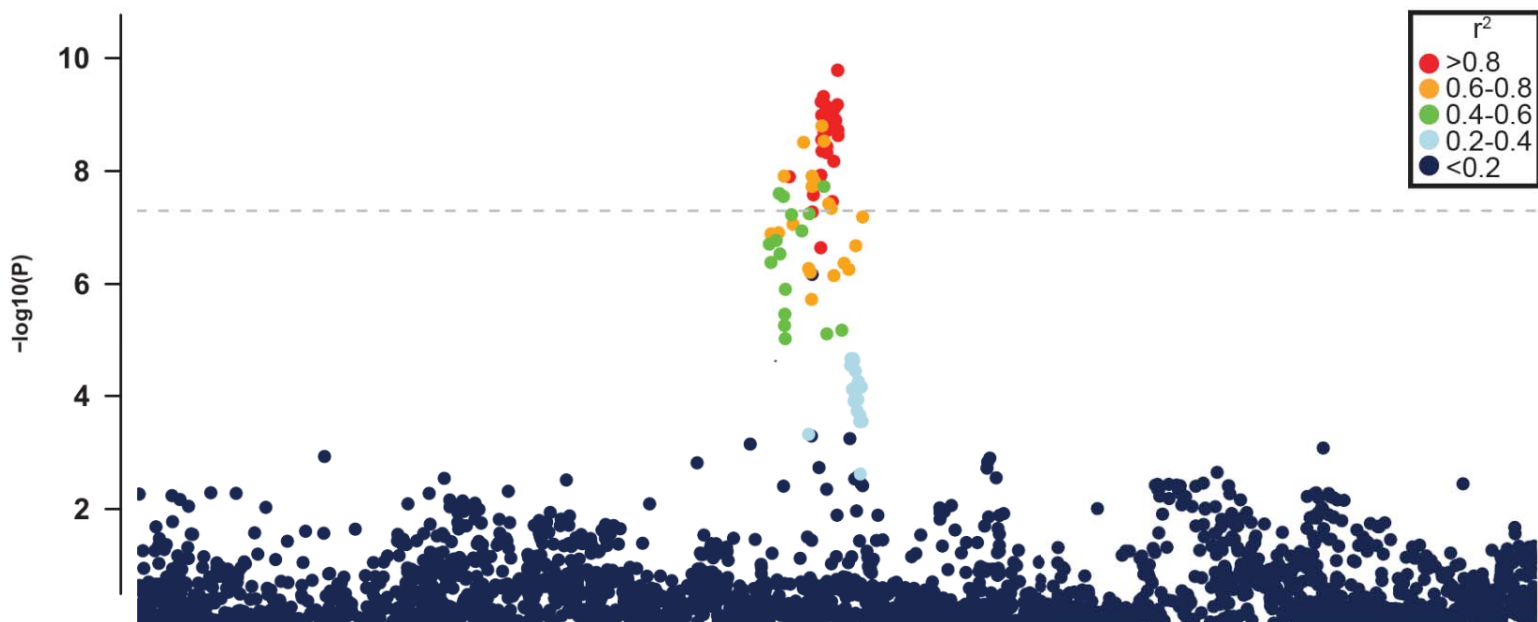
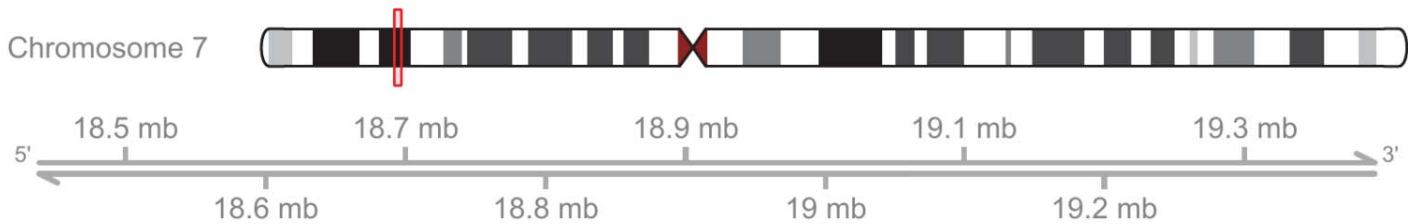
Uncorrected



Zscore of the effect allele: T

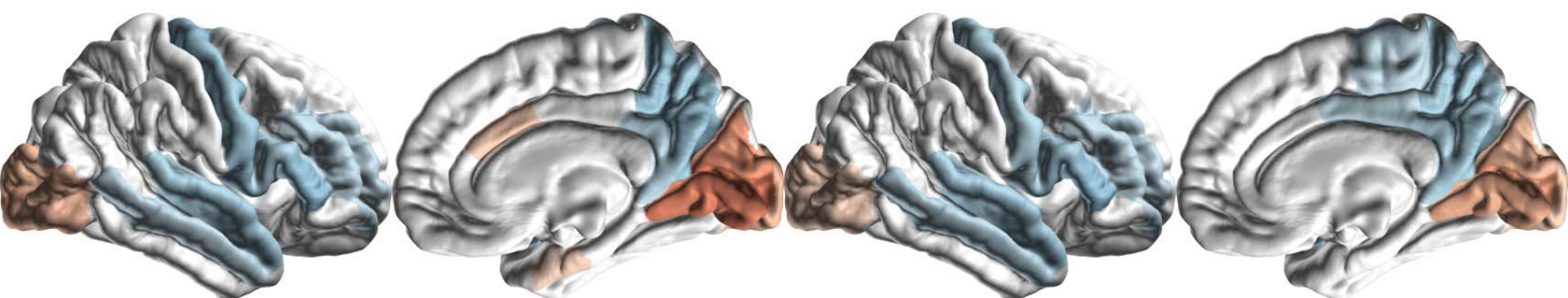


Lingual SA: rs10237366

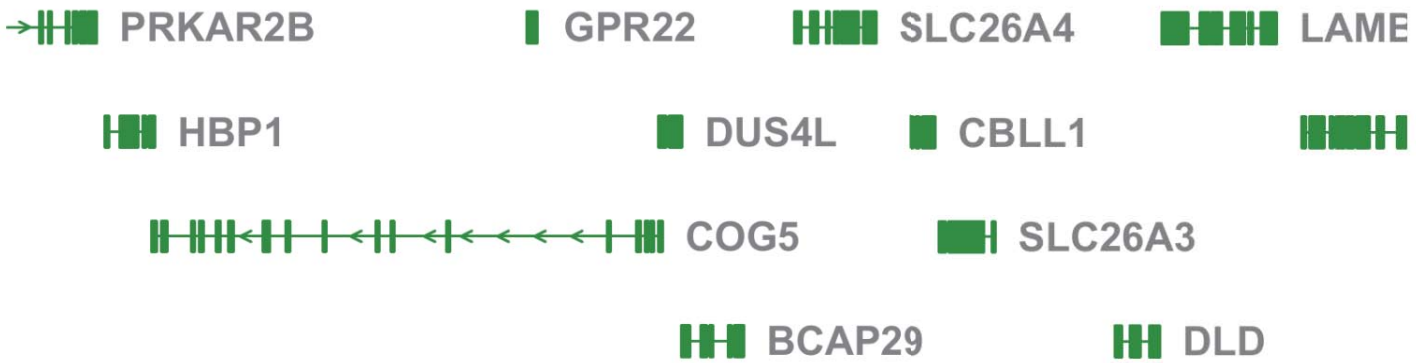
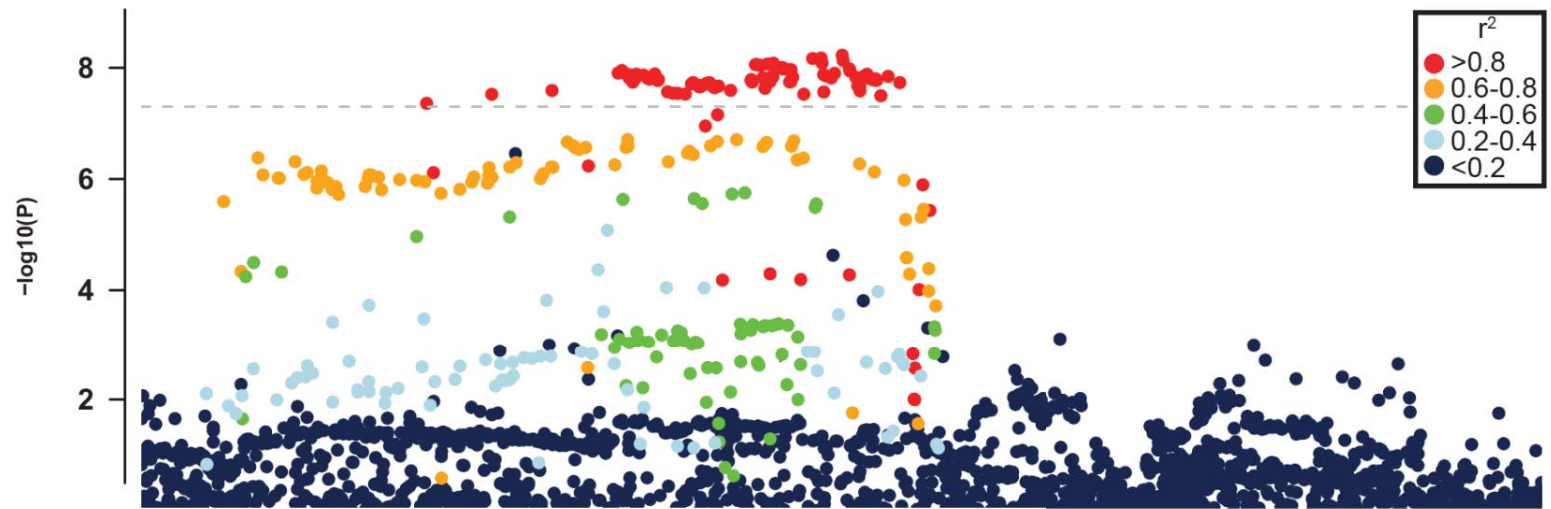
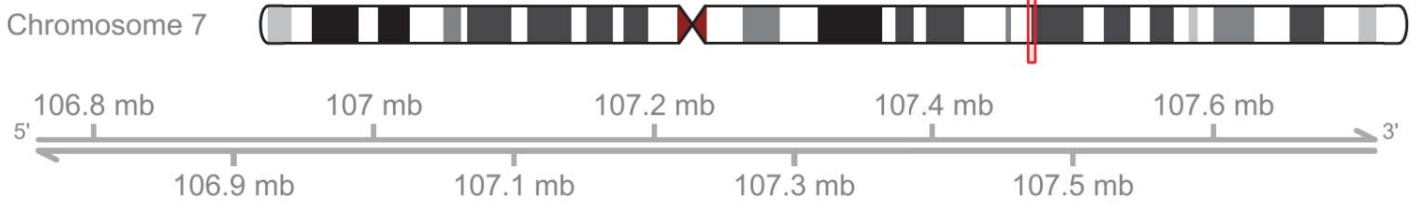


Corrected for total SA

Uncorrected

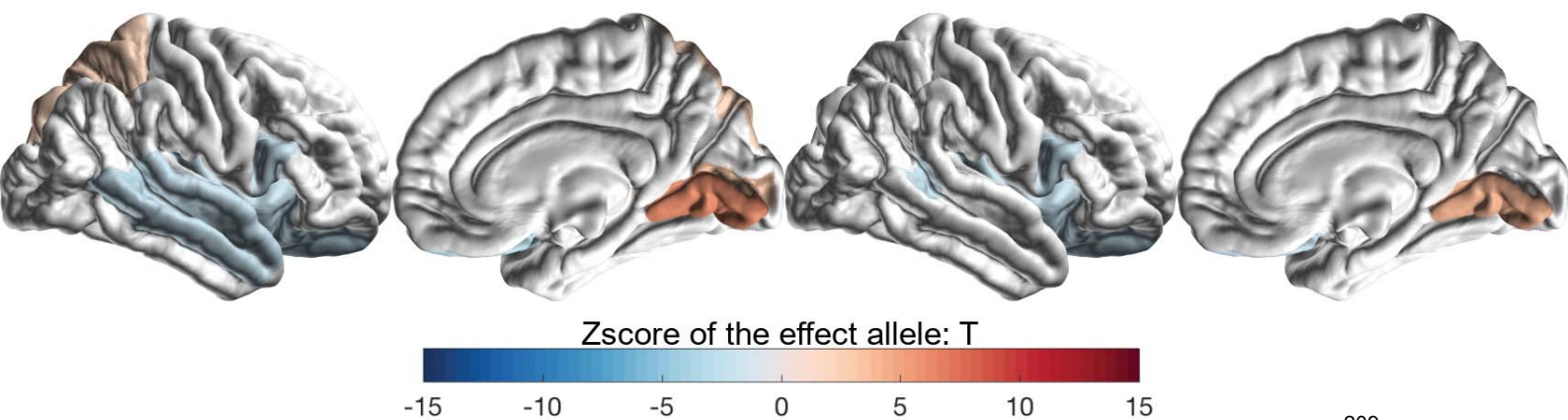


Lingual SA: rs7809950

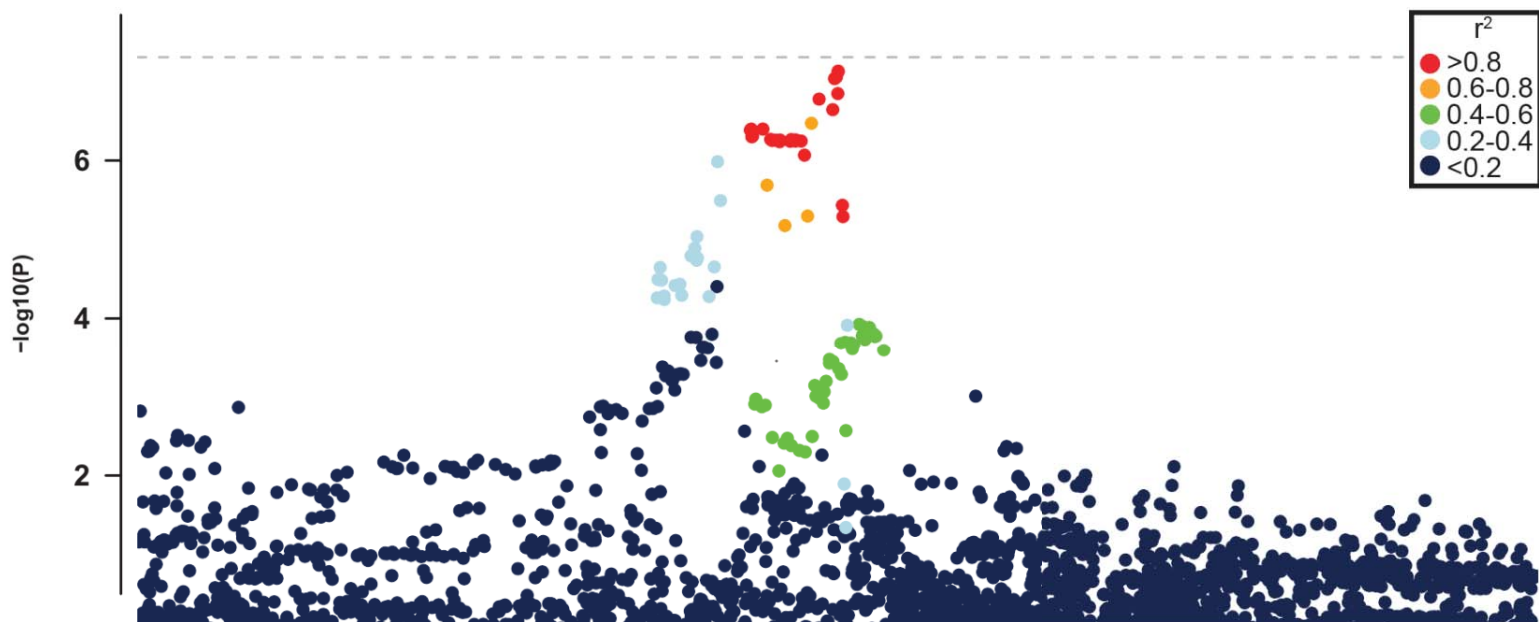
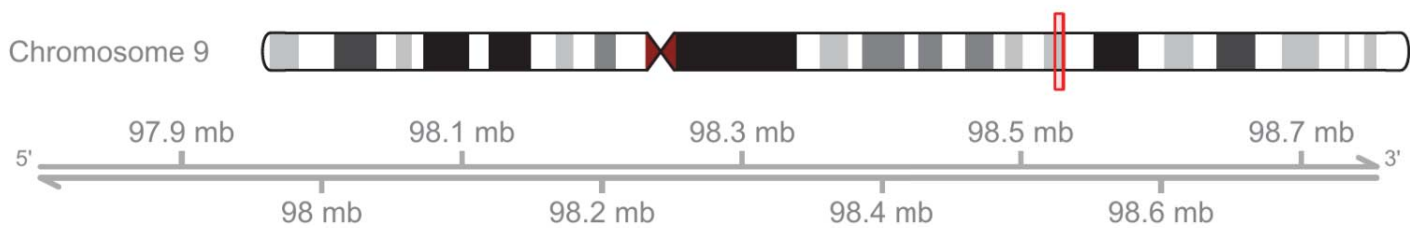


Corrected for total SA

Uncorrected



Lingual SA: rs28410513

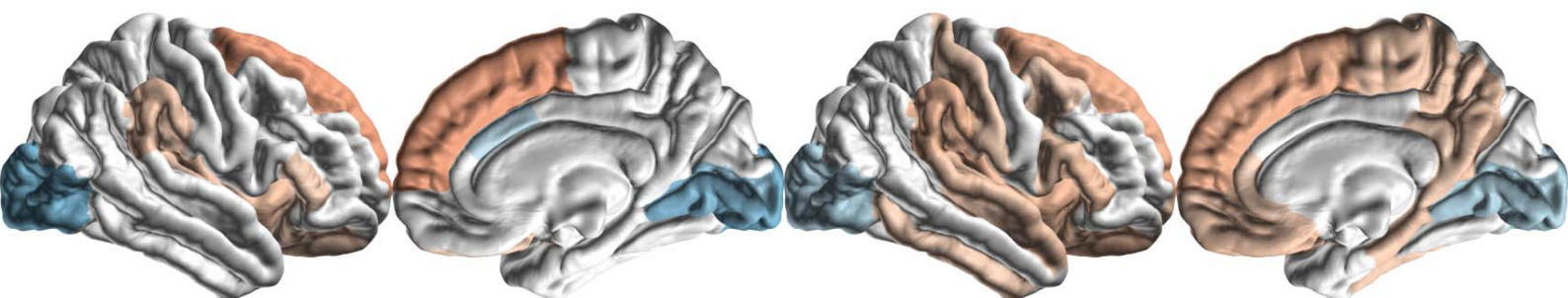


GZ>CP
CP>GZ



Corrected for total SA

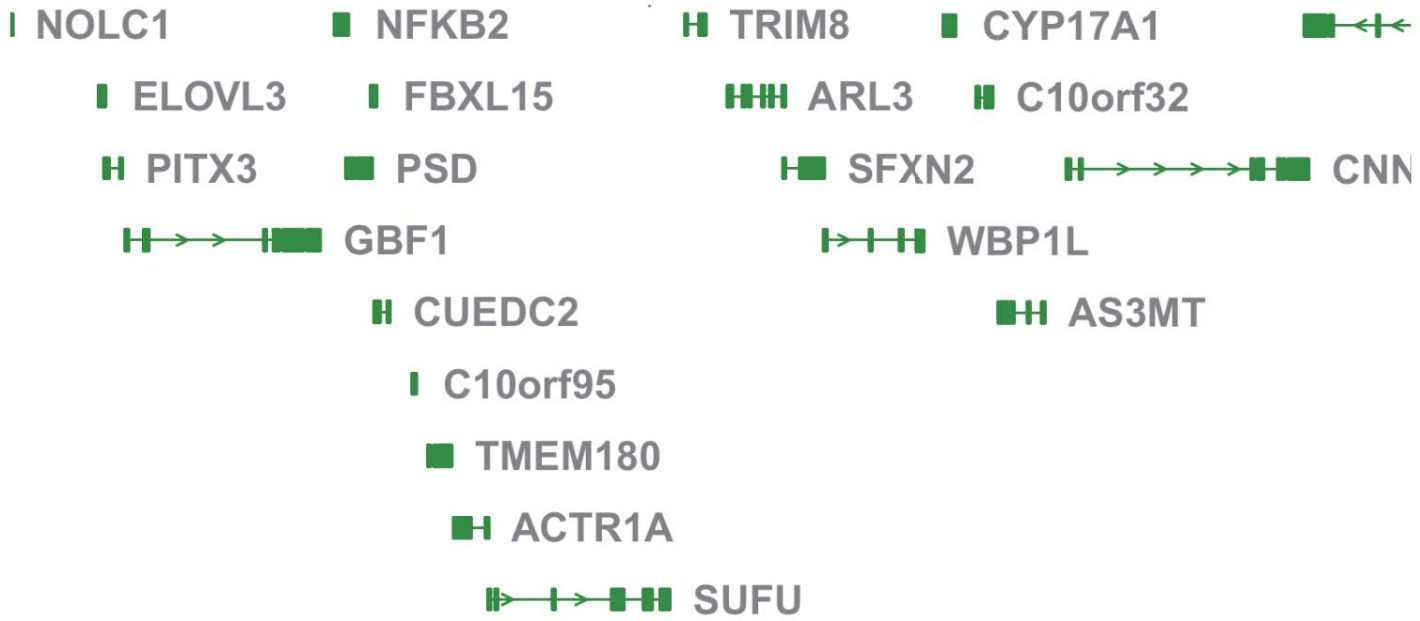
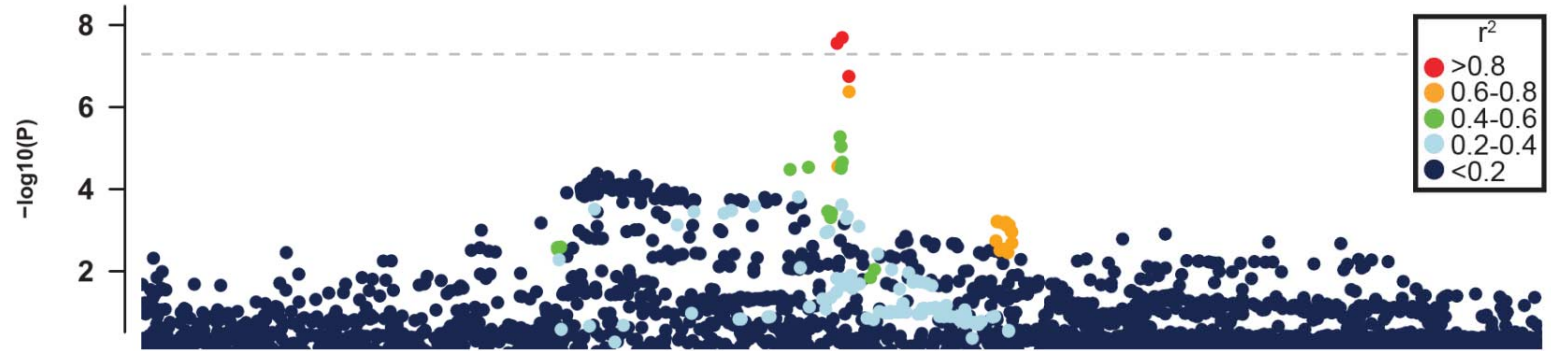
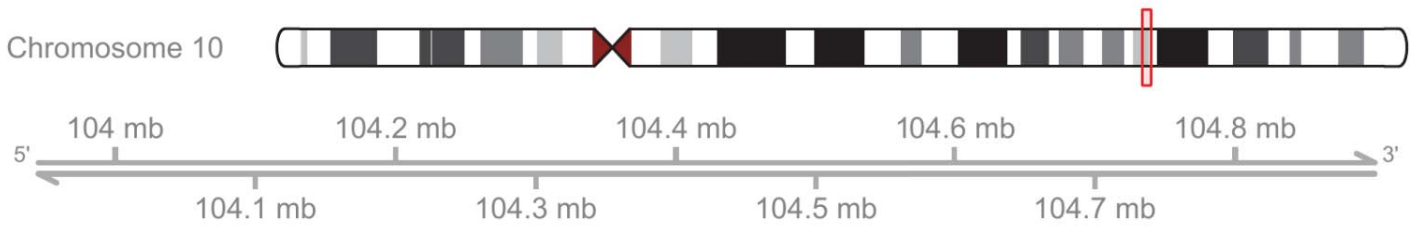
Uncorrected



Zscore of the effect allele: T



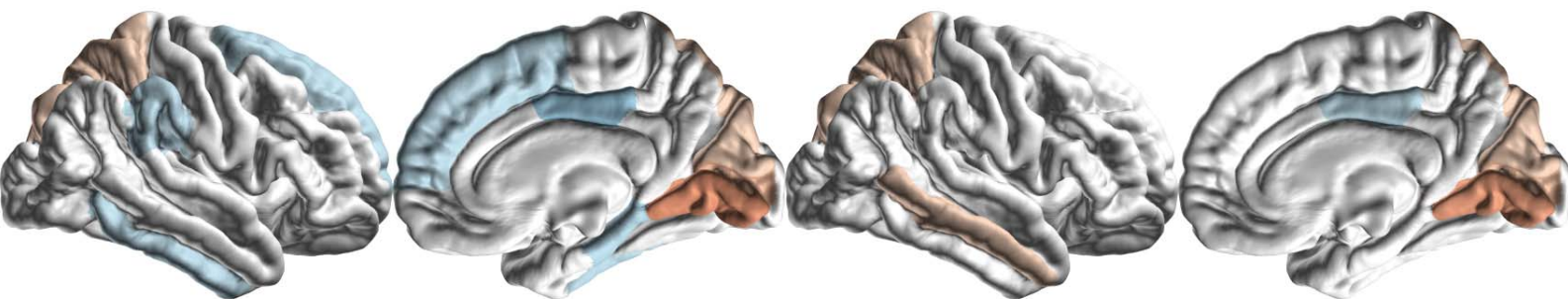
Lingual SA: rs7914158



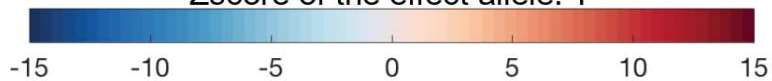
CP>GZGZ>CP

Corrected for total SA

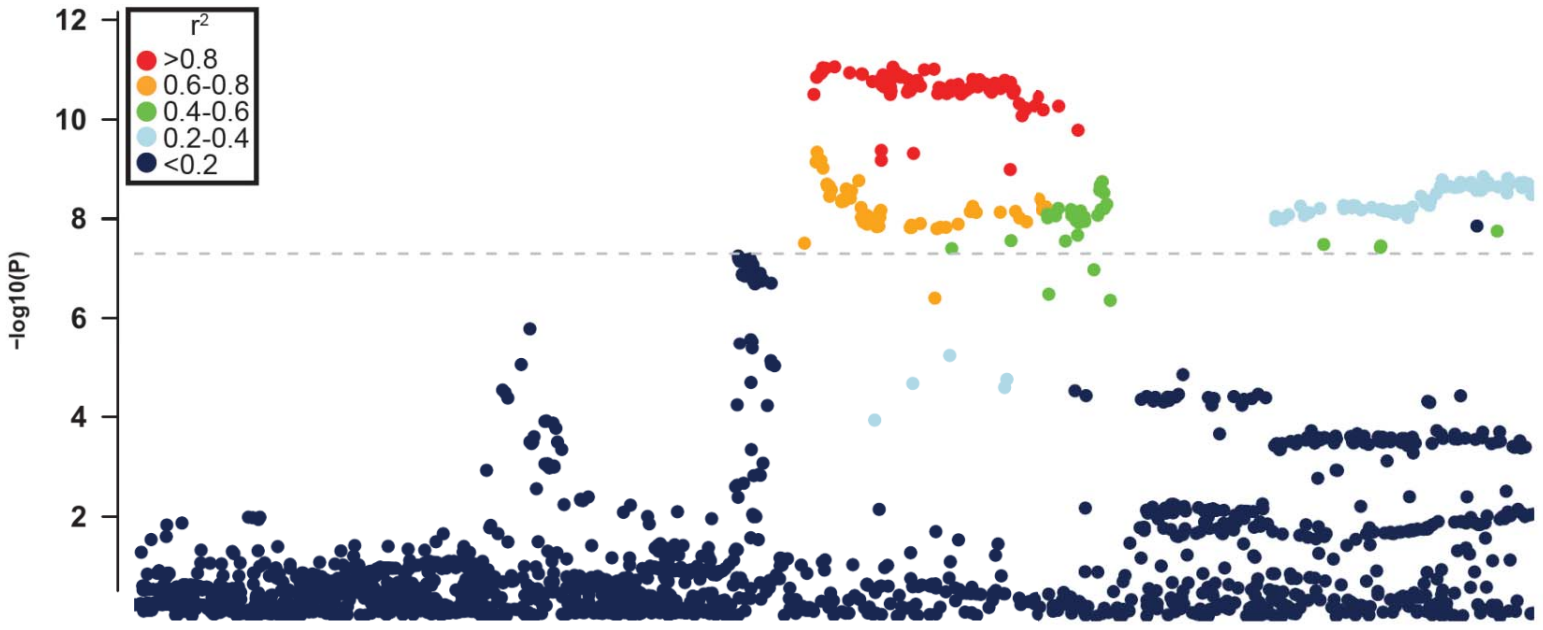
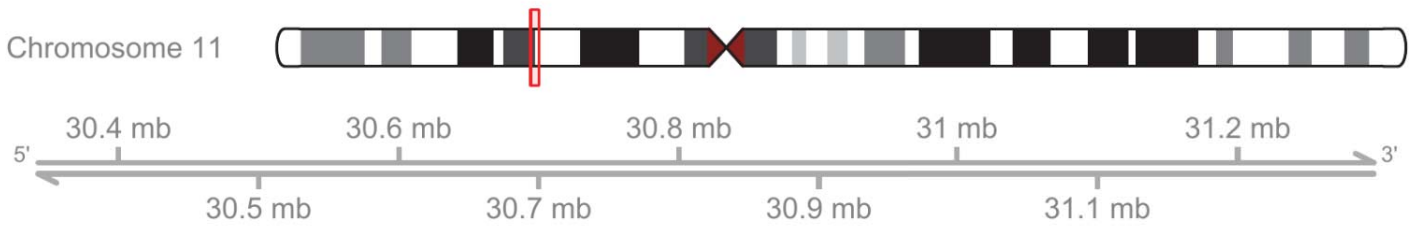
Uncorrected



Zscore of the effect allele: T



Lingual SA: rs2022130



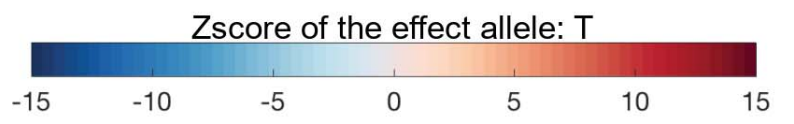
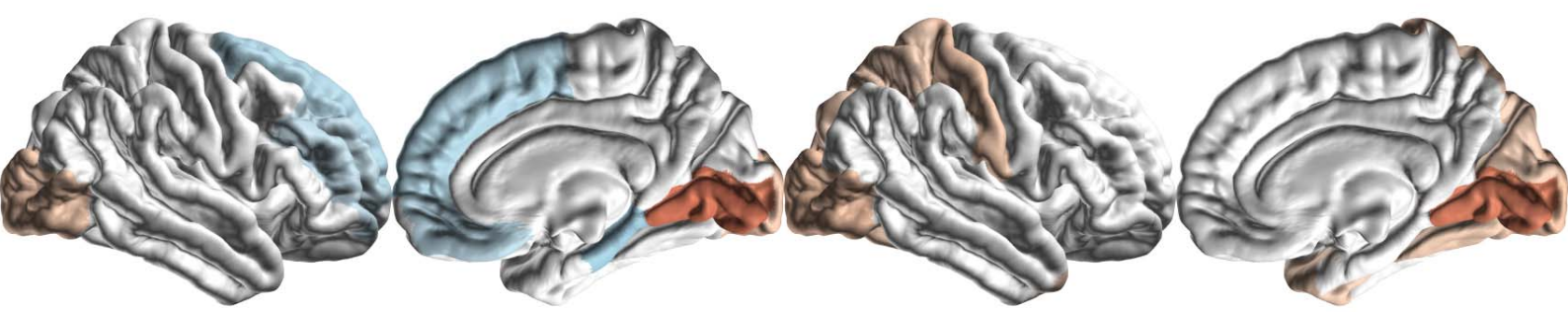
ARL14EP

MPPED2



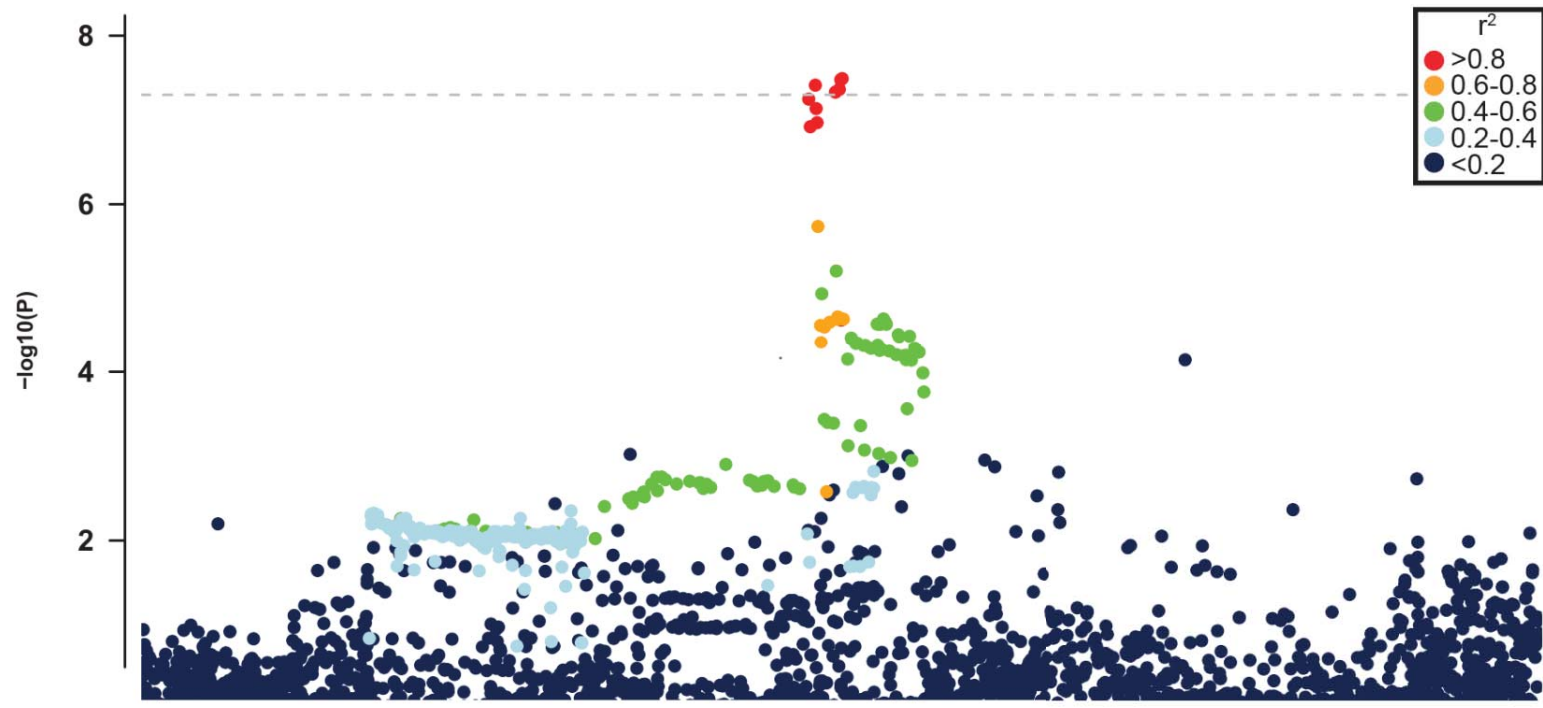
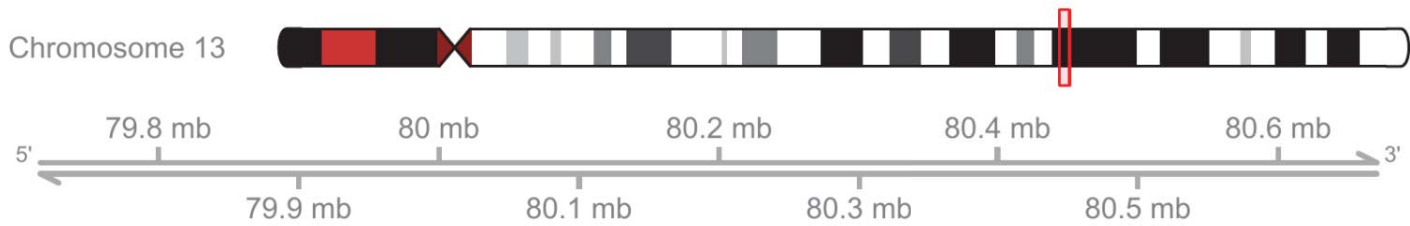
Corrected for total SA

Uncorrected



Lingual SA: rs9545158

Chromosome 13

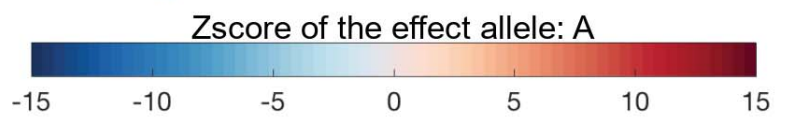
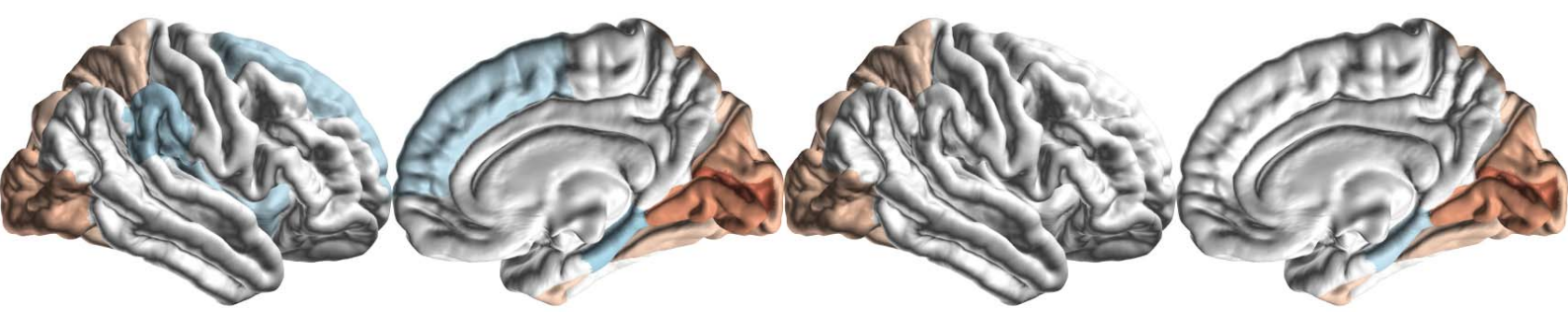


RBM26 NDFIP2



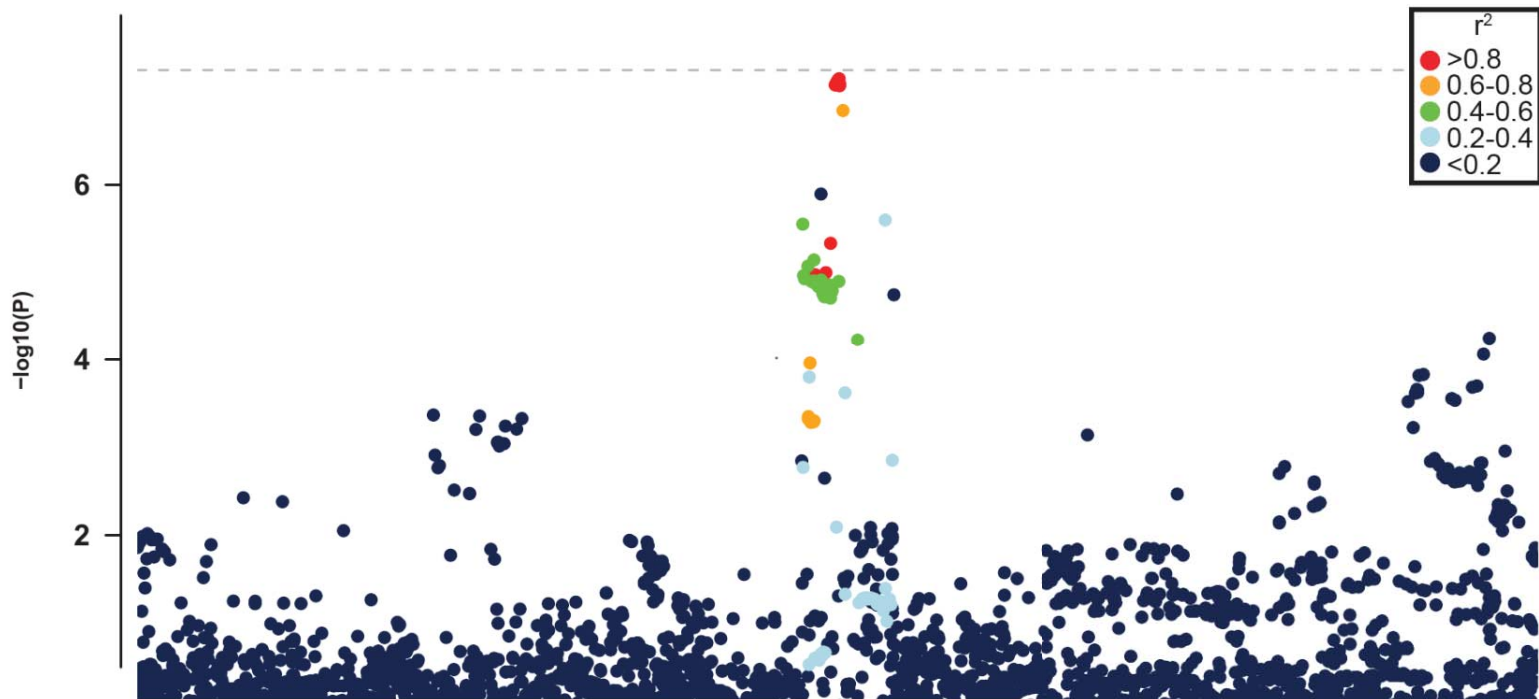
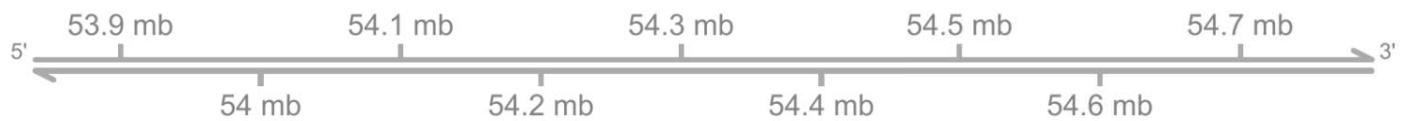
Corrected for total SA

Uncorrected



Lingual SA: rs10431624

Chromosome 14



BMP4

GZ>CP

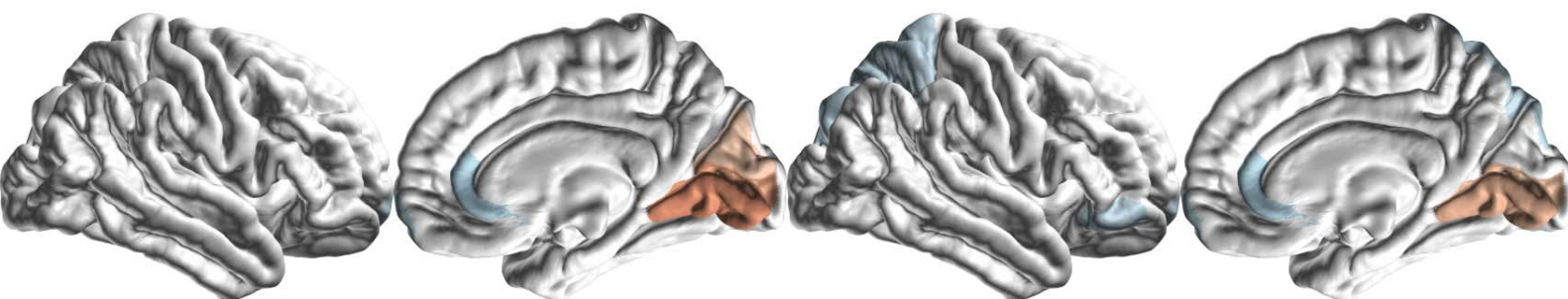


CP>GZ



Corrected for total SA

Uncorrected

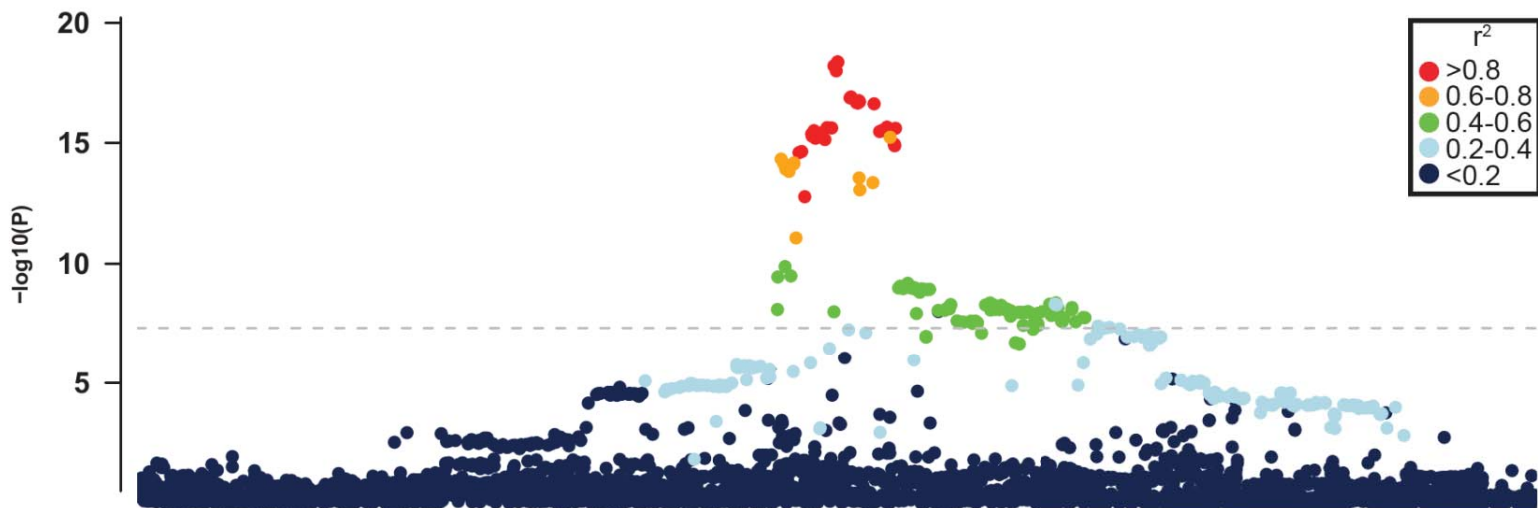
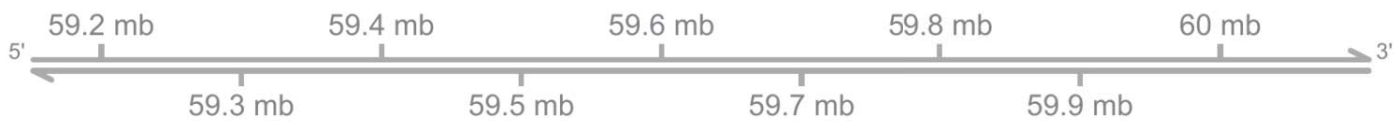


Zscore of the effect allele: T



Lingual SA: rs74826997

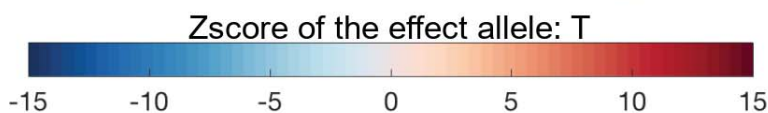
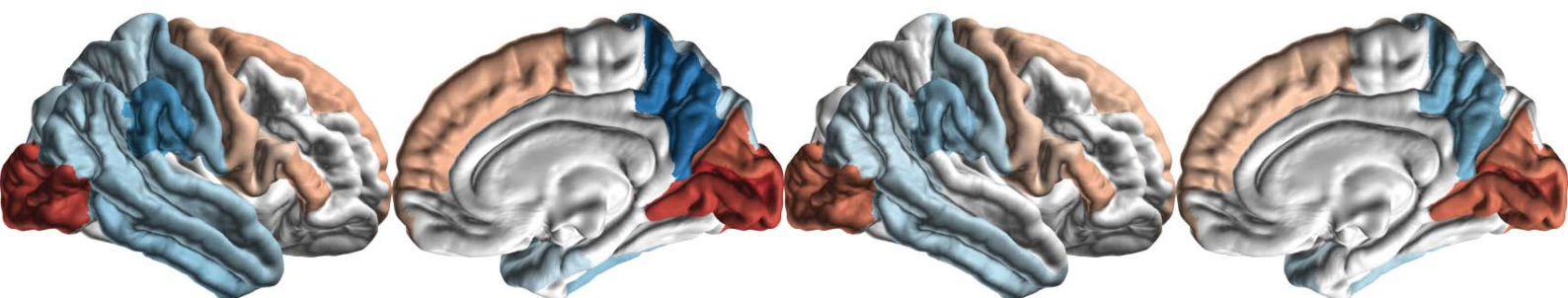
Chromosome 14



CP>GZ
GZ>CP

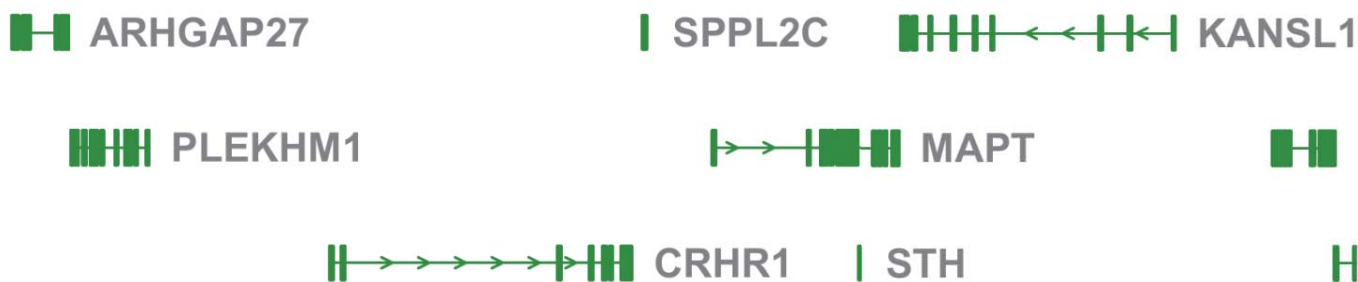
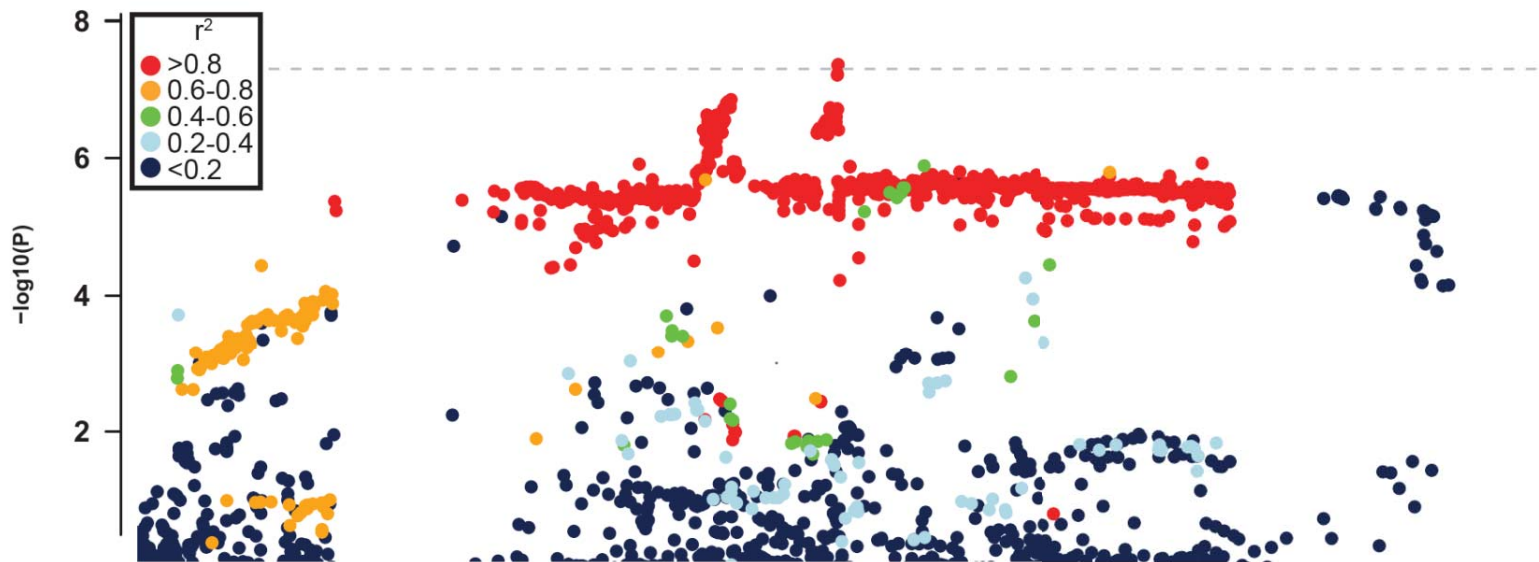
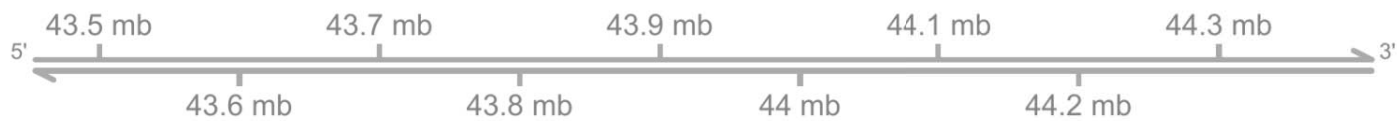
Corrected for total SA

Uncorrected



Lingual SA: rs17690987

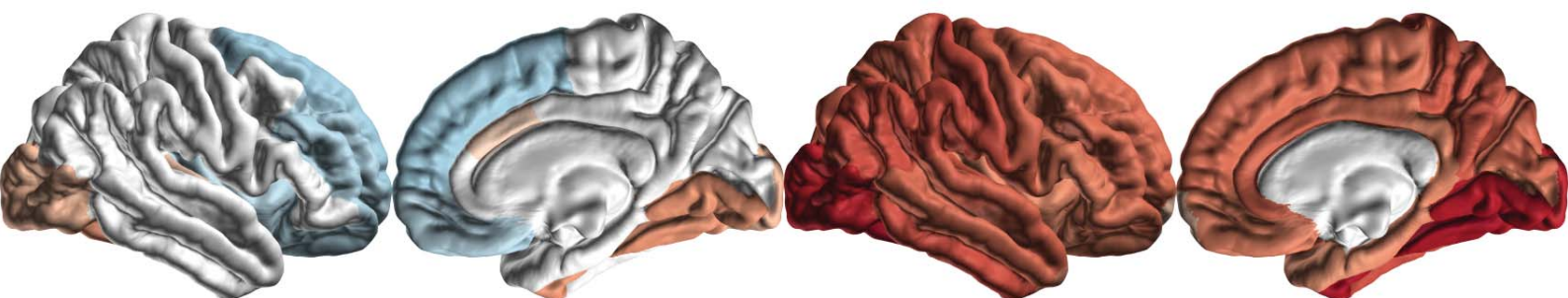
Chromosome 17



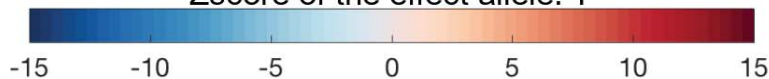
GZ>CP
CP>GZ

Corrected for total SA

Uncorrected

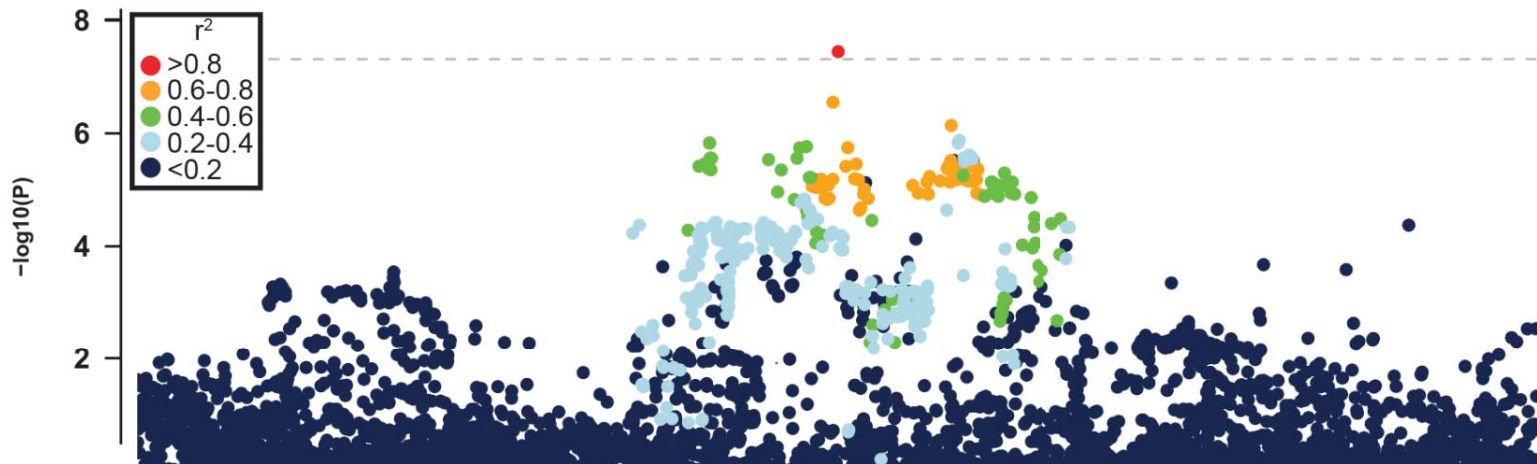
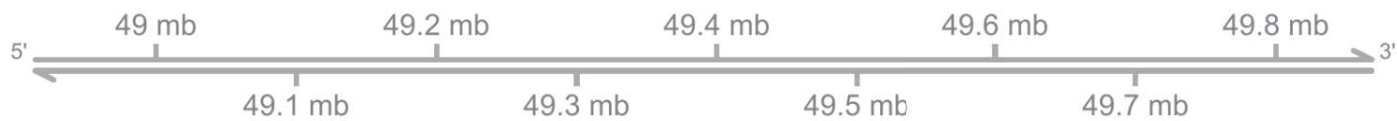


Zscore of the effect allele: T



Lingual TH: rs10495963

Chromosome 2



STON1-GTF2A1L

GTF2A1L

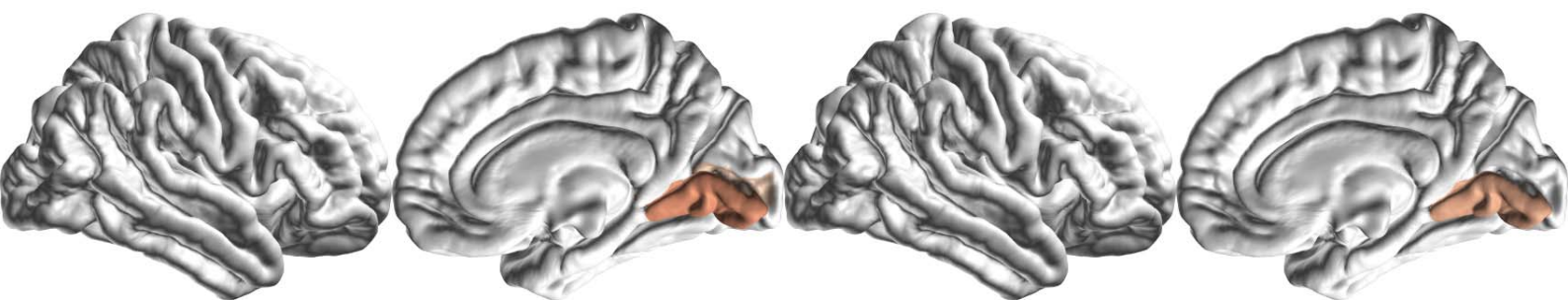
LHCGR

Angular Temporal dIPFC Cingulate

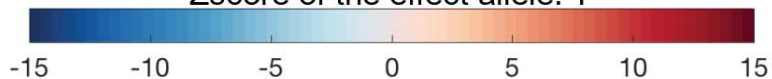


Corrected for average TH

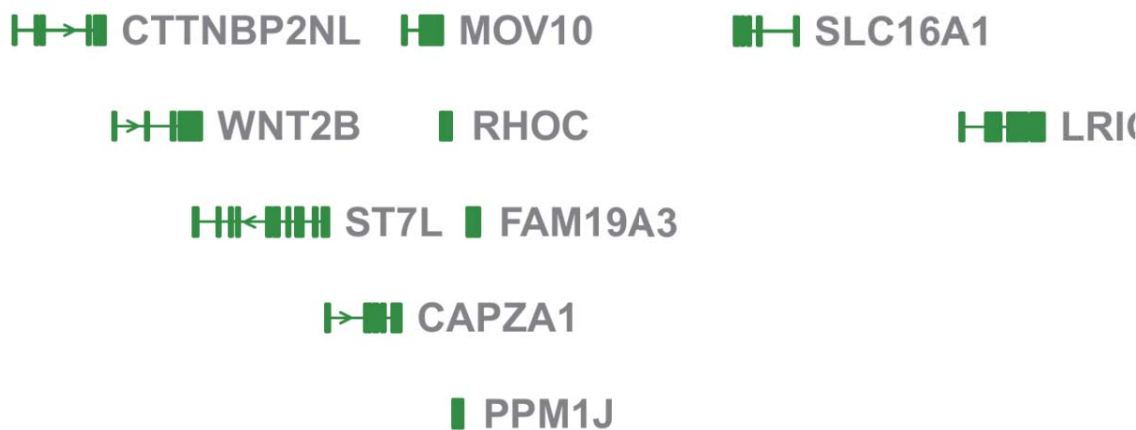
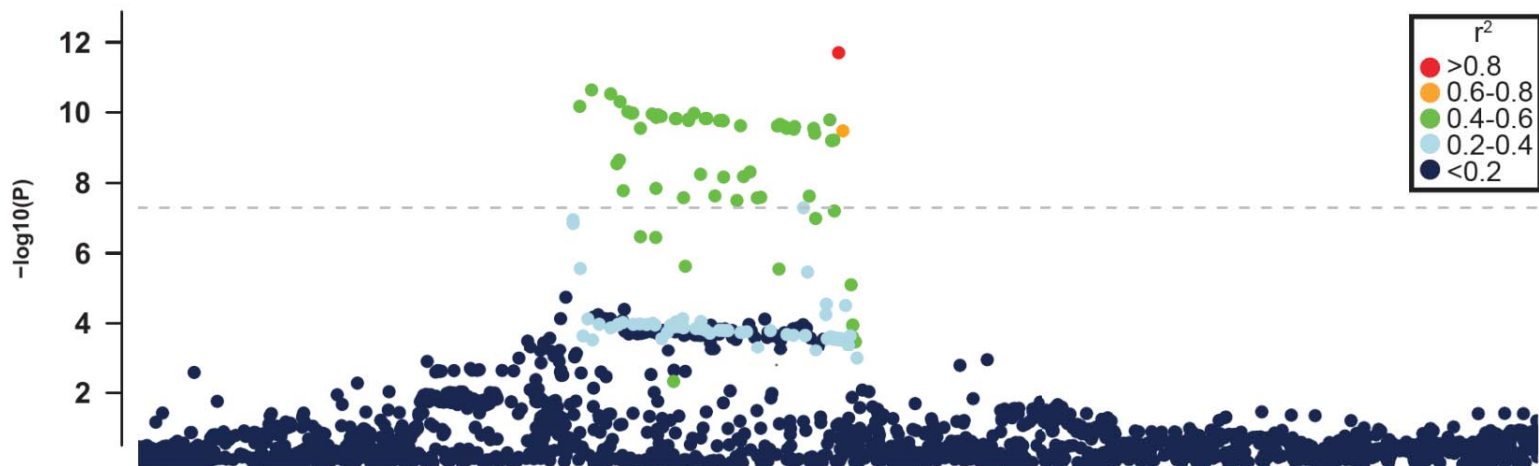
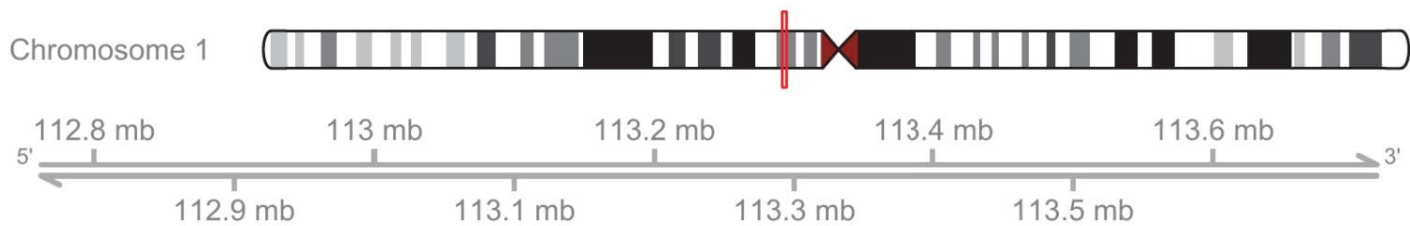
Uncorrected



Zscore of the effect allele: T



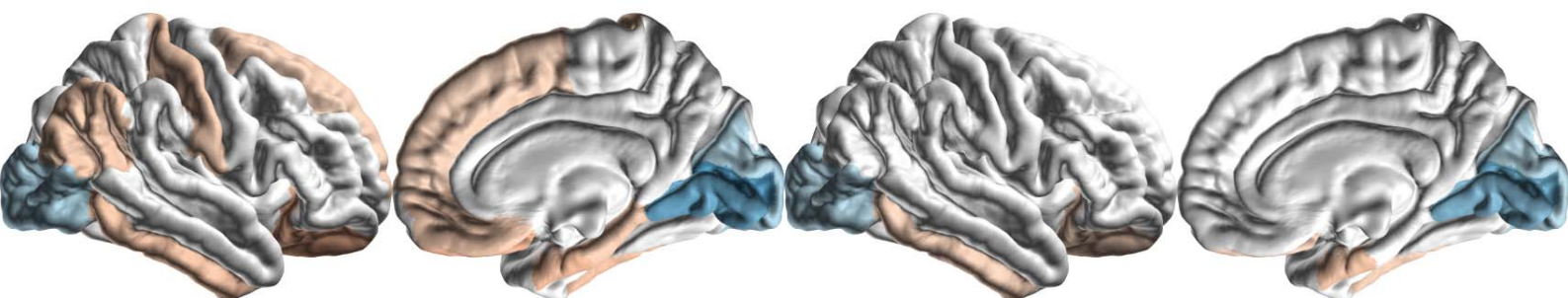
Pericalcarine SA: rs2999158



CP>GZ GZ>CP

Corrected for total SA

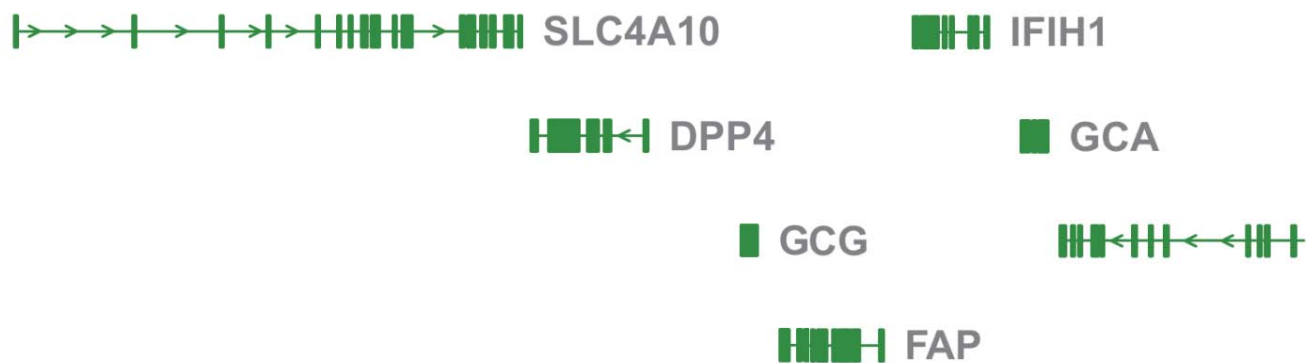
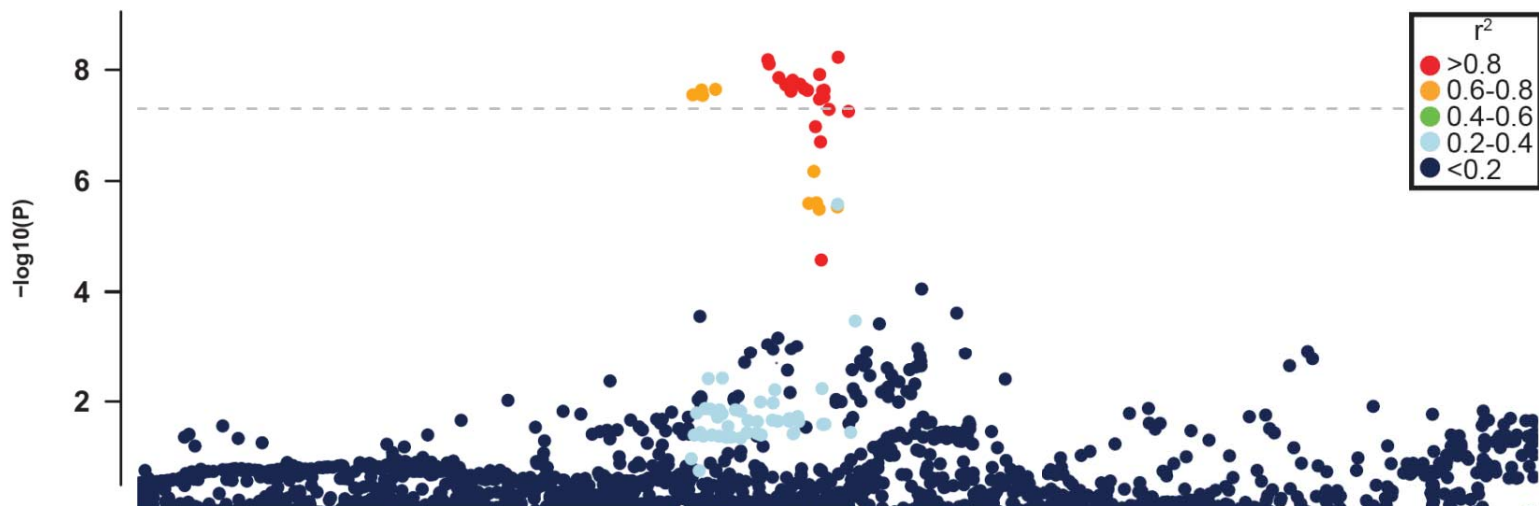
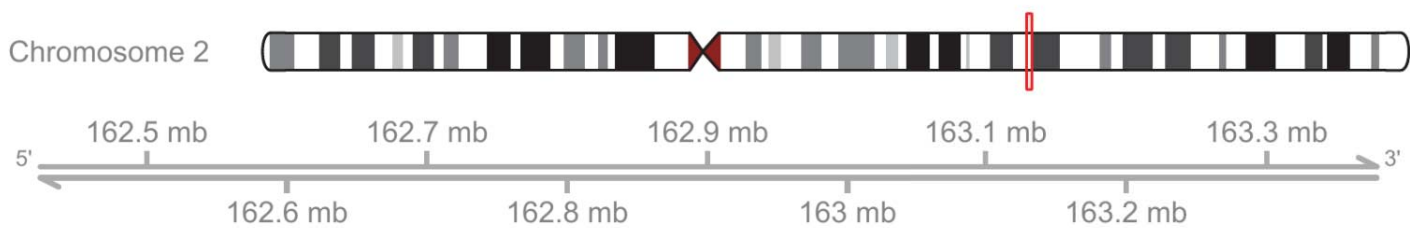
Uncorrected



Zscore of the effect allele: T

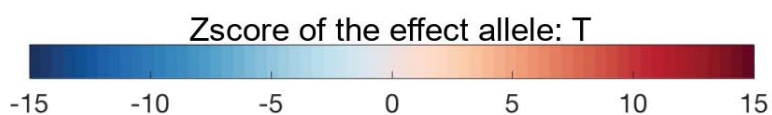
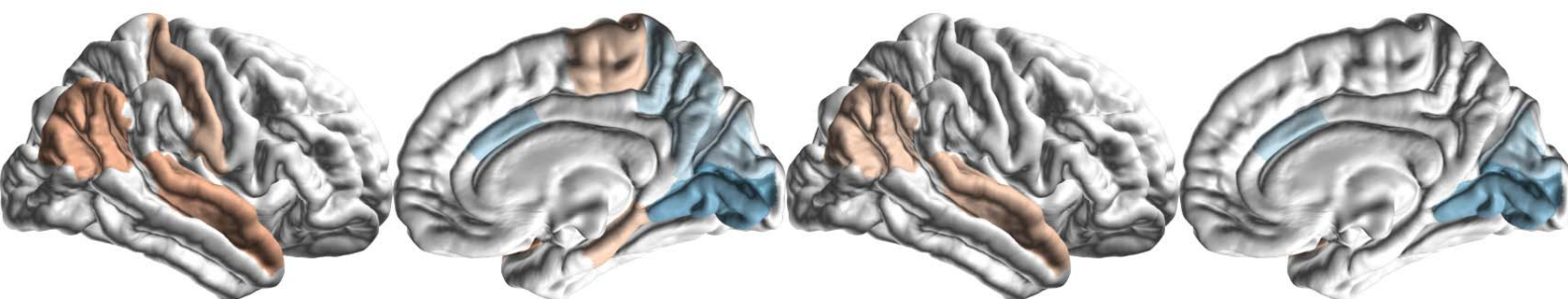


Pericalcarine SA: rs16822665



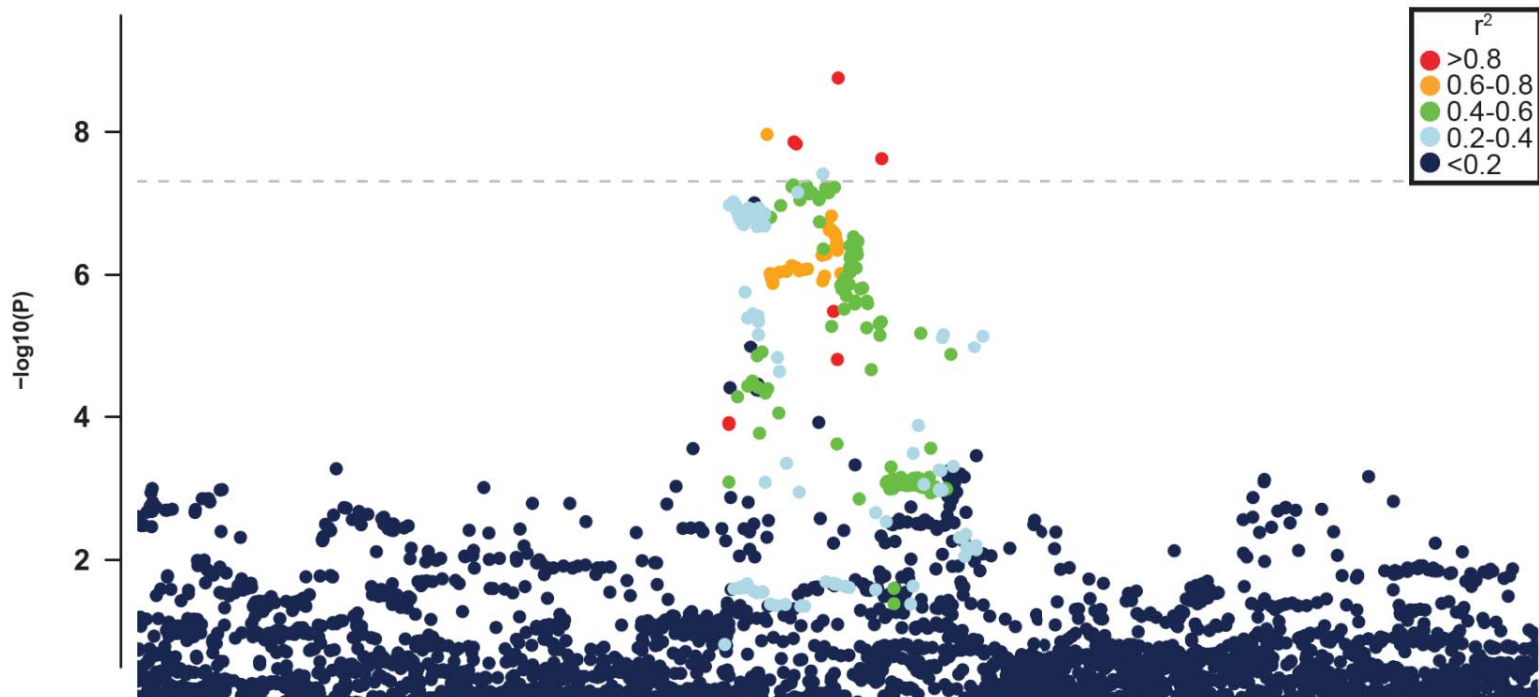
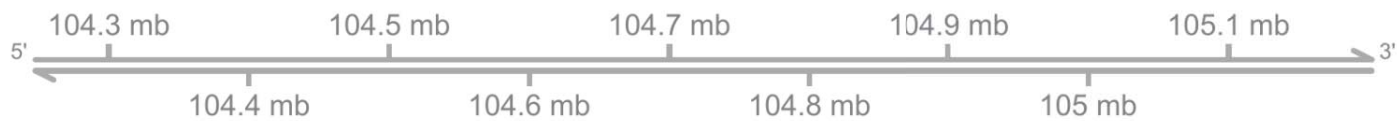
Corrected for total SA

Uncorrected



Pericalcarine SA: rs971550

Chromosome 3



GZ>CP

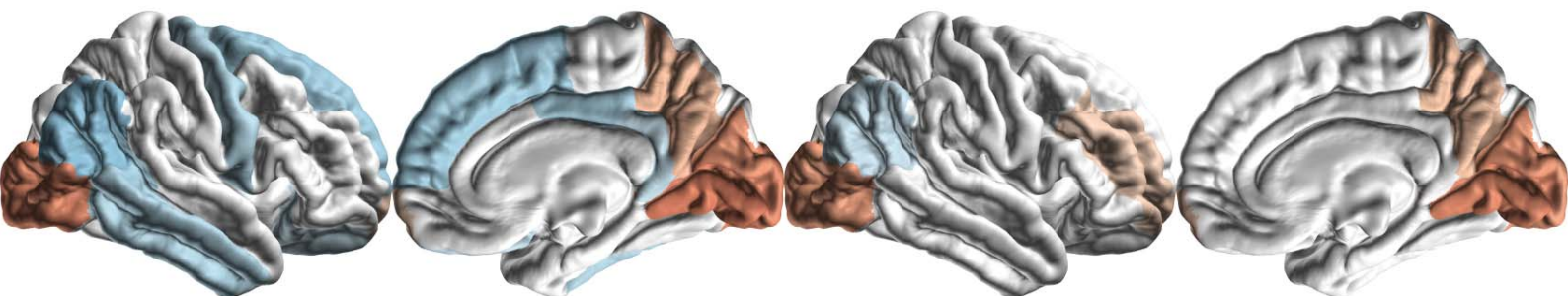


CP>GZ

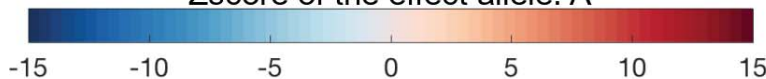


Corrected for total SA

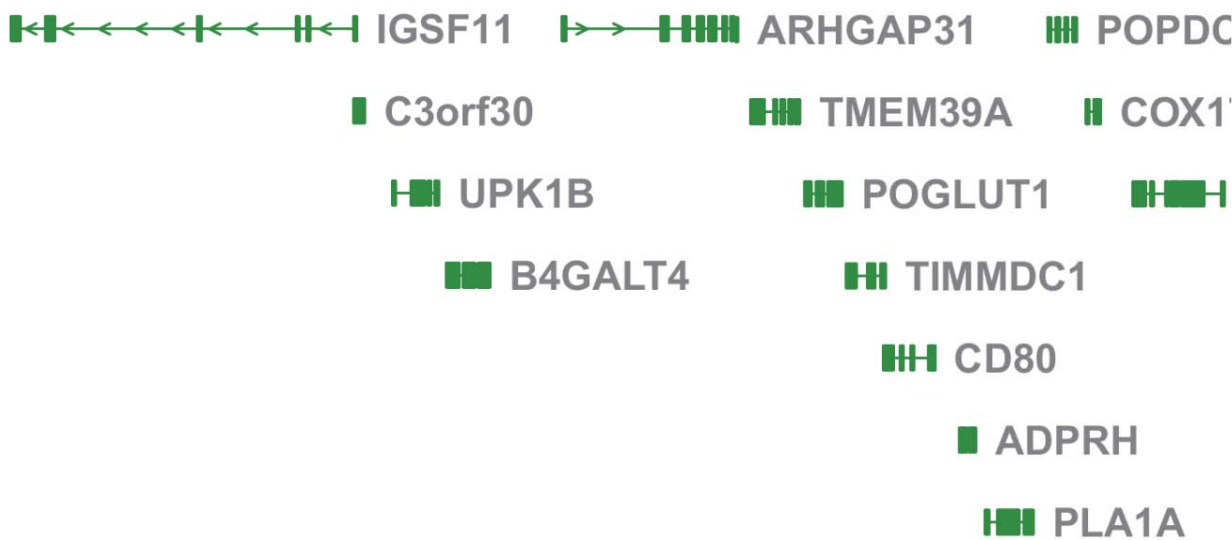
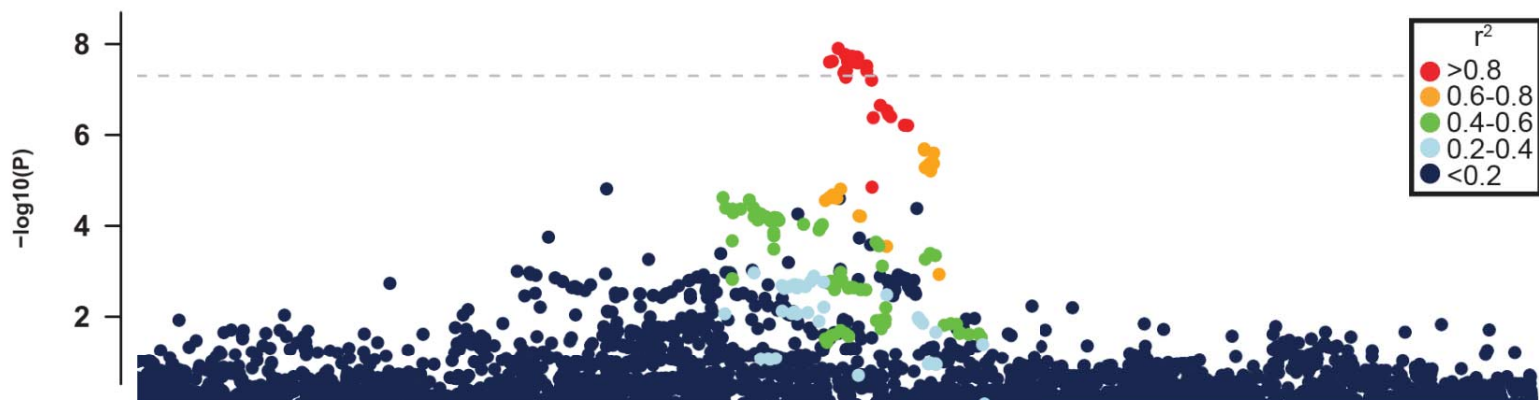
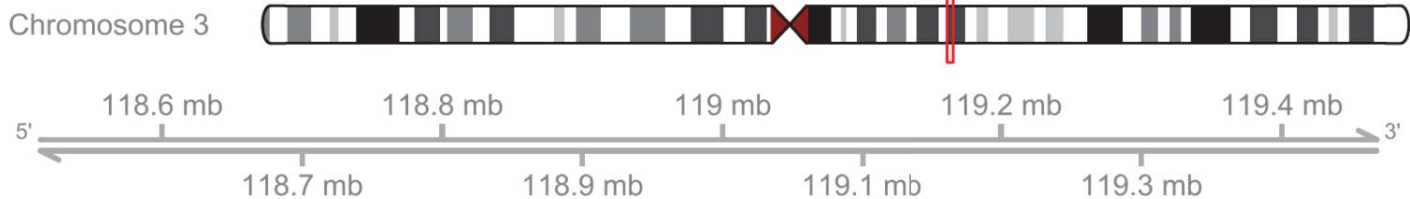
Uncorrected



Zscore of the effect allele: A

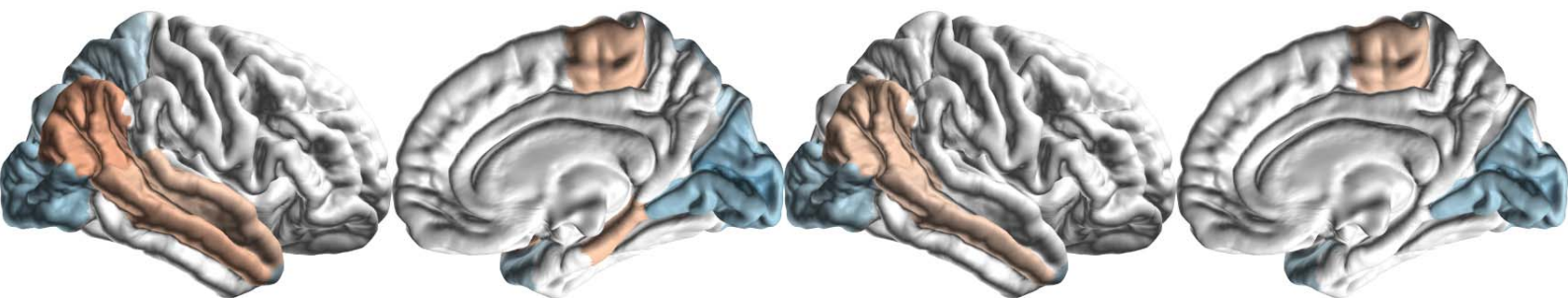


Pericalcarine SA: rs139834736



Corrected for total SA

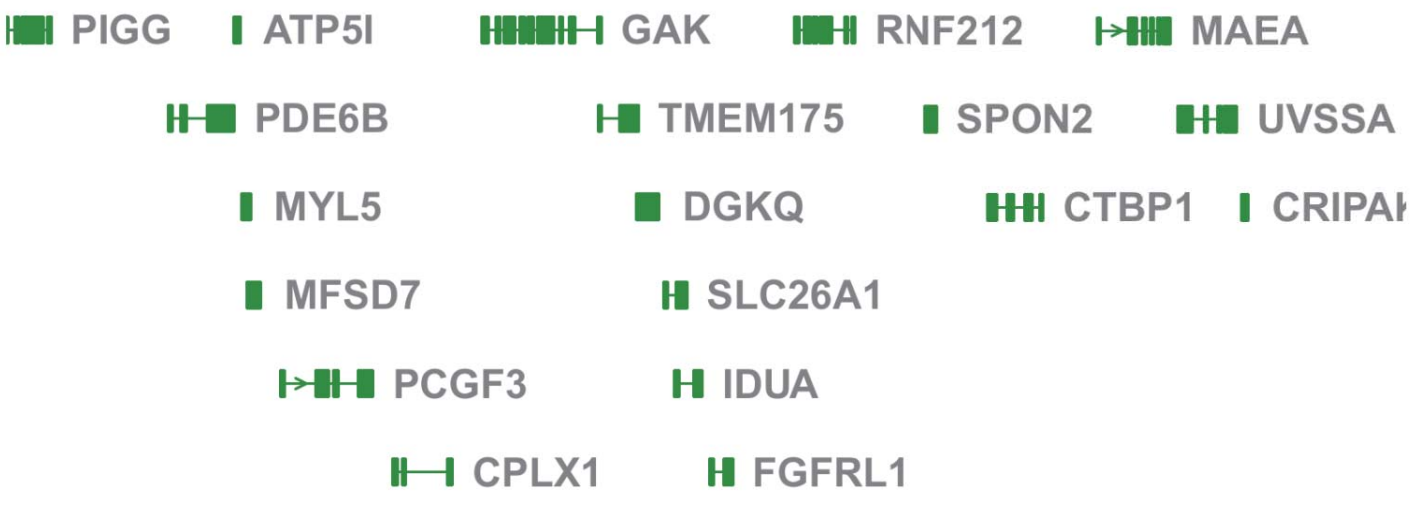
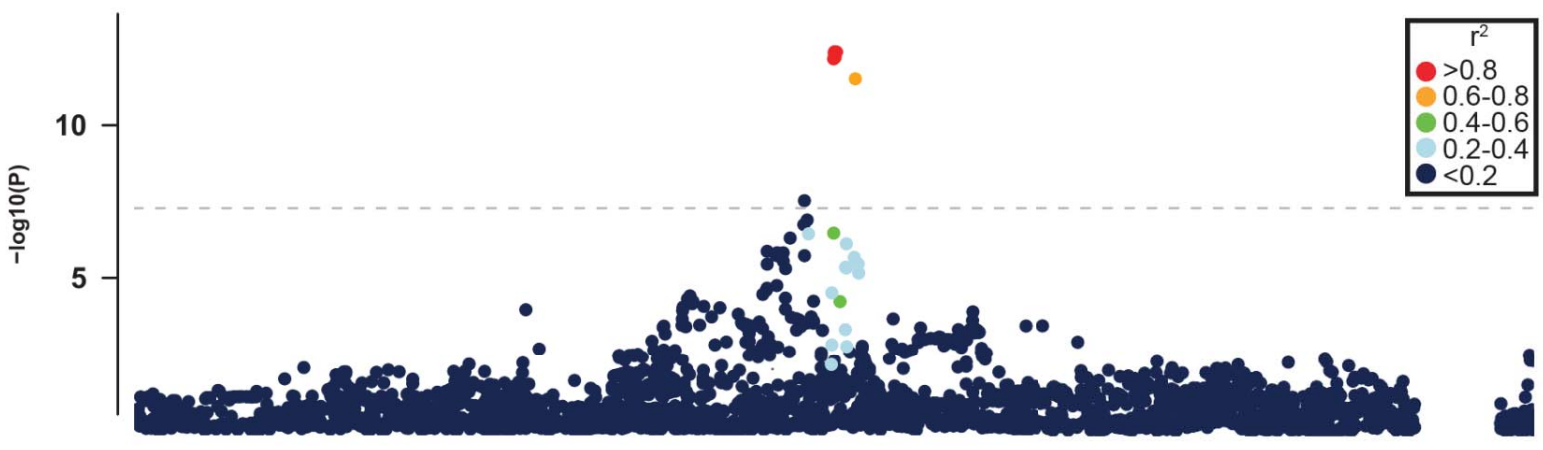
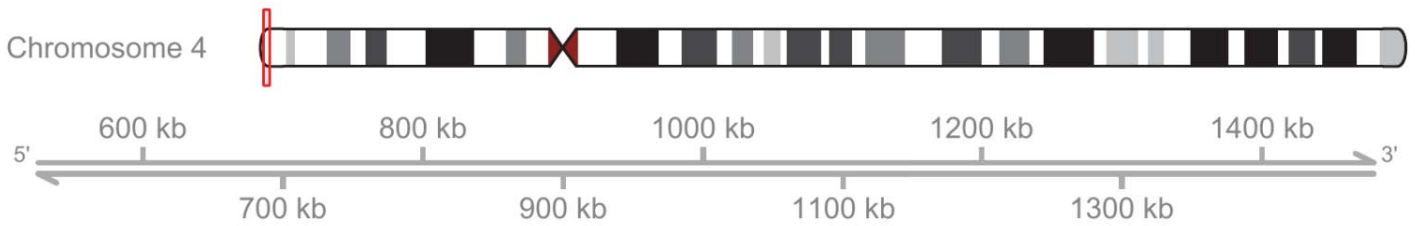
Uncorrected



Zscore of the effect allele: A



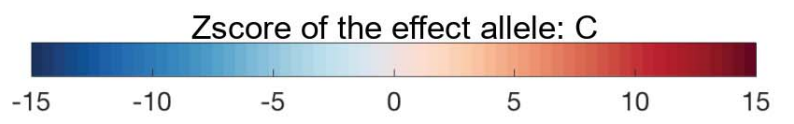
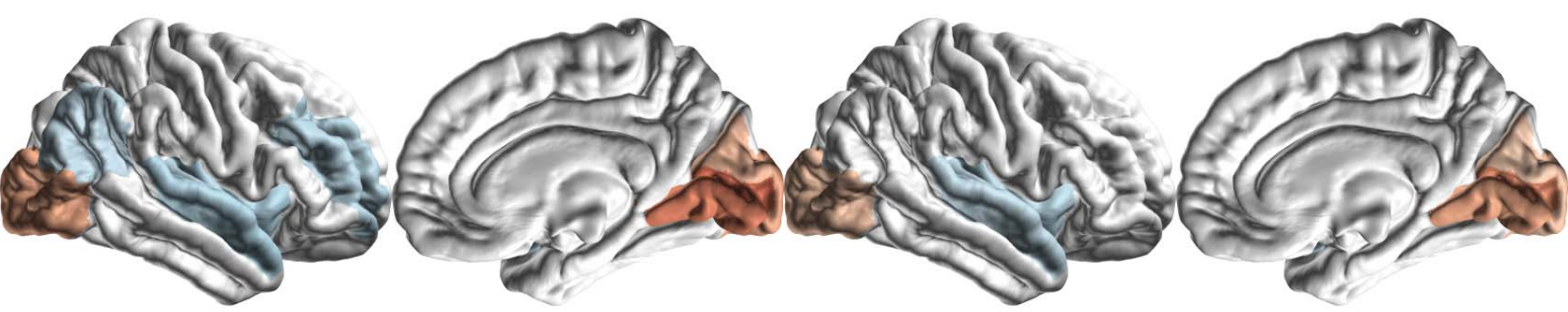
Pericalcarine SA: rs6812278



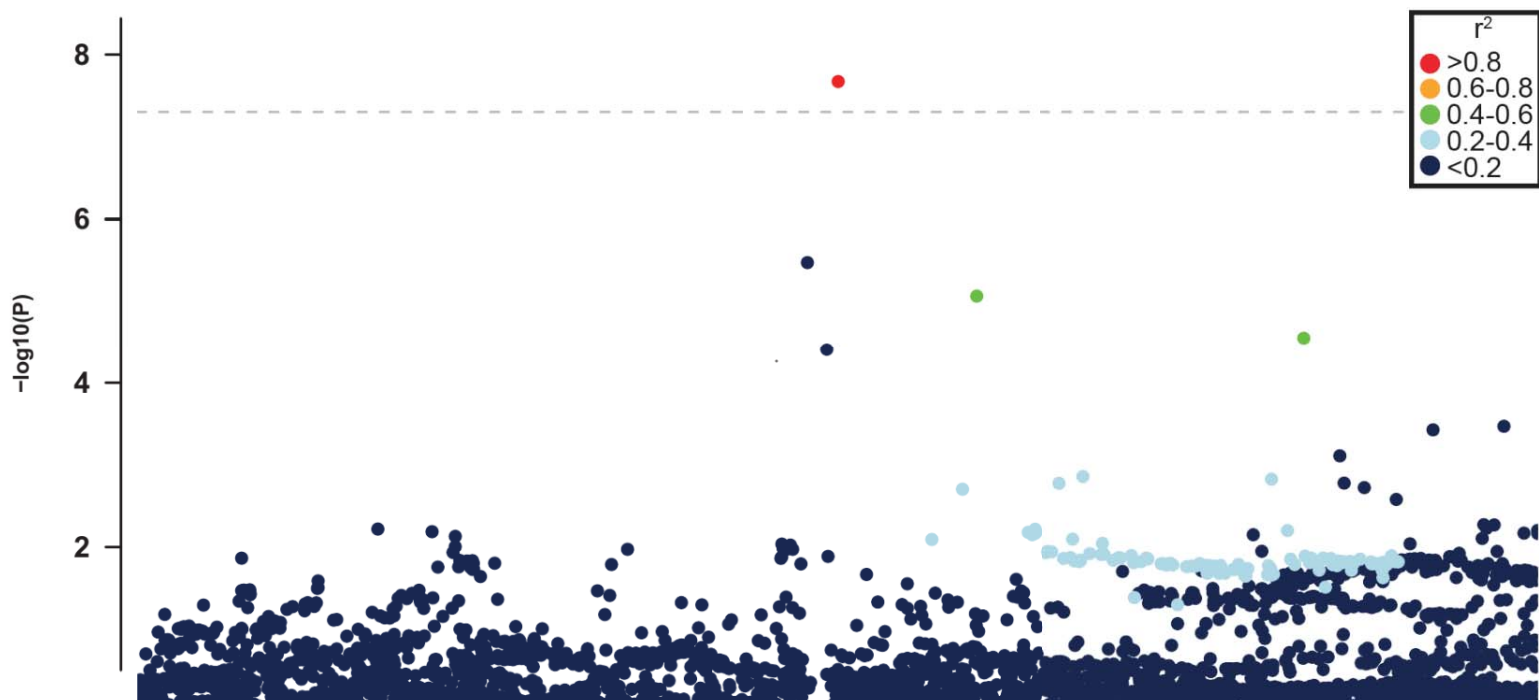
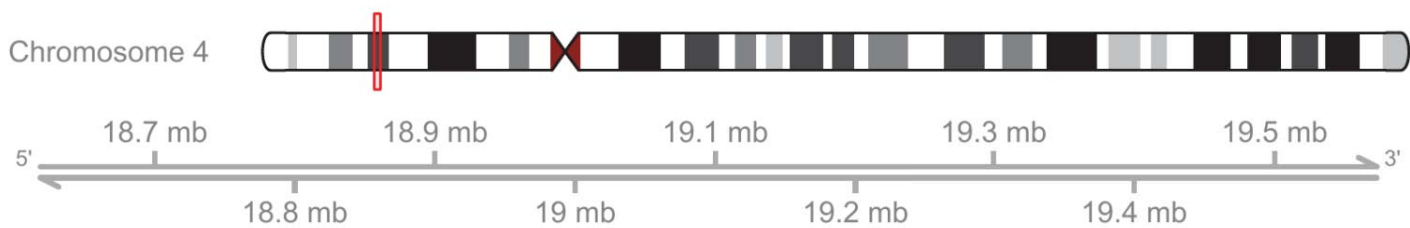
CP>GZ GZ>CP

Corrected for total SA

Uncorrected

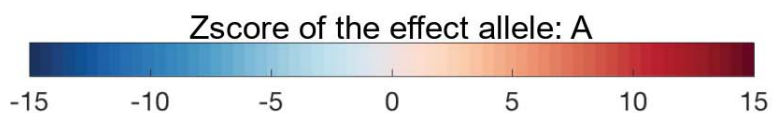
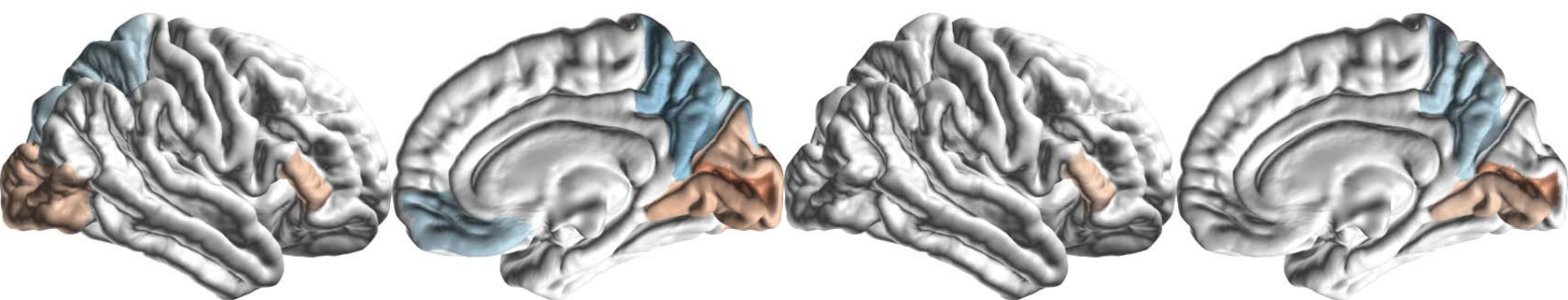


Pericalcarine SA: rs13115025

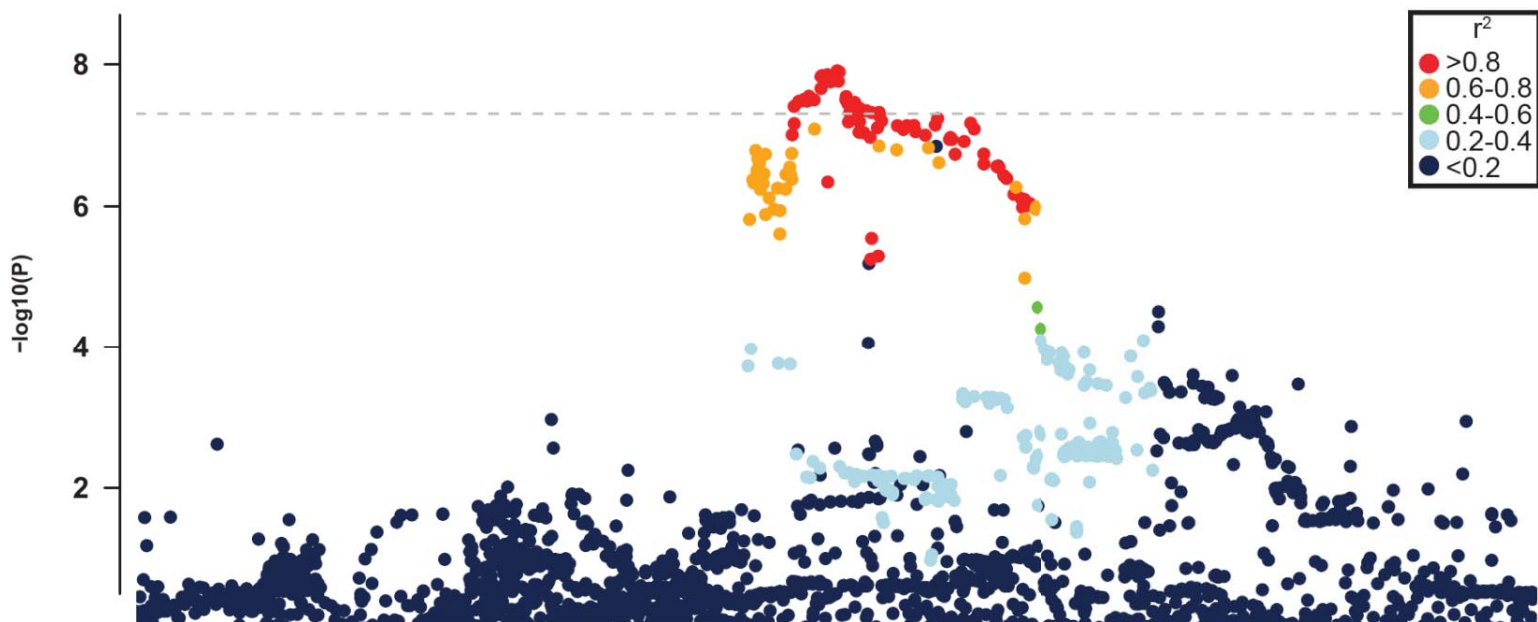
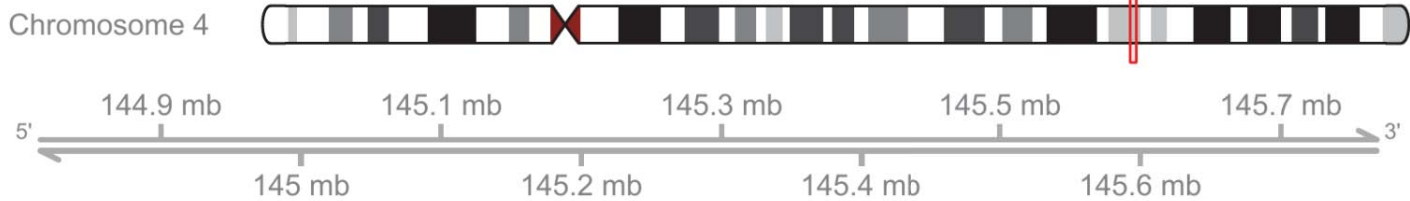


Corrected for total SA

Uncorrected

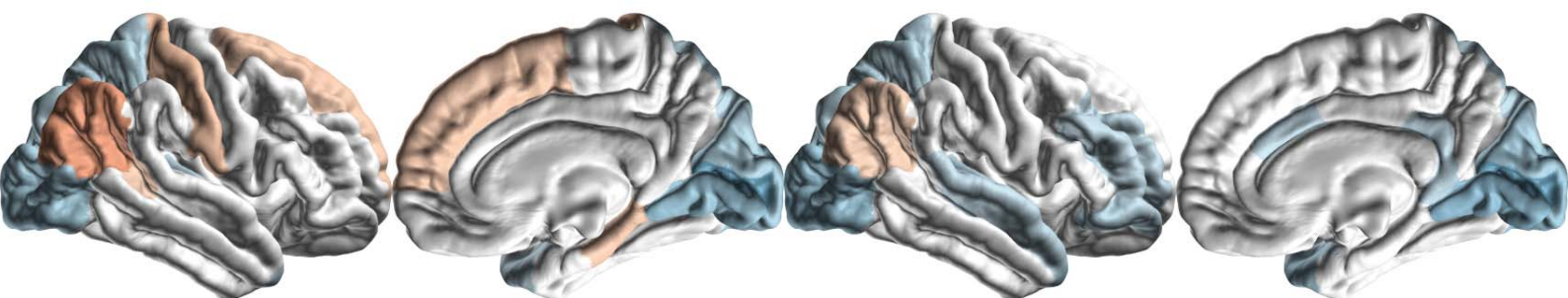


Pericalcarine SA: rs4465995



Corrected for total SA

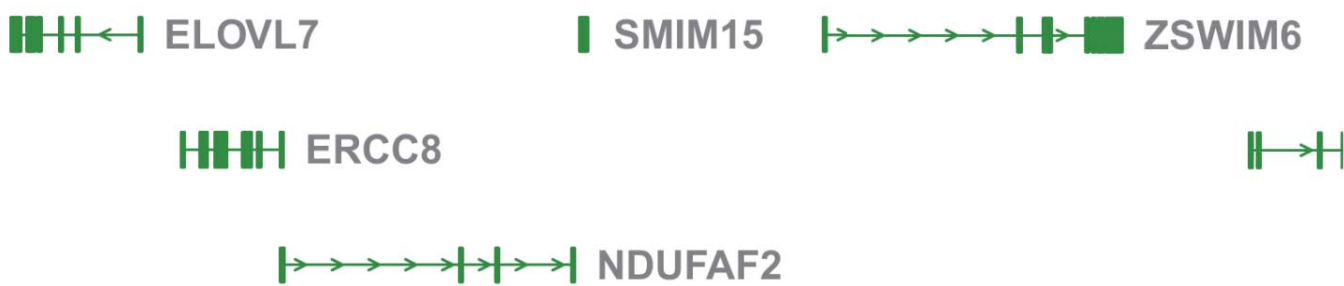
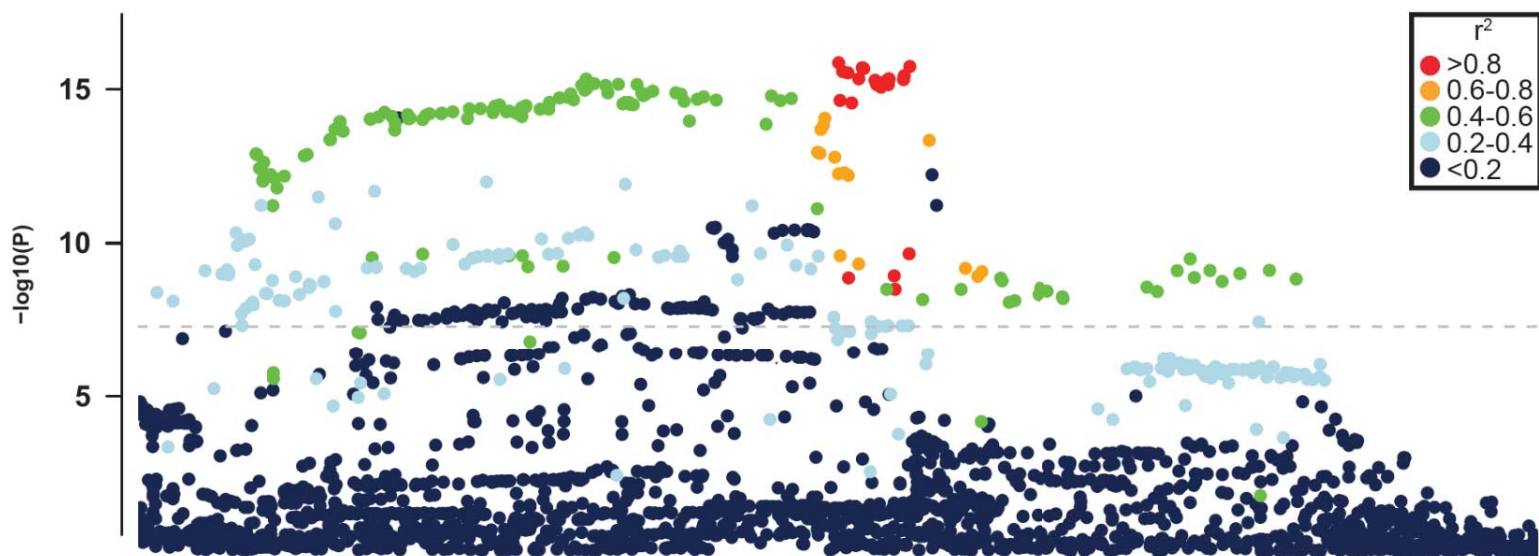
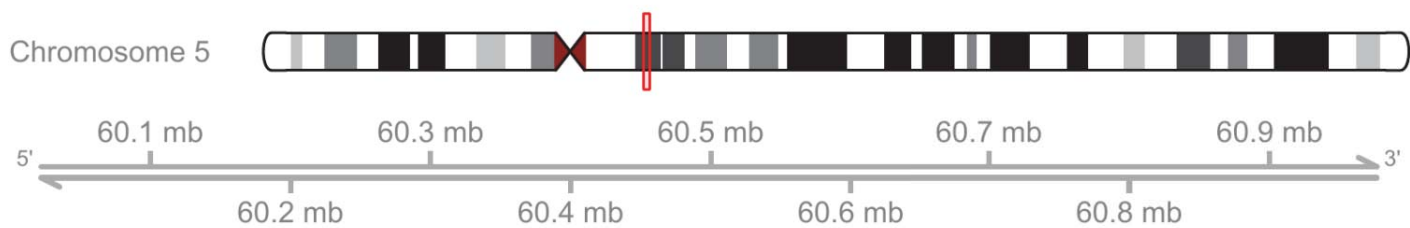
Uncorrected



Zscore of the effect allele: T

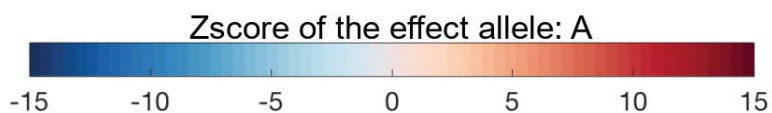
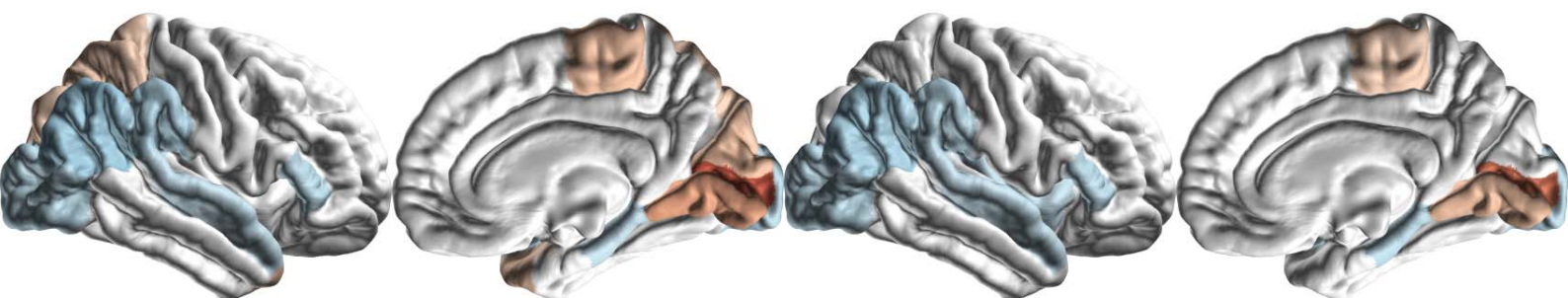


Pericalcarine SA: rs194369

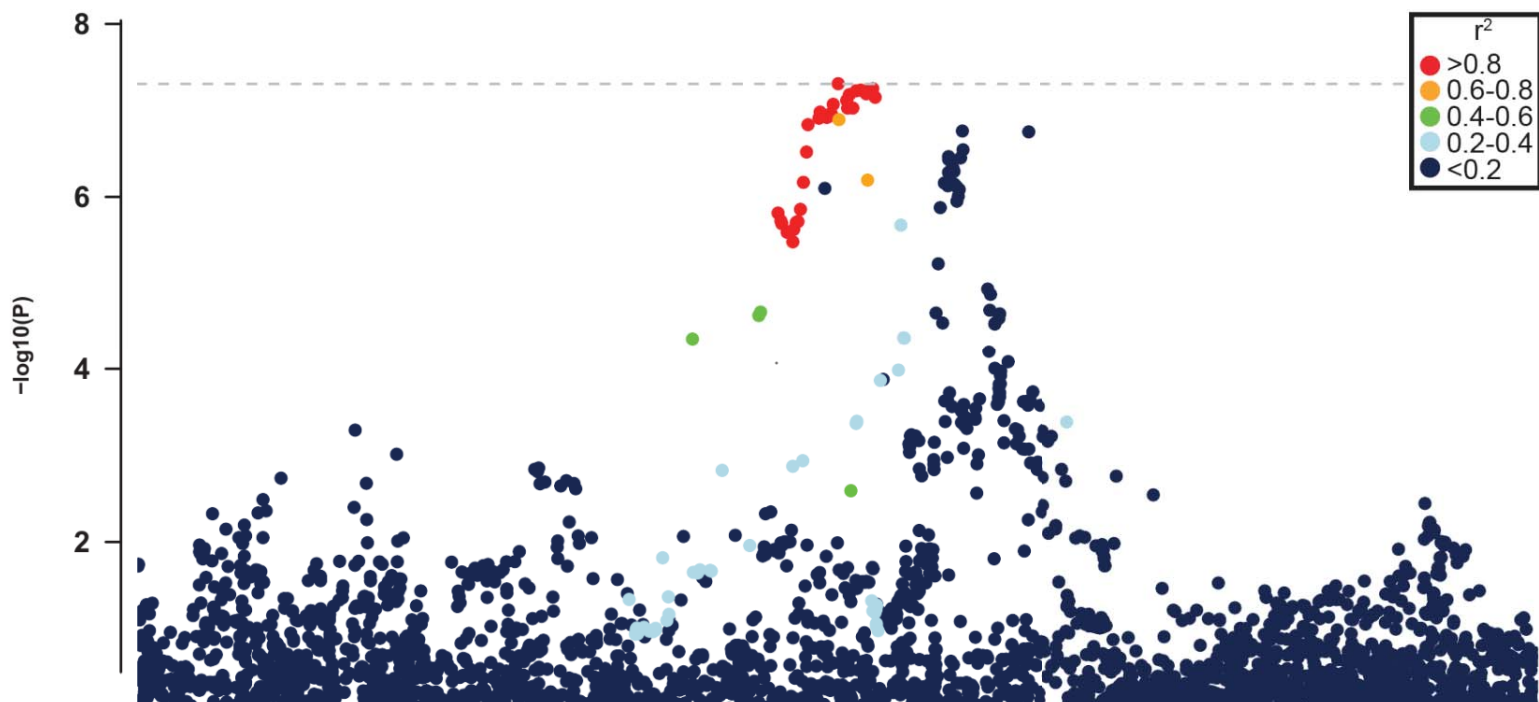


Corrected for total SA

Uncorrected



Pericalcarine SA: rs57063427



ZNF608

GZ>CP

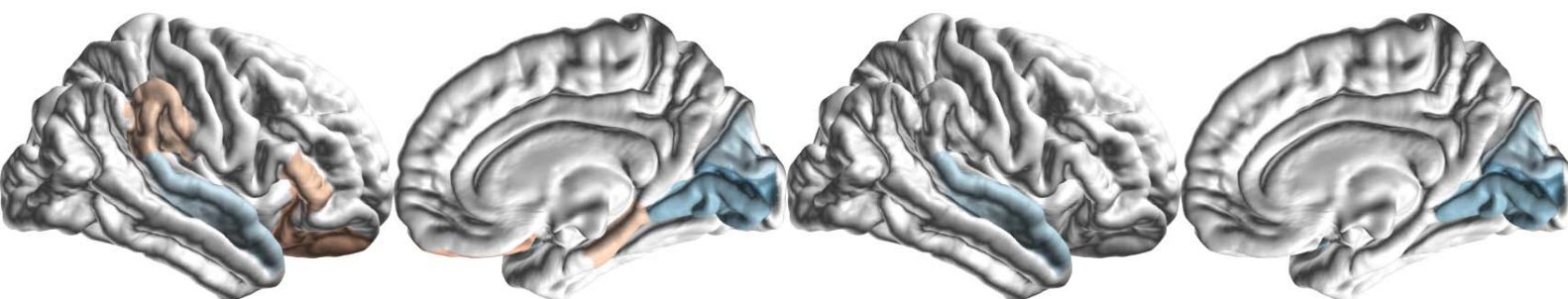


CP>GZ



Corrected for total SA

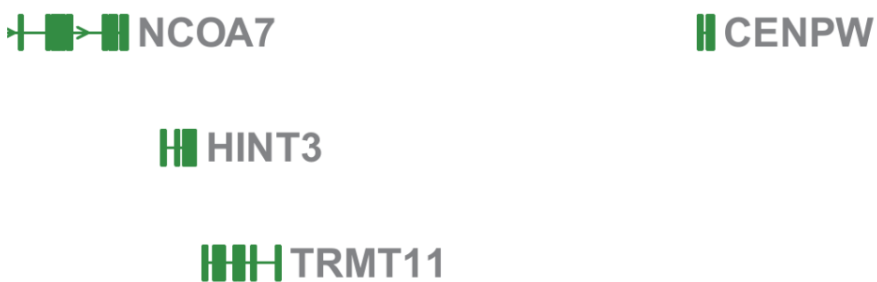
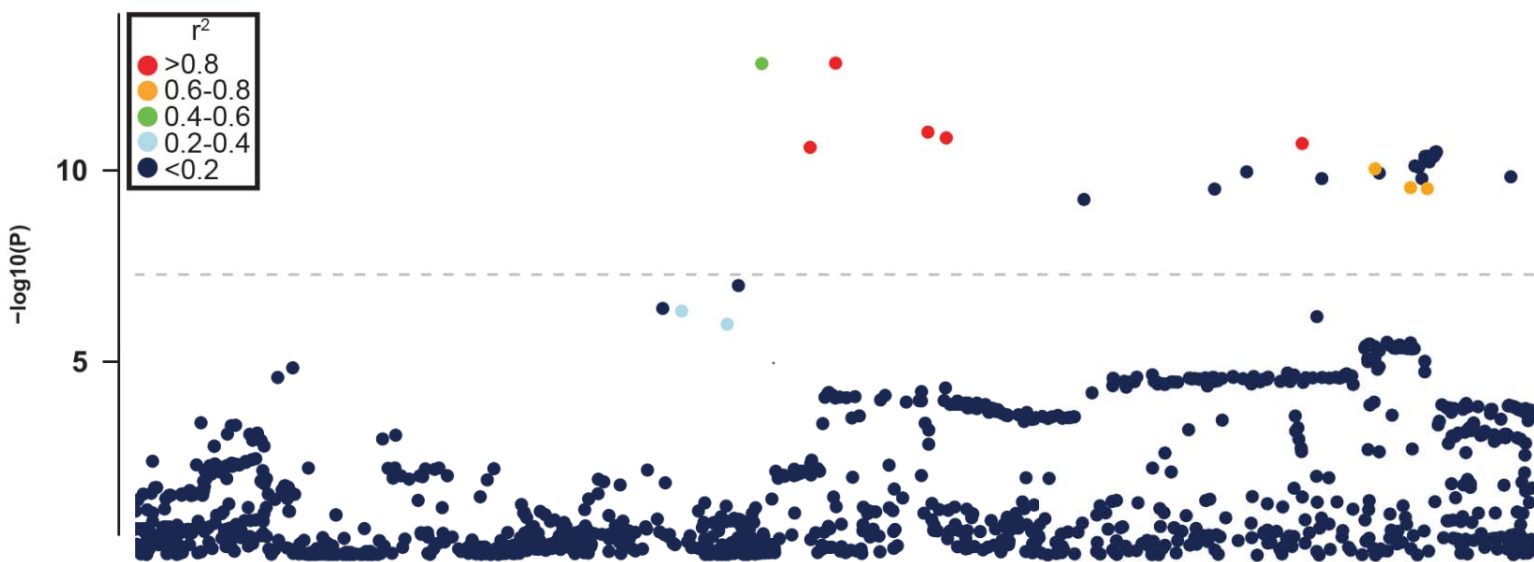
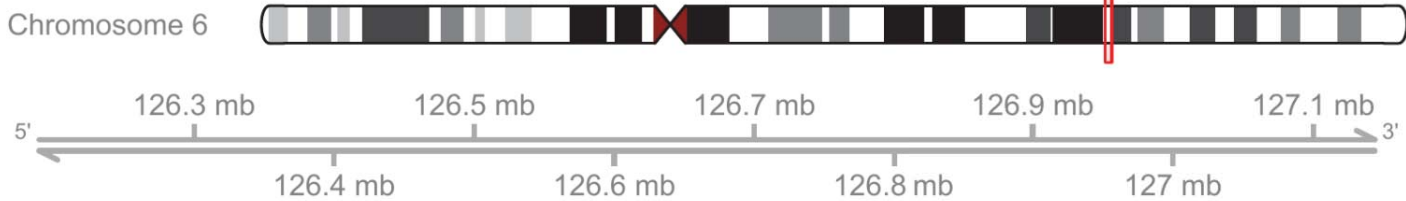
Uncorrected



Zscore of the effect allele: T

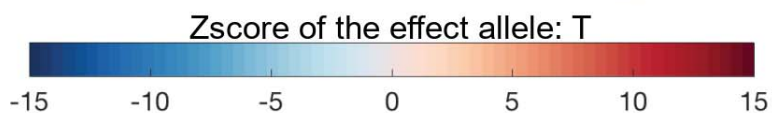
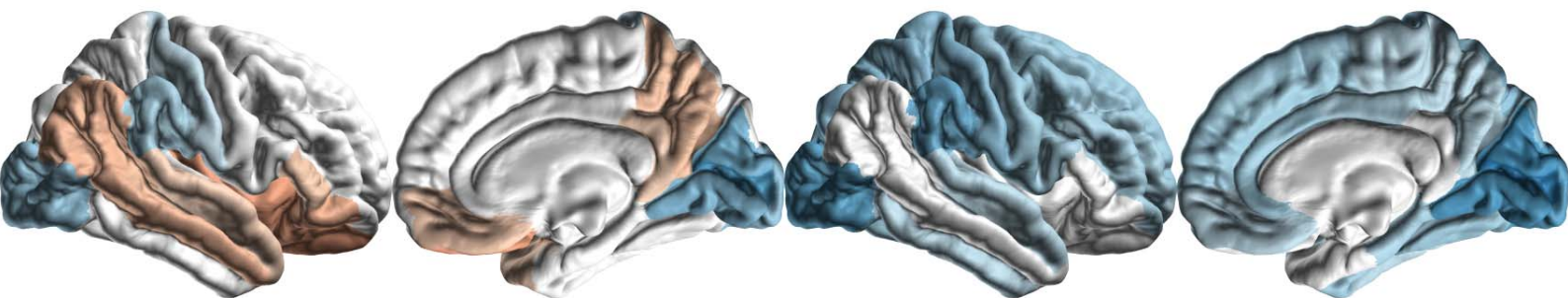


Pericalcarine SA: rs117892760

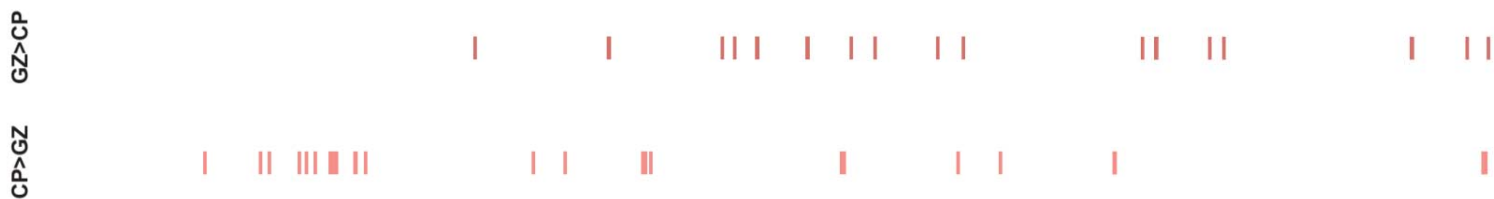
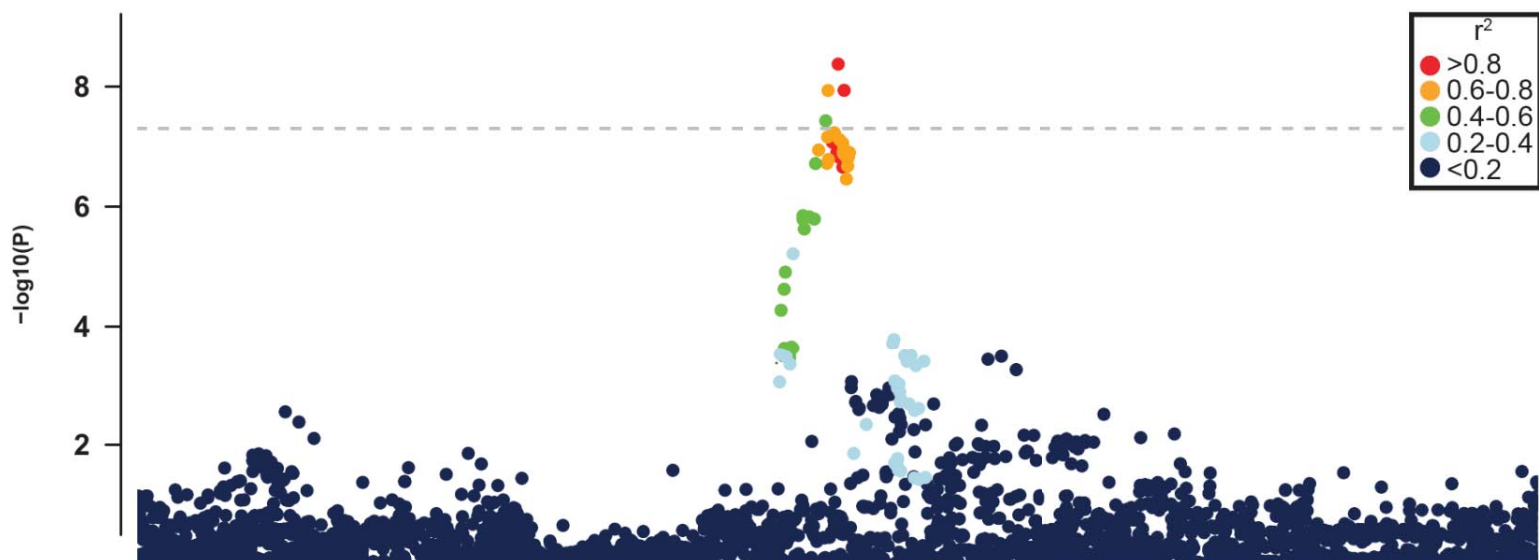
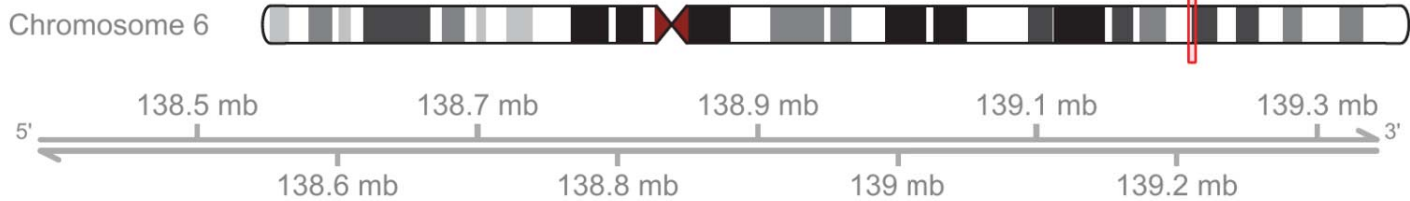


Corrected for total SA

Uncorrected

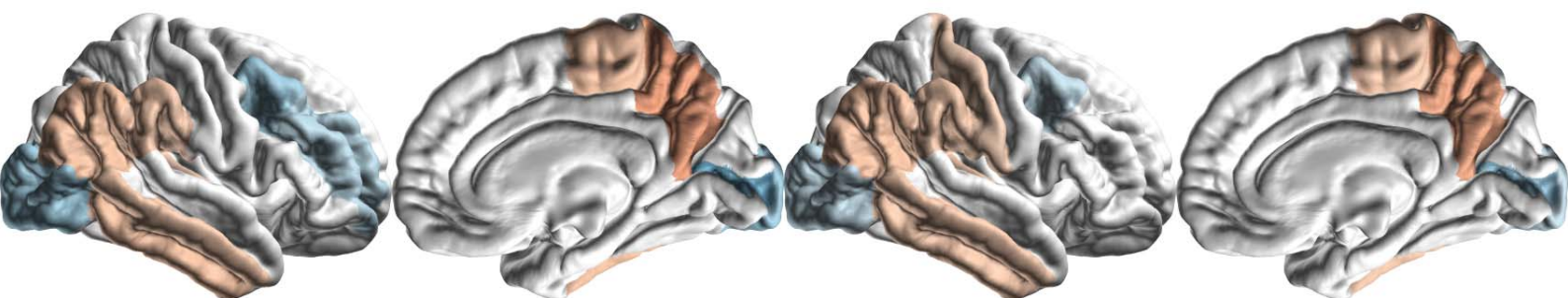


Pericalcarine SA: rs489532



Corrected for total SA

Uncorrected

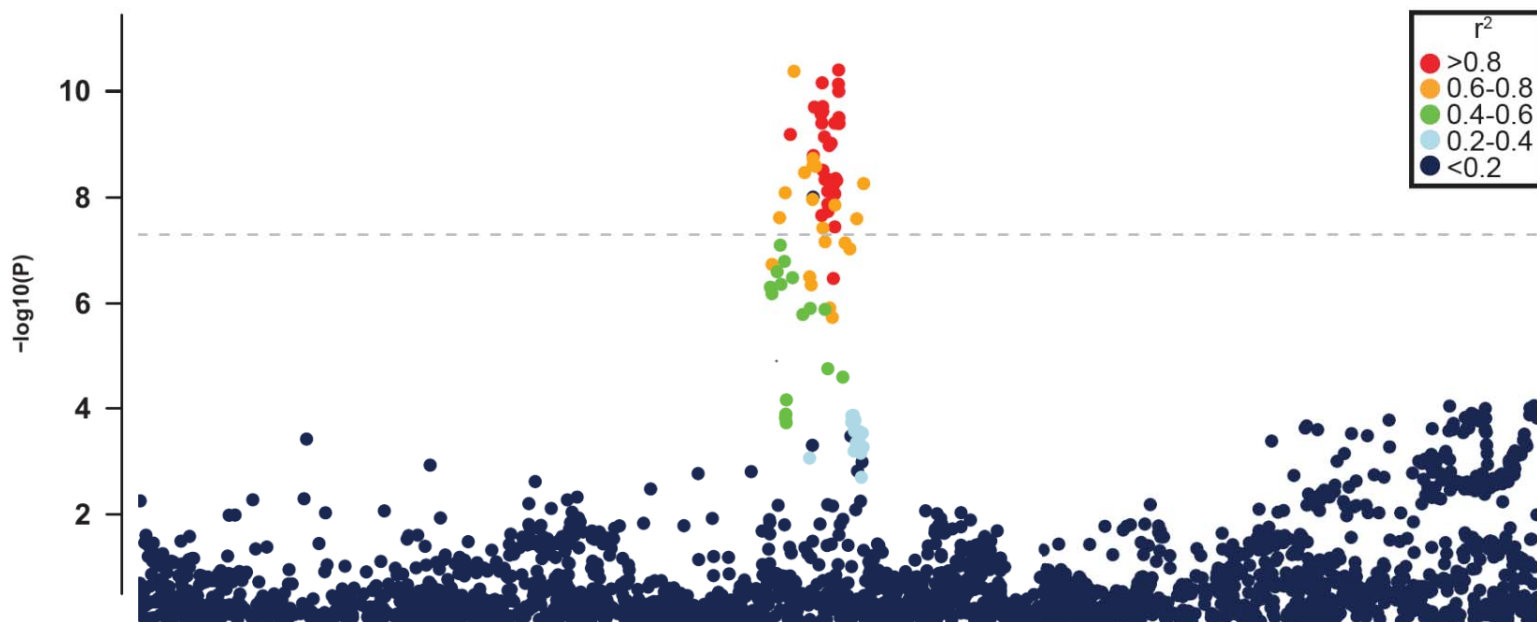
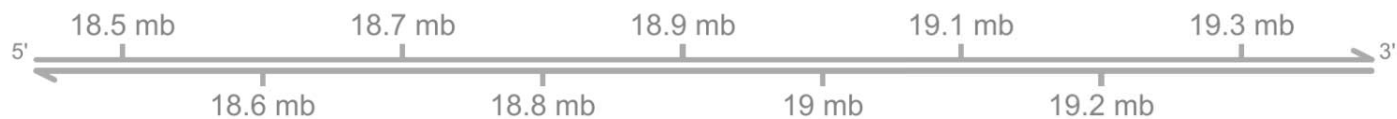


Zscore of the effect allele: T



Pericalcarine SA: rs10237366

Chromosome 7



TWIST1

GZ>CP

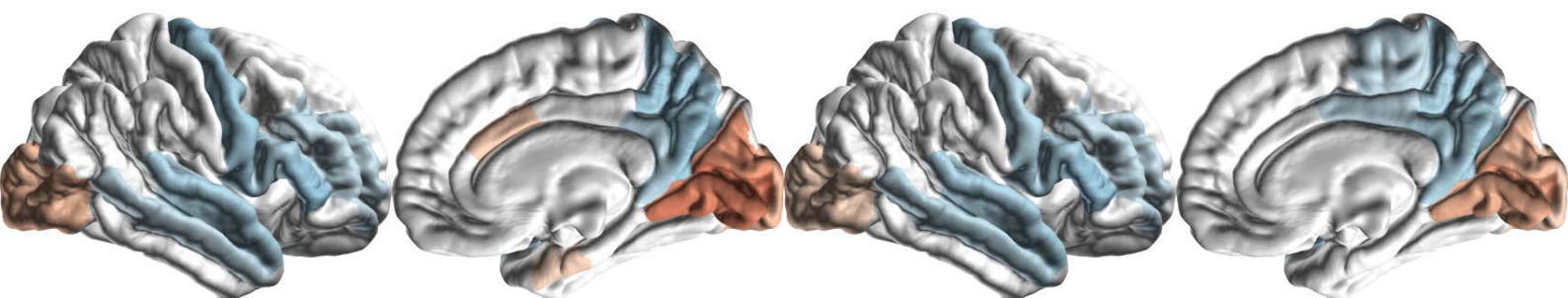


CP>GZ

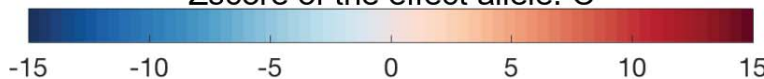


Corrected for total SA

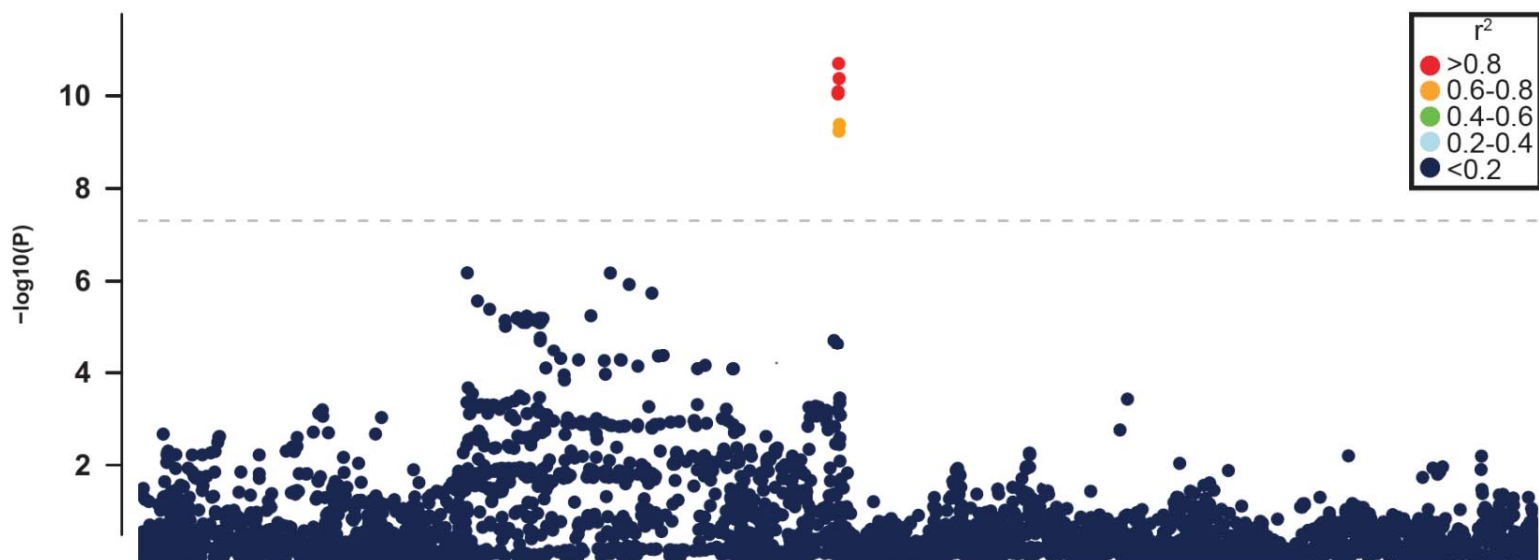
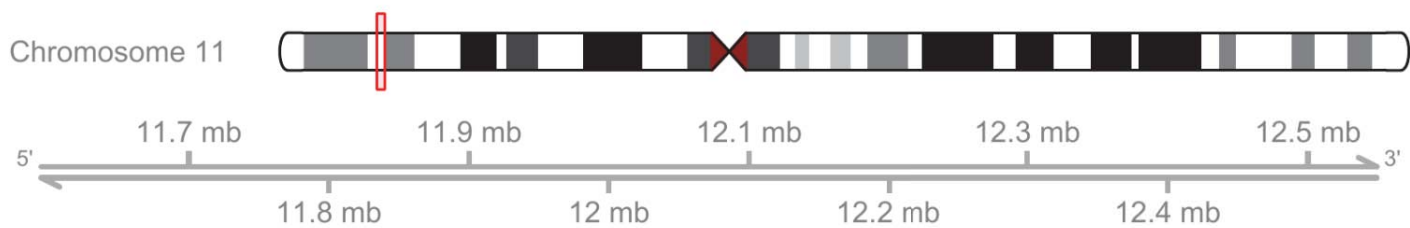
Uncorrected



Zscore of the effect allele: C

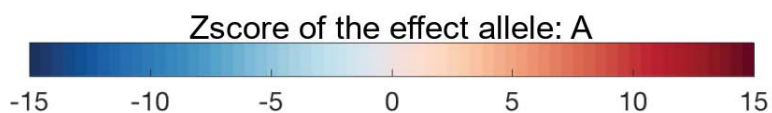
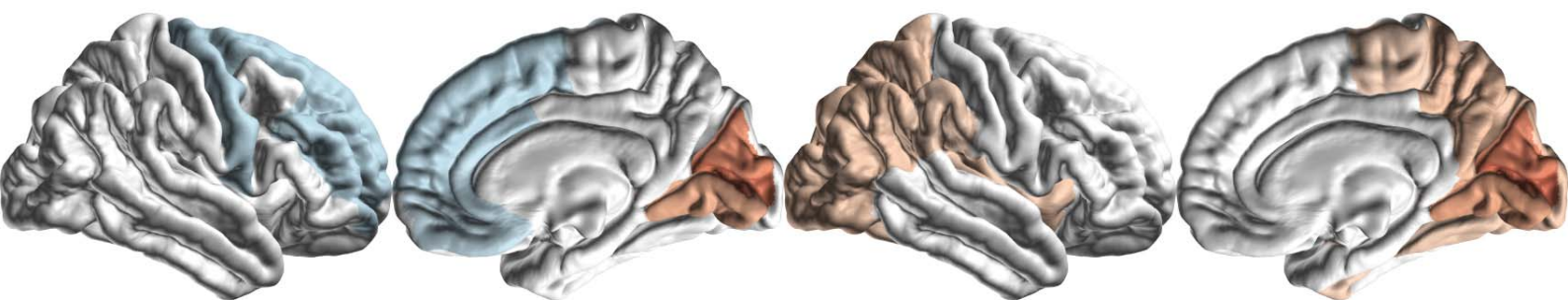


Pericalcarine SA: rs10765918

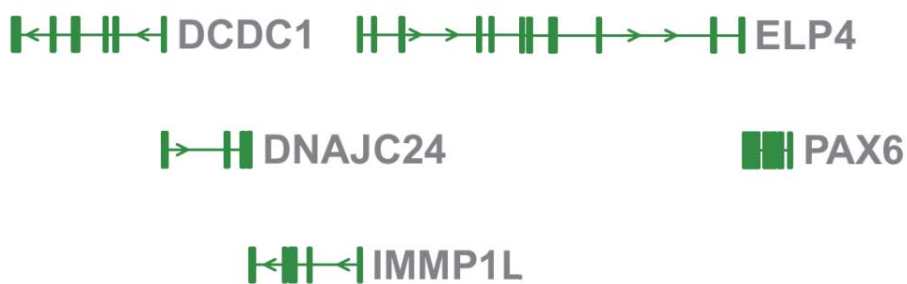
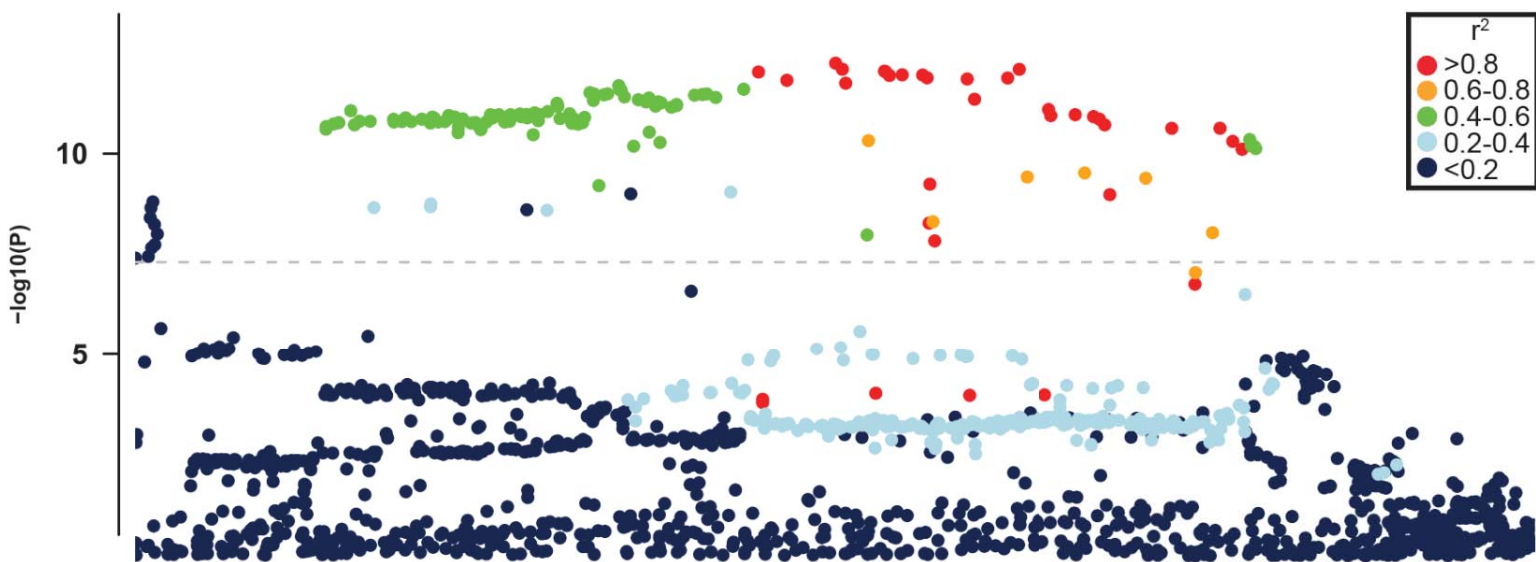
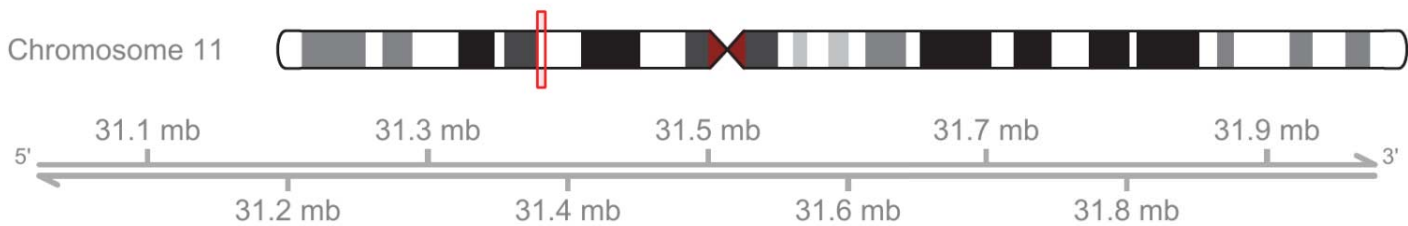


Corrected for total SA

Uncorrected

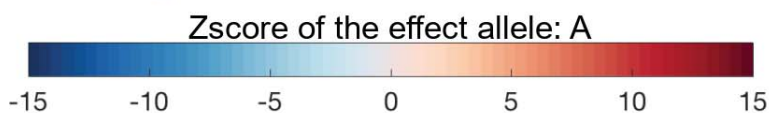
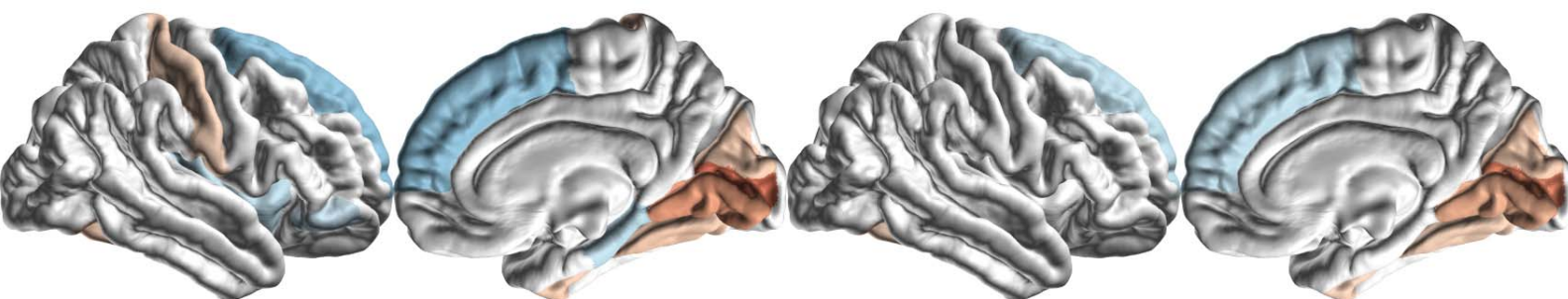


Pericalcarine SA: rs1223090

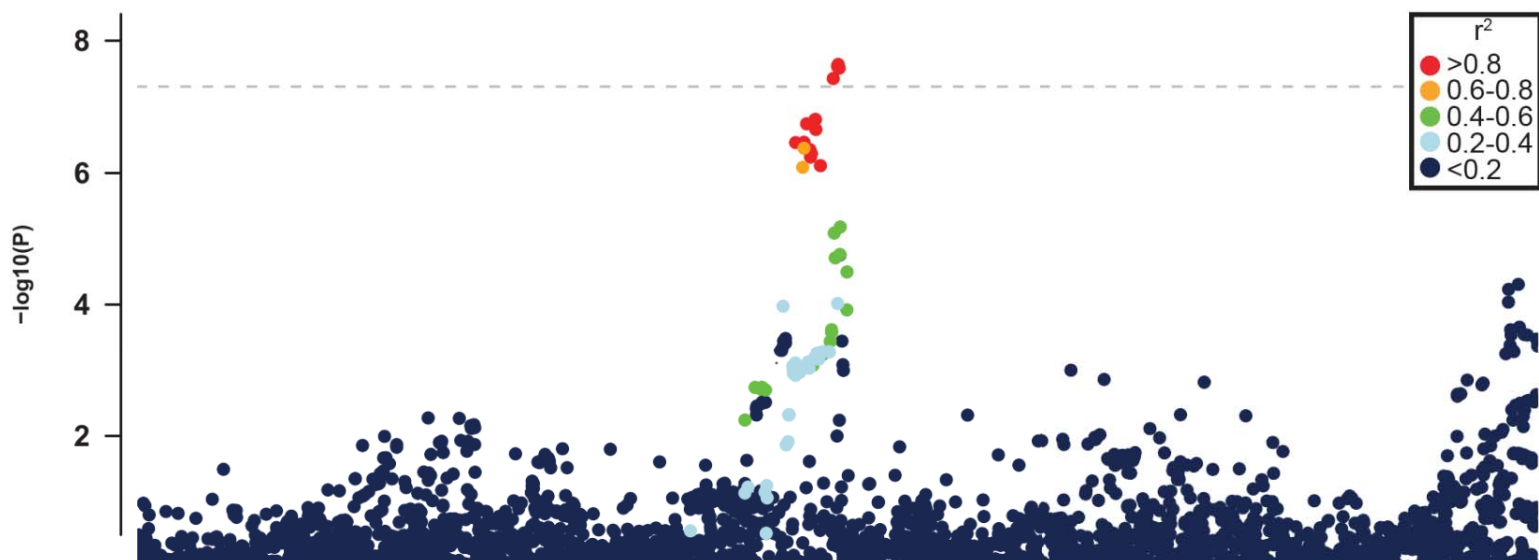
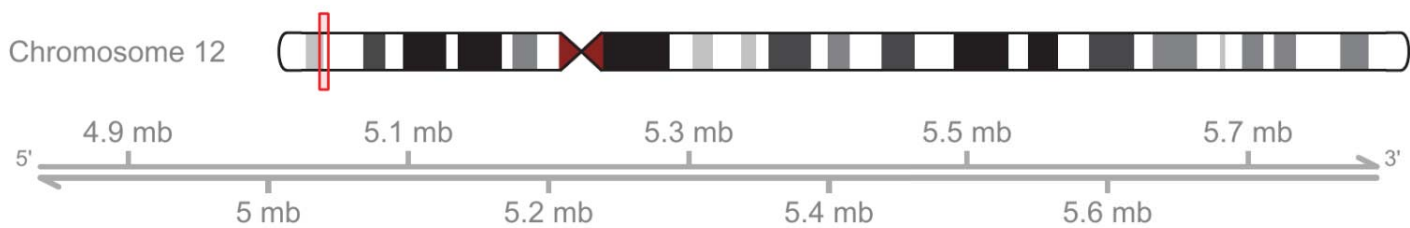


Corrected for total SA

Uncorrected



Pericalcarine SA: rs17179798



GALNT8

KCNA5

NTF3

KCNA6

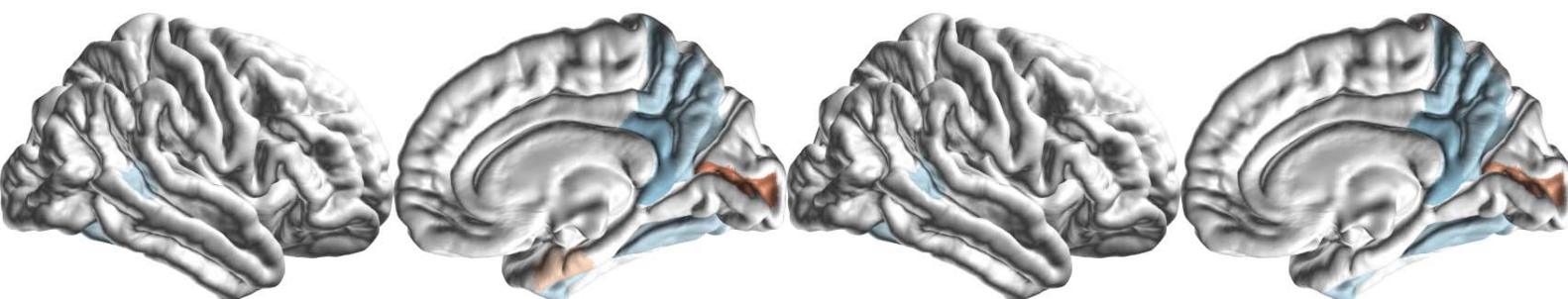
KCNA1

GZ>CP

CP>GZ

Corrected for total SA

Uncorrected

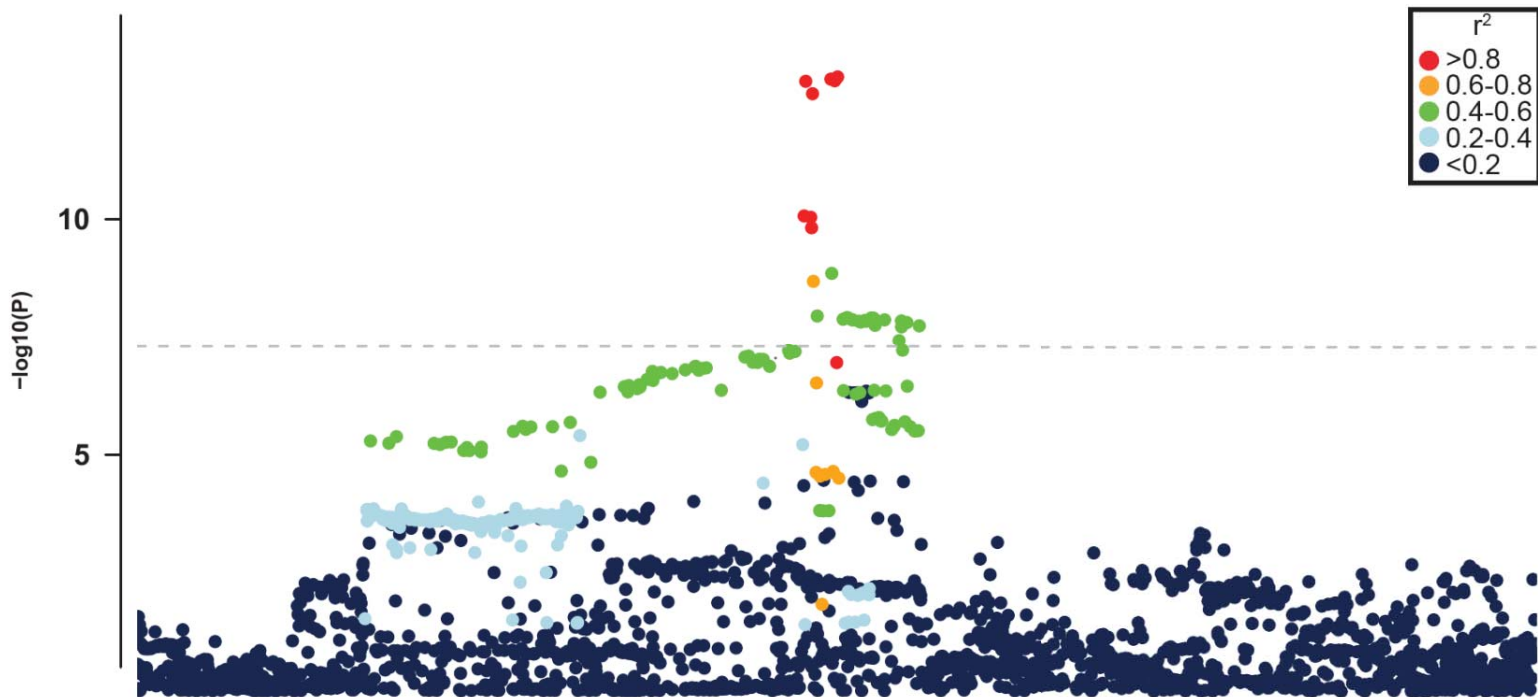


Zscore of the effect allele: A



Pericalcarine SA: rs9545158

Chromosome 13



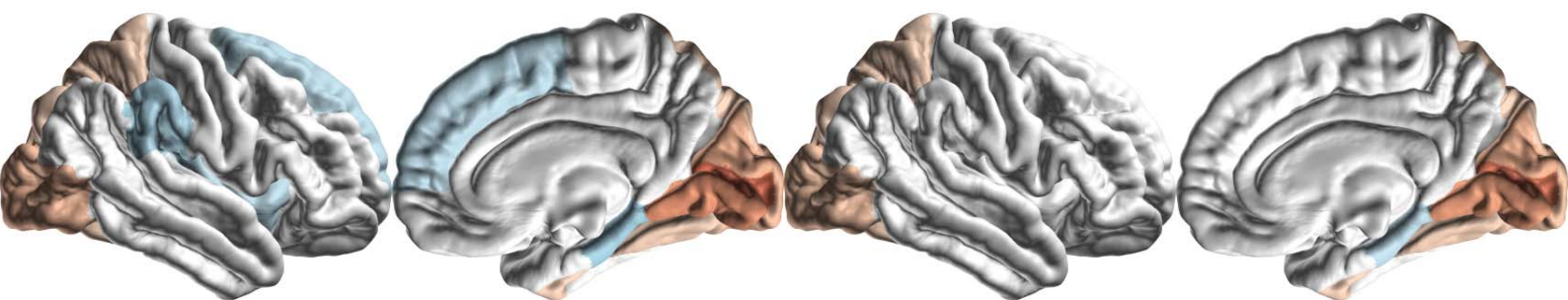
RBM26 NDFIP2

GZ>CP

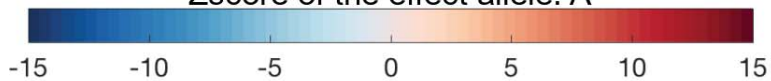
CP>GZ

Corrected for total SA

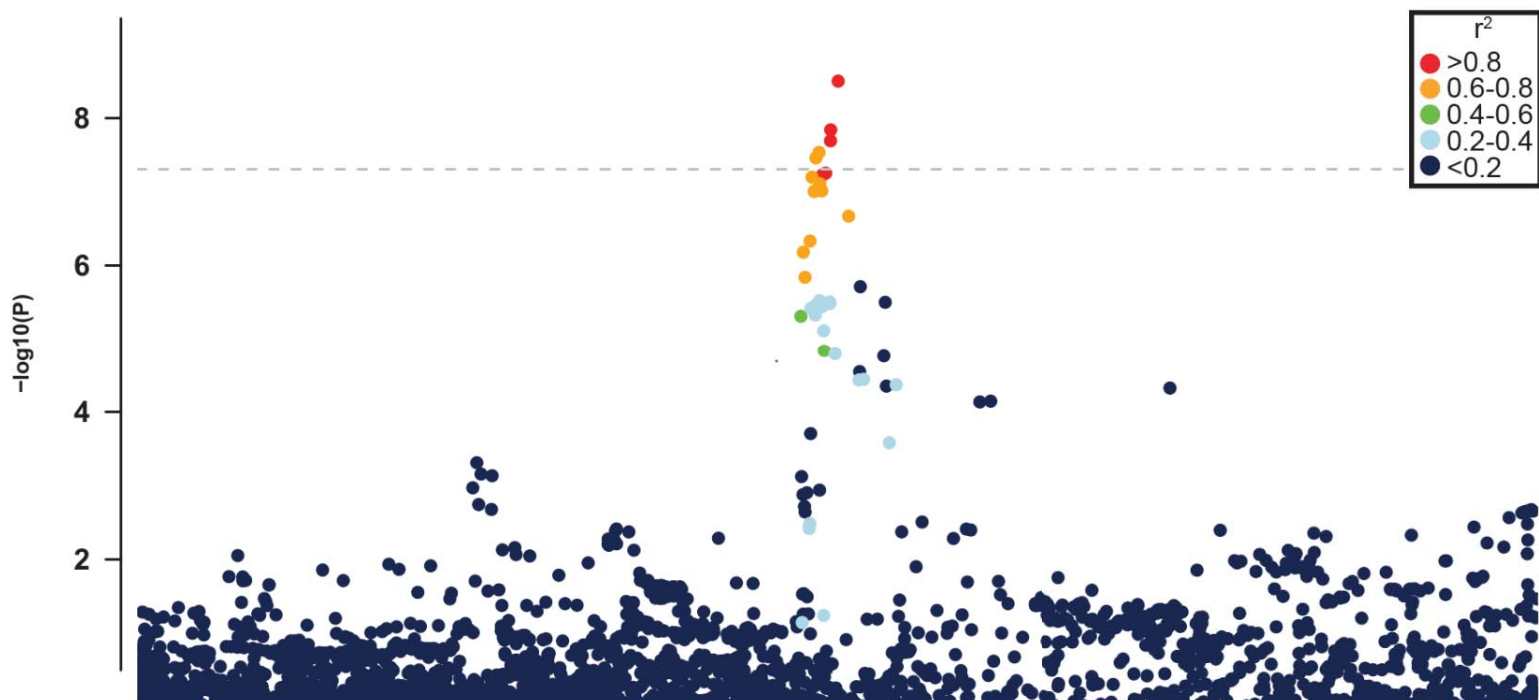
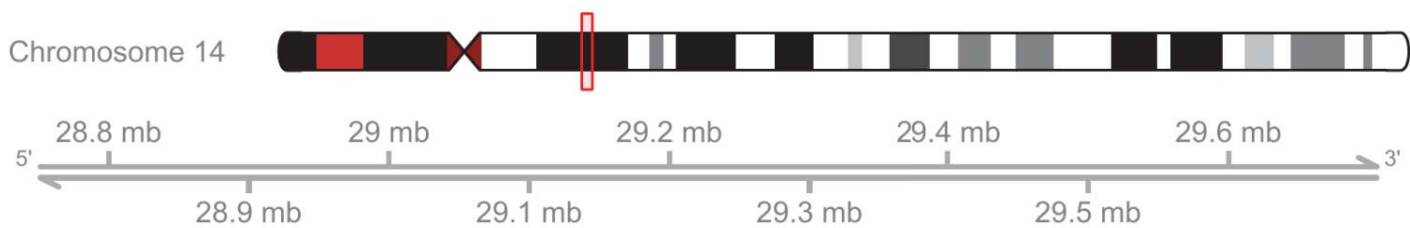
Uncorrected



Zscore of the effect allele: A



Pericalcarine SA: rs35342371



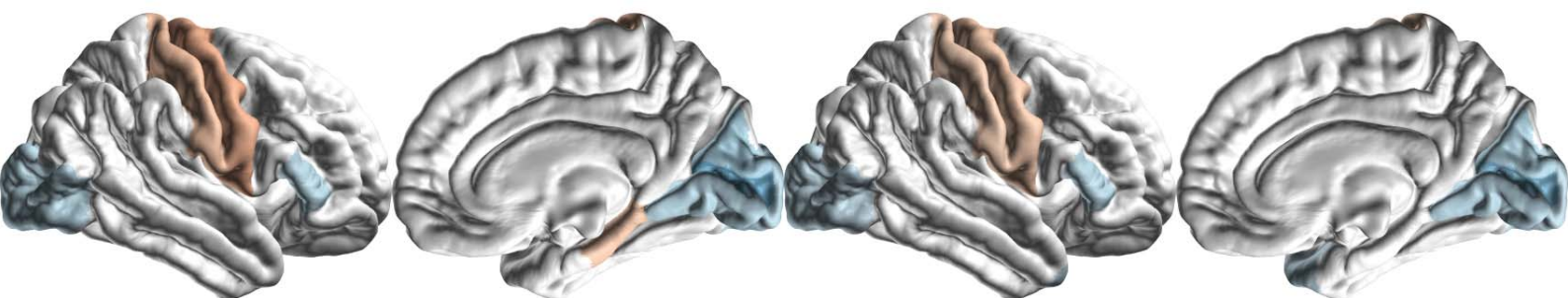
FOXG1

GZ>CP

CP>GZ

Corrected for total SA

Uncorrected

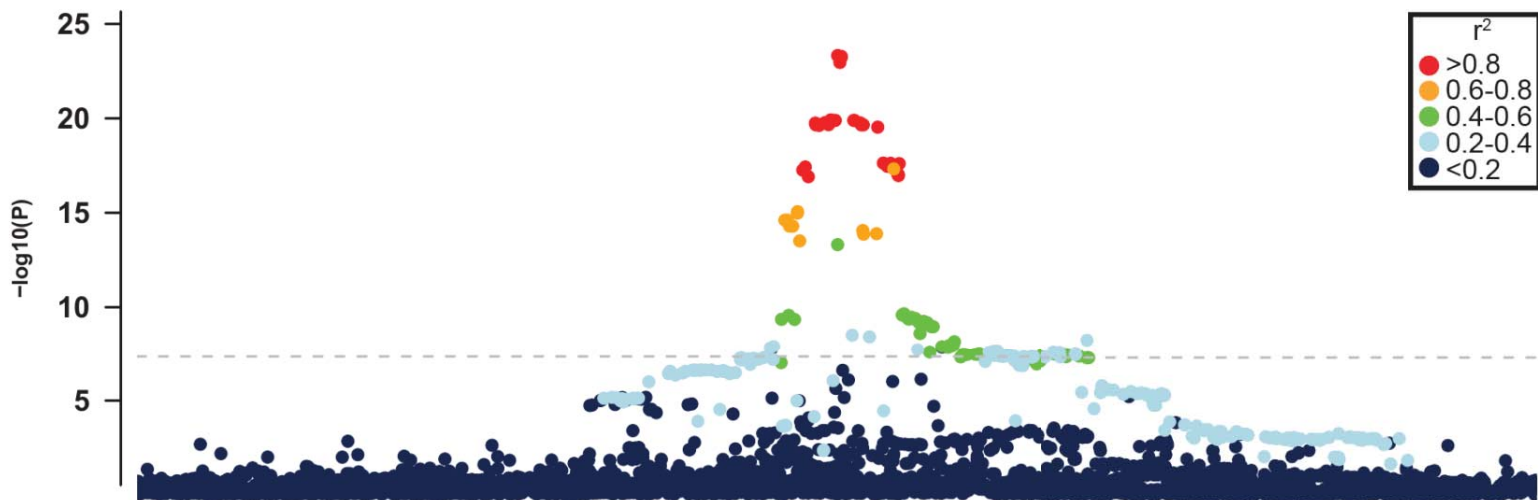
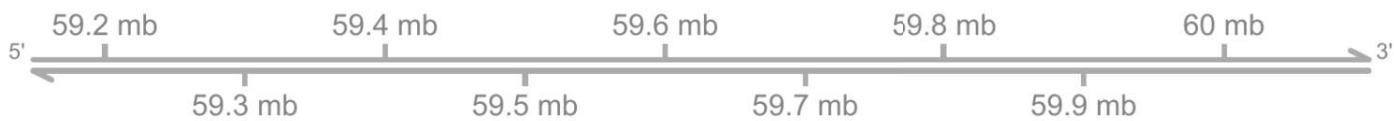


Zscore of the effect allele: A



Pericalcarine SA: rs73313052

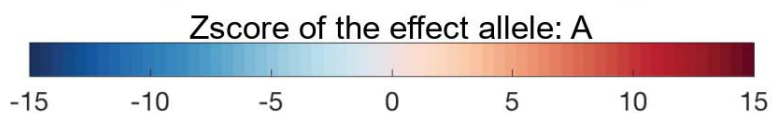
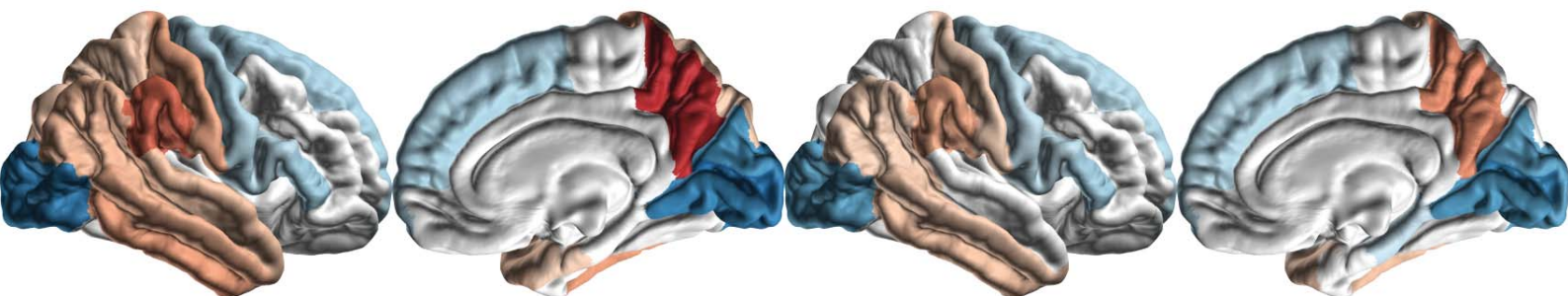
Chromosome 14



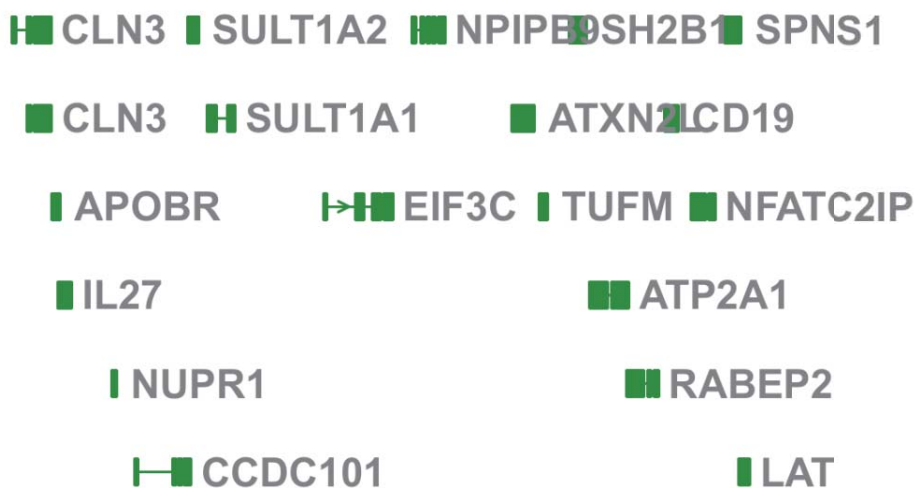
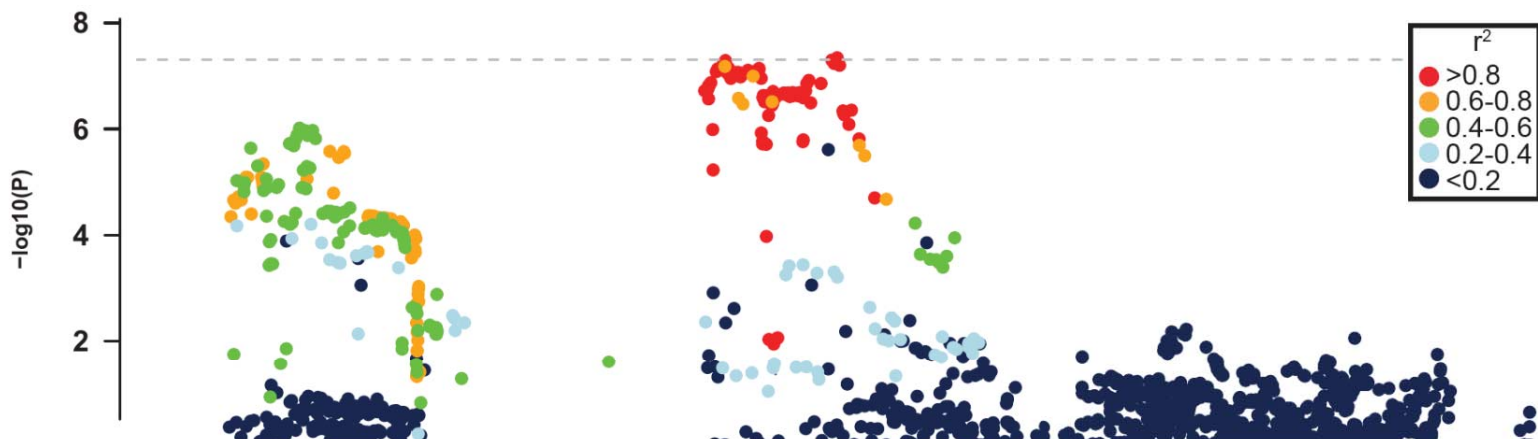
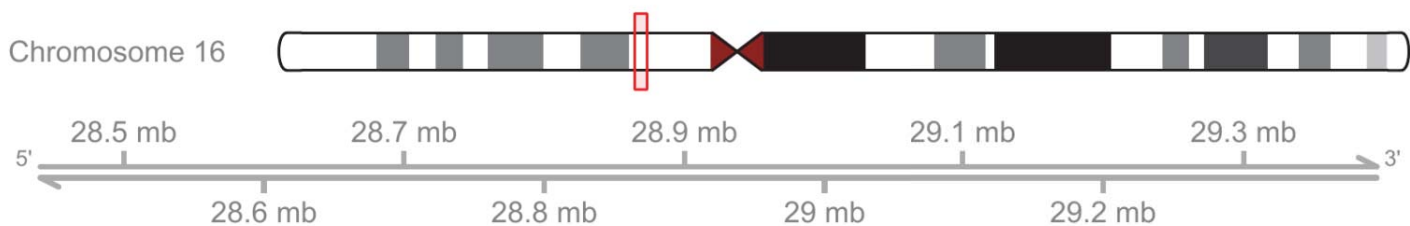
CP>GZ
GZ>CP

Corrected for total SA

Uncorrected



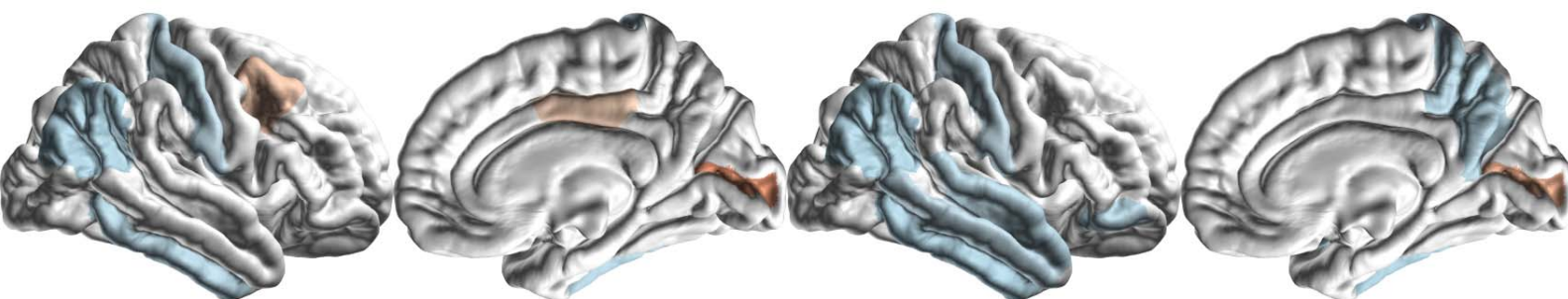
Pericalcarine SA: rs7188071



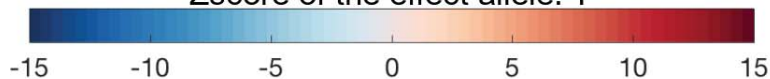
CP>GZ GZ>CP

Corrected for total SA

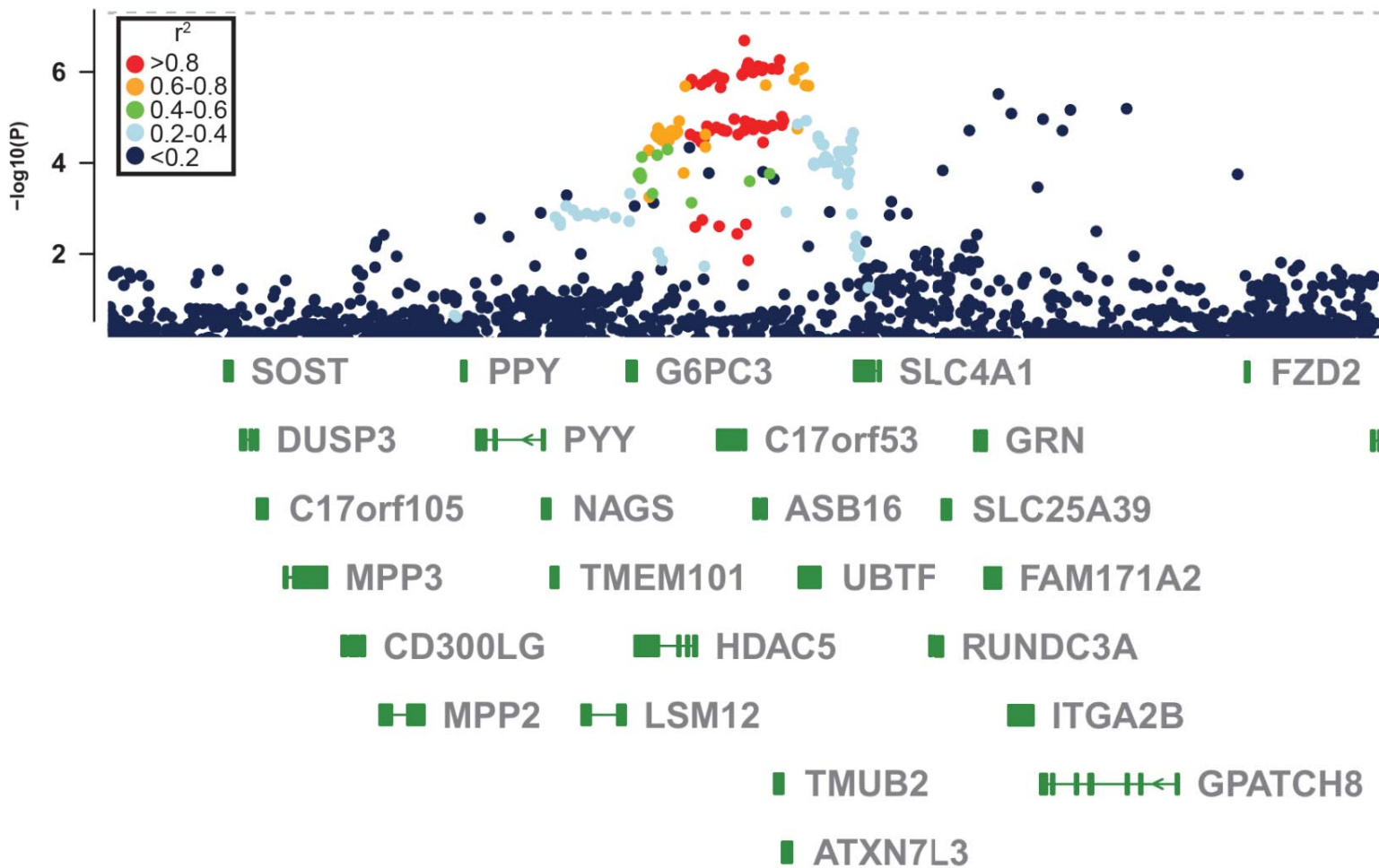
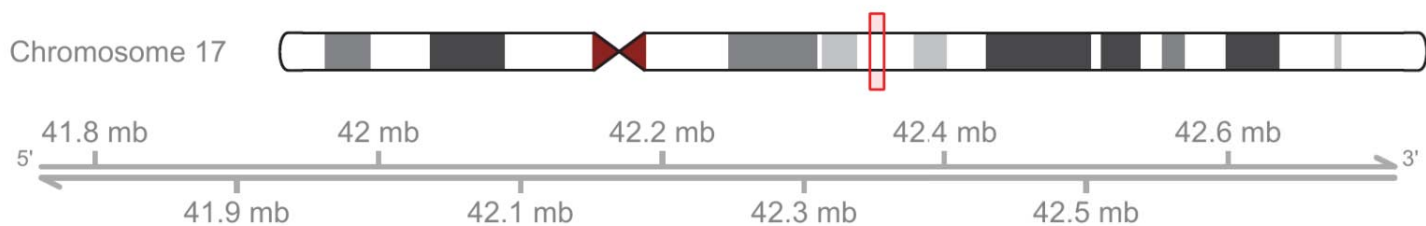
Uncorrected



Zscore of the effect allele: T



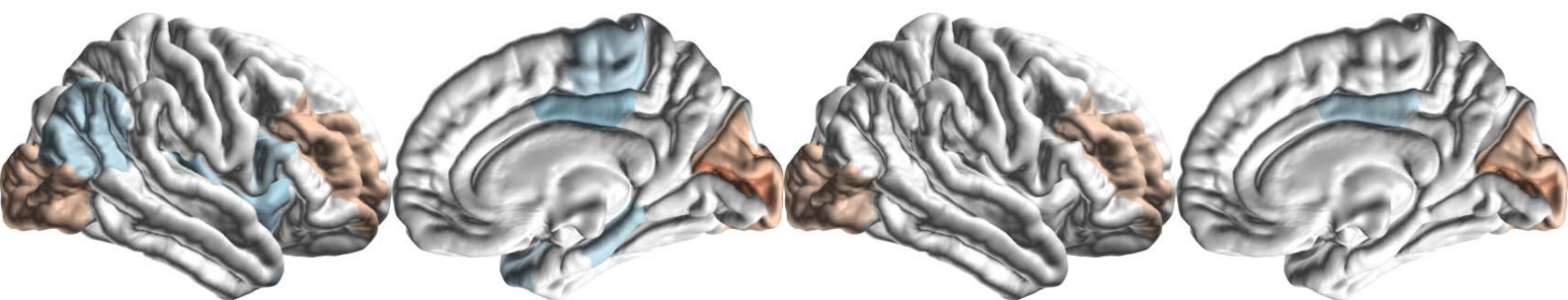
Pericalcarine SA: rs13726



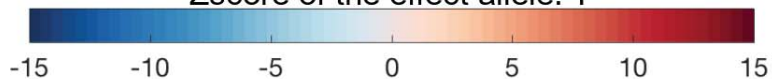
CP>GZGZ>CP

Corrected for total SA

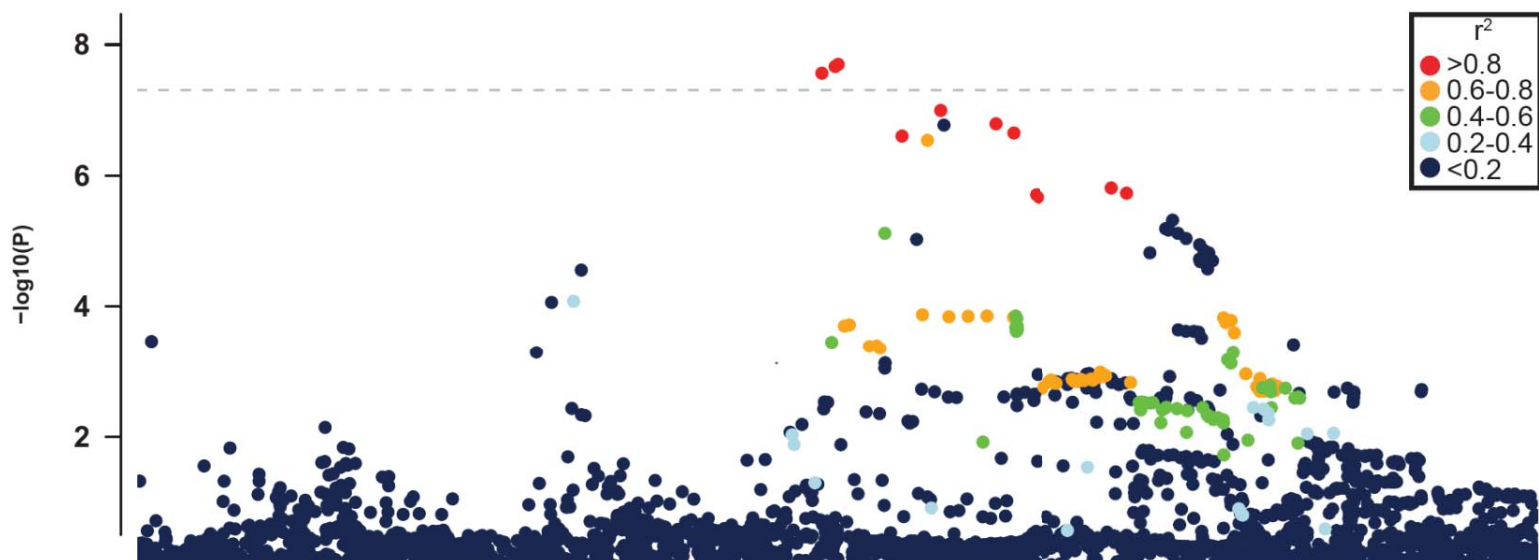
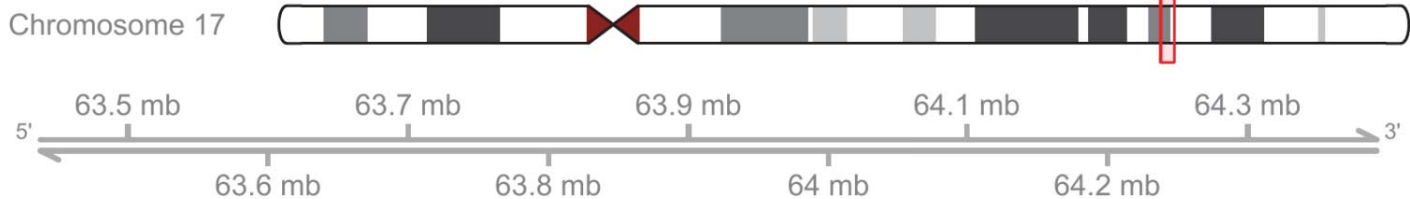
Uncorrected



Zscore of the effect allele: T



Pericalcarine SA: rs1420791



AXIN2

APOH

CEP112

GZ>CP

||

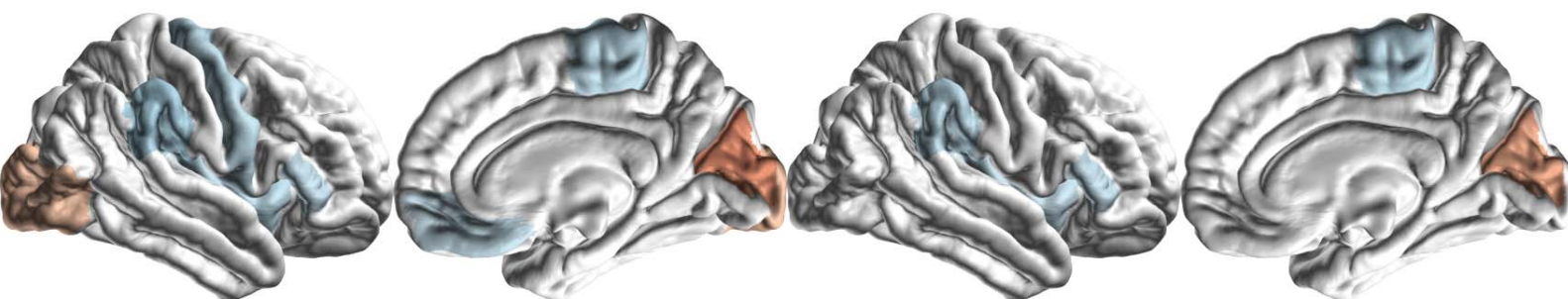
||

CP>GZ

|

Corrected for total SA

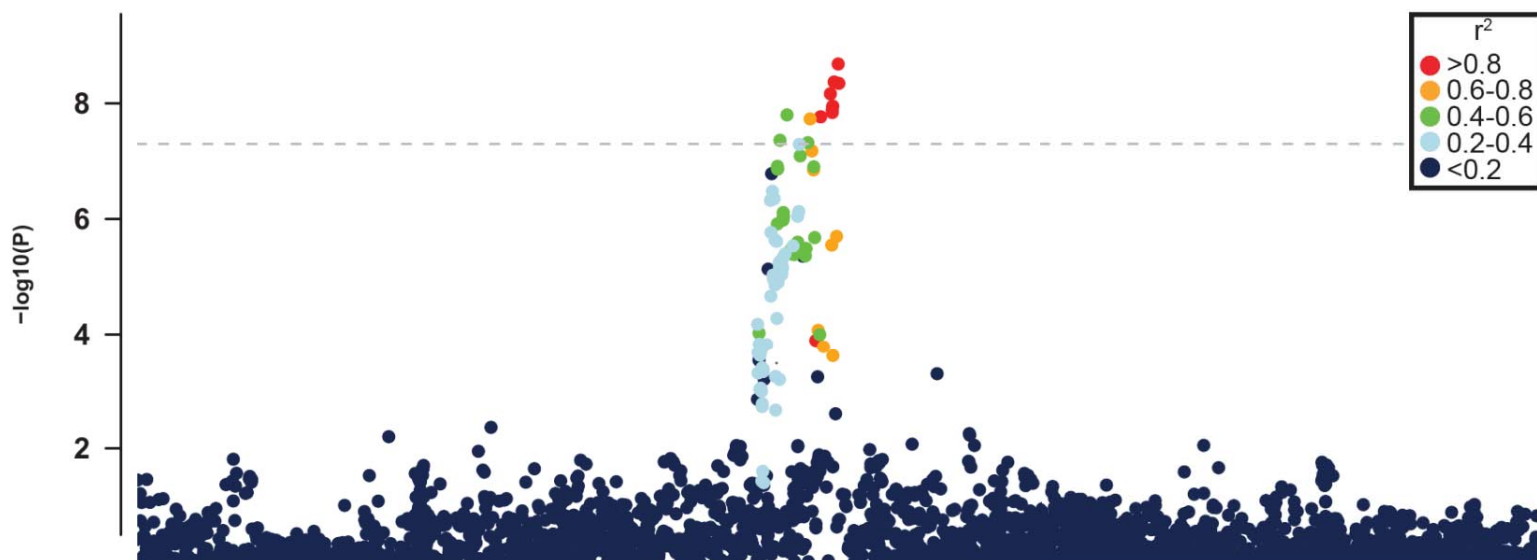
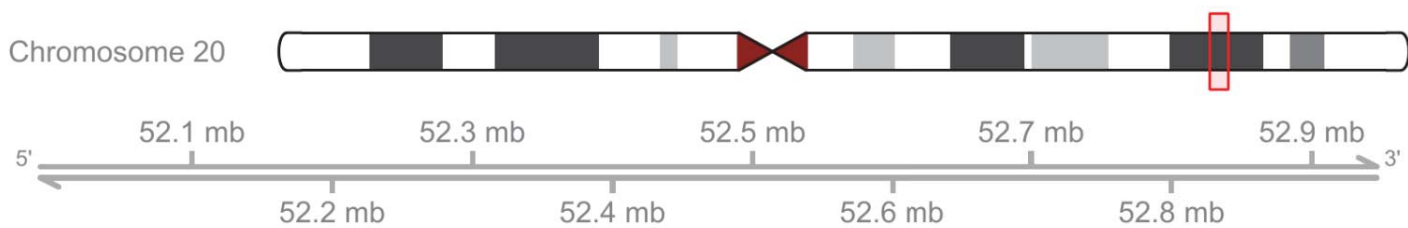
Uncorrected



Zscore of the effect allele: A



Pericalcarine SA: rs4811476



TSHZ2

BCAS1

ZNF217

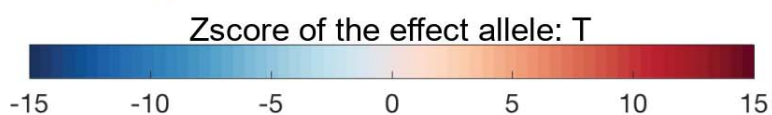
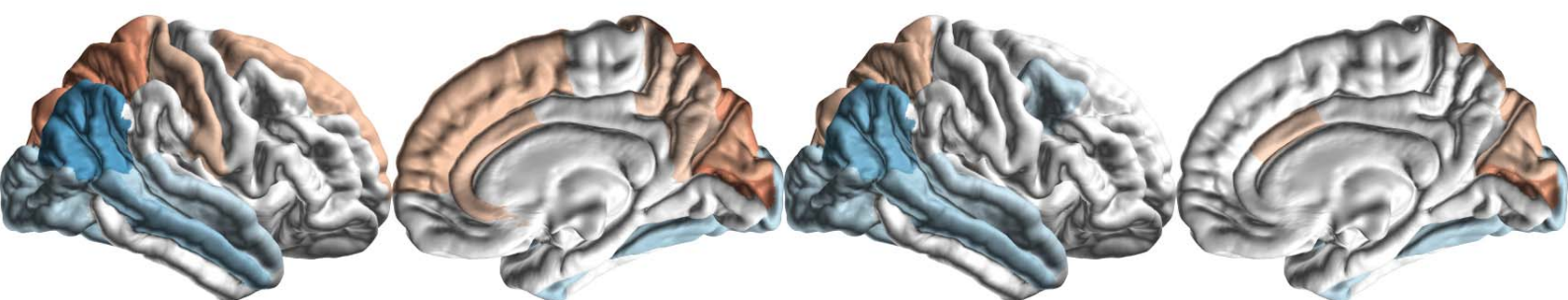
CYP24A1

PFDN4

CP>GZ
GZ>CP

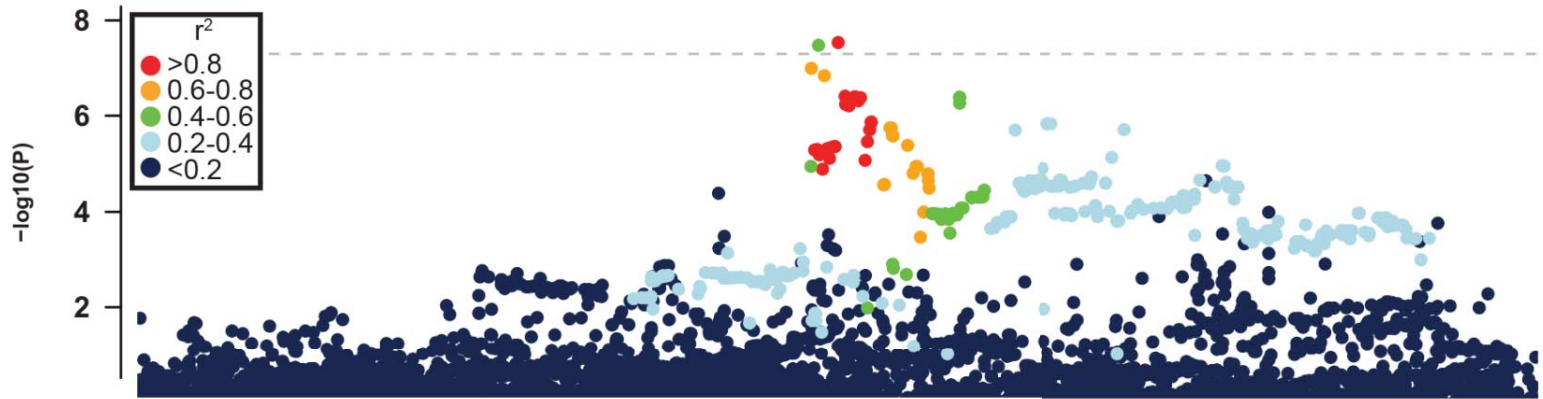
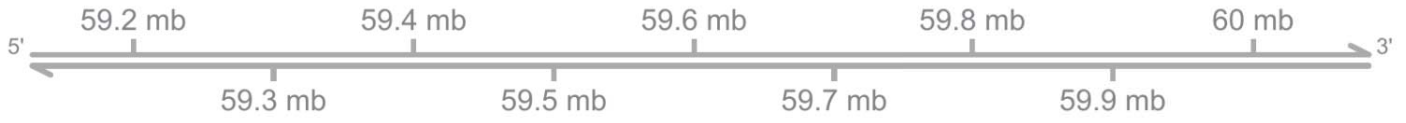
Corrected for total SA

Uncorrected



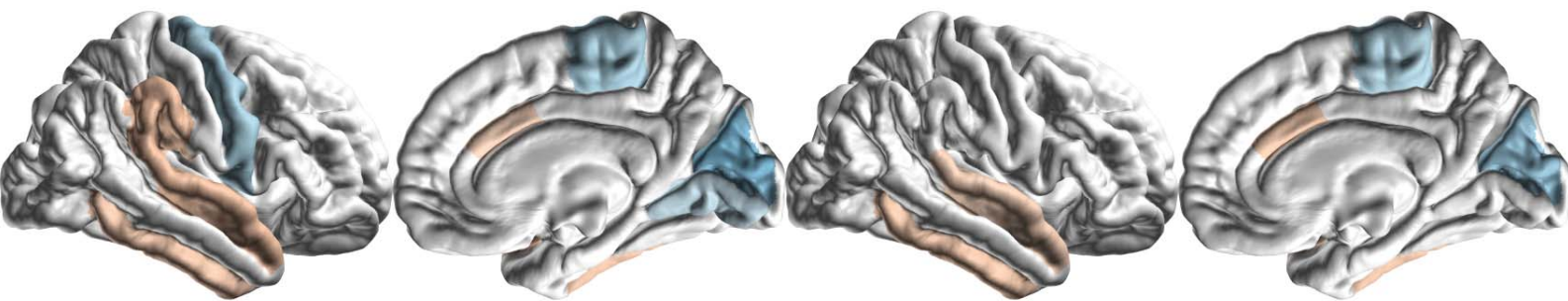
Pericalcarine TH: rs117461235

Chromosome 14

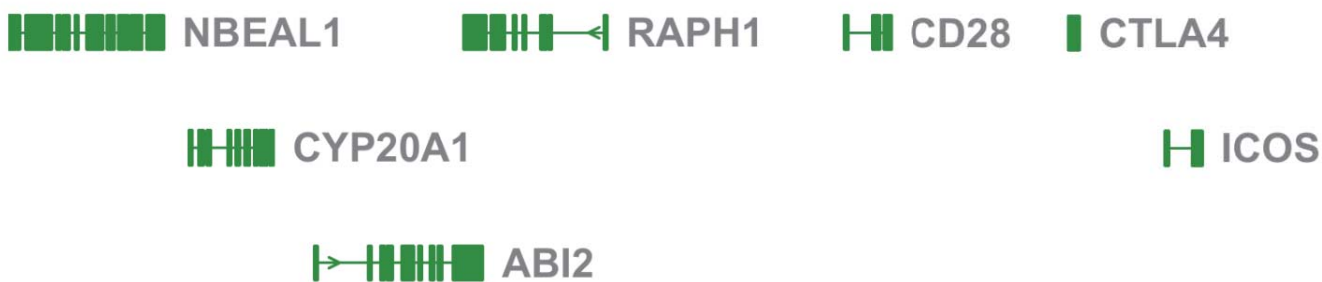
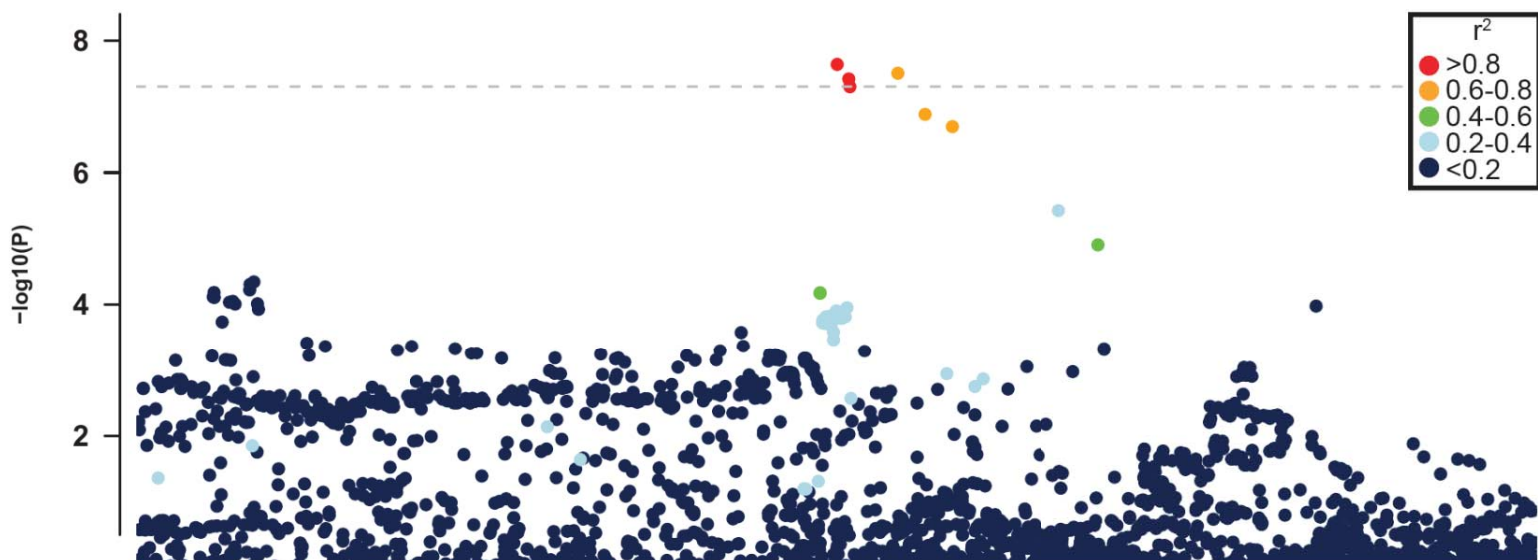


Corrected for average TH

Uncorrected

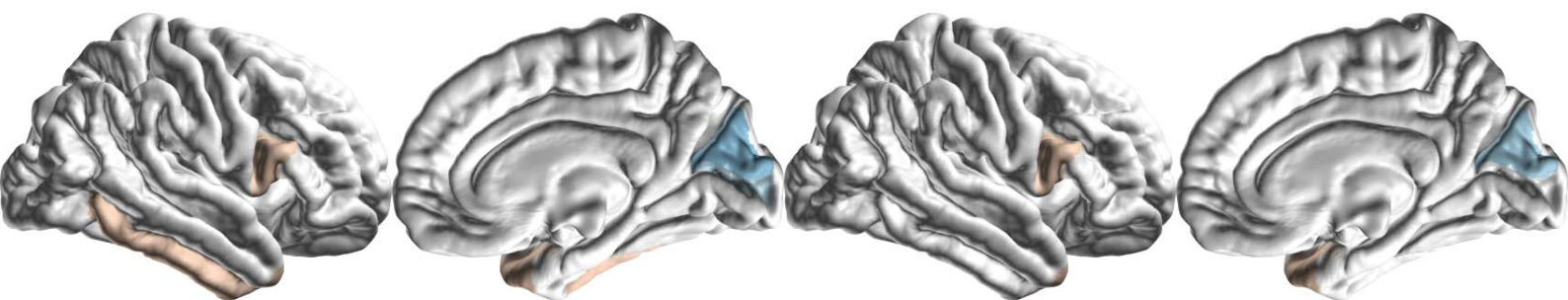


Cuneus SA: rs7585601



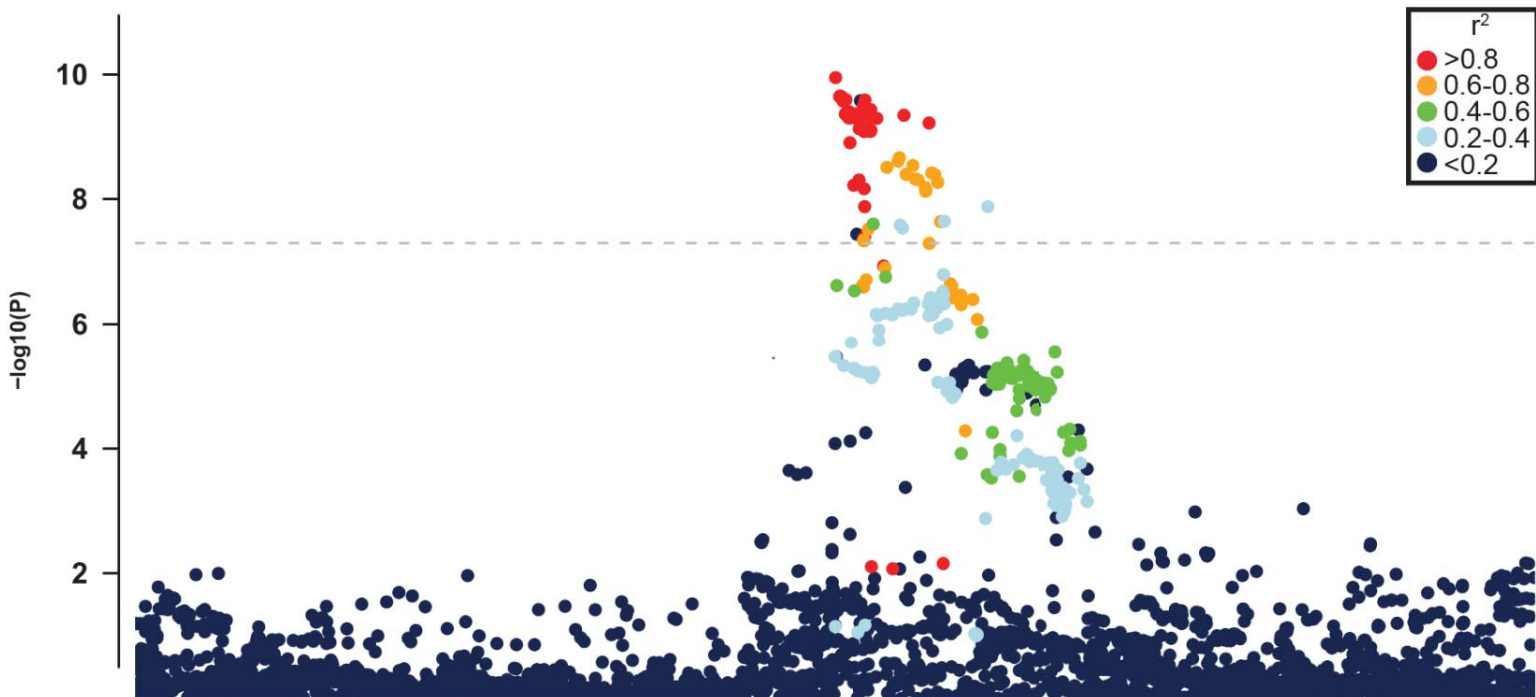
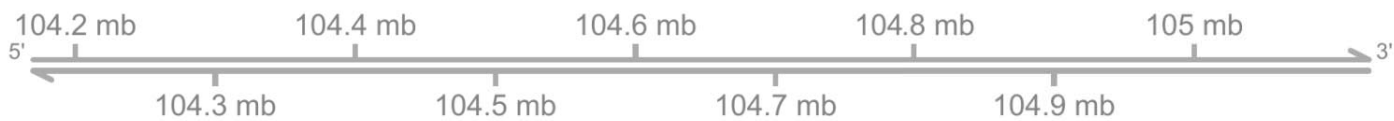
Corrected for total SA

Uncorrected



Cuneus SA: rs4895120

Chromosome 3



GZ>CP

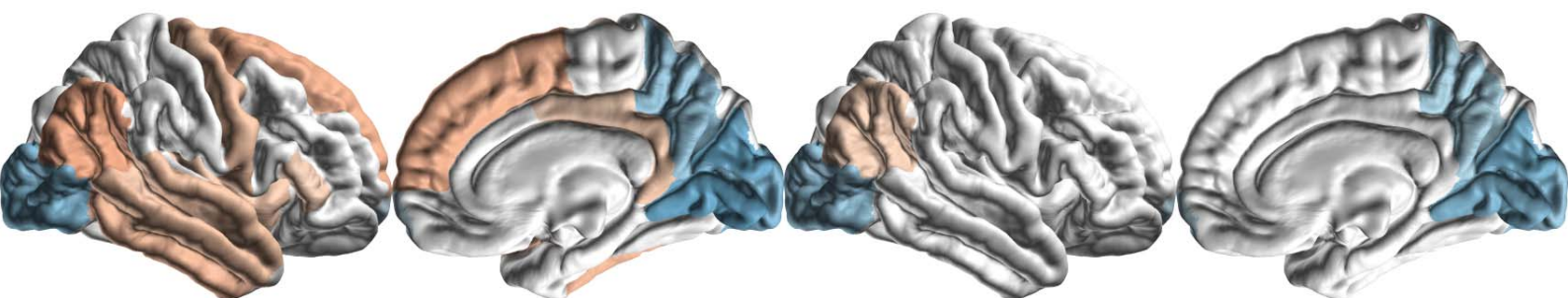


CP>GZ

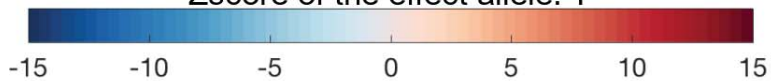


Corrected for total SA

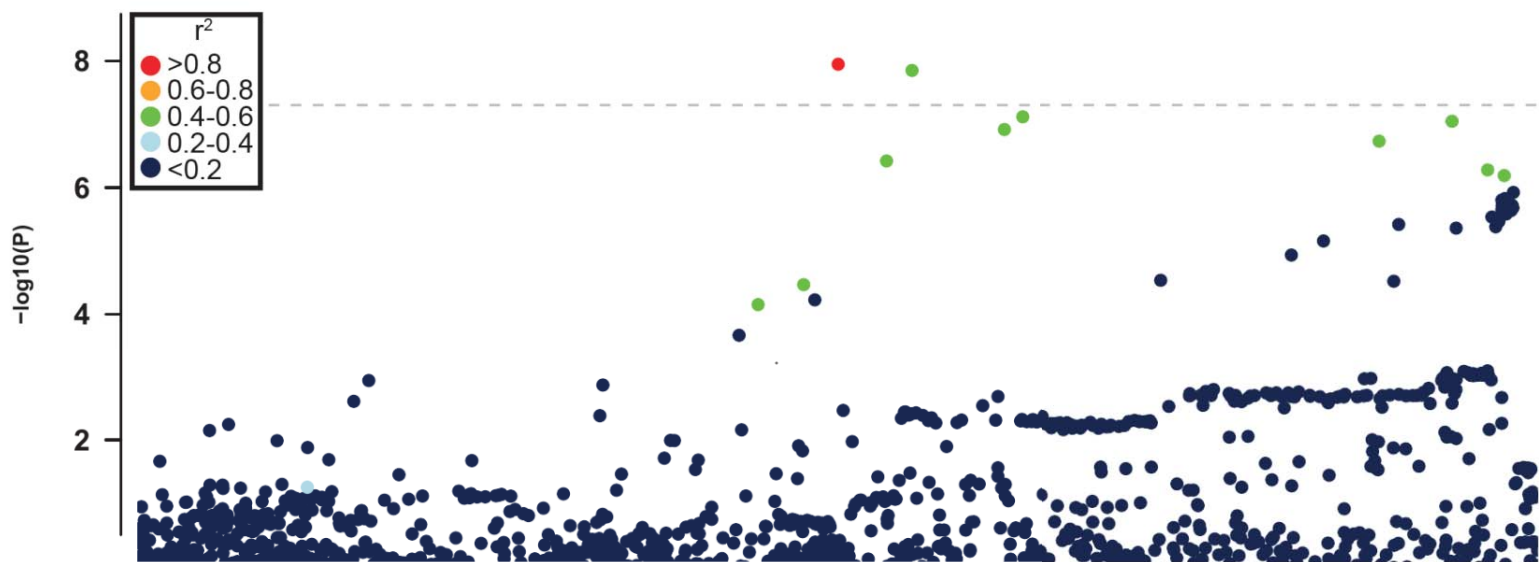
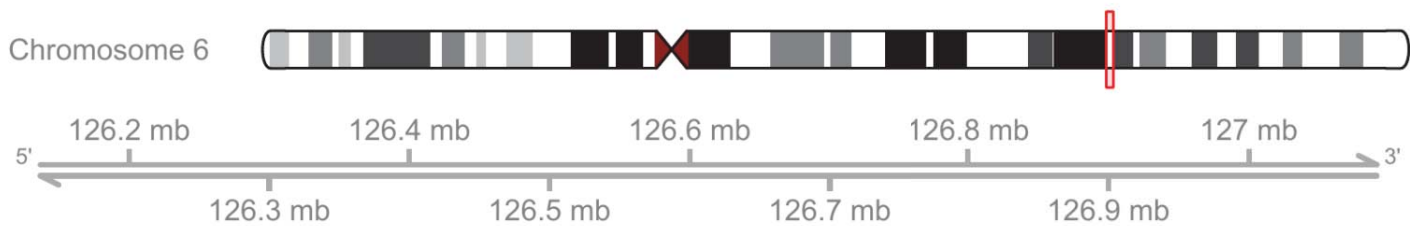
Uncorrected



Zscore of the effect allele: T



Cuneus SA: rs76470478



NCOA7 CENPW

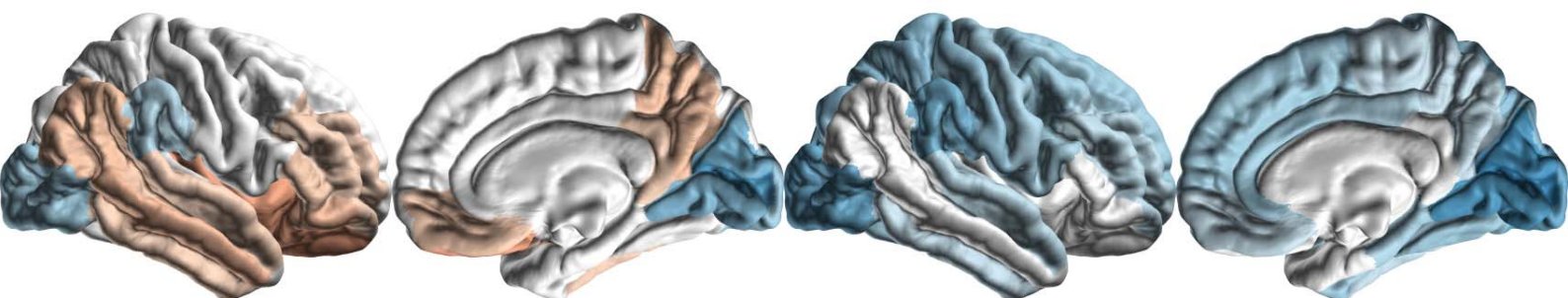
HINT3

TRMT11

GZ>CP
CP>GZ

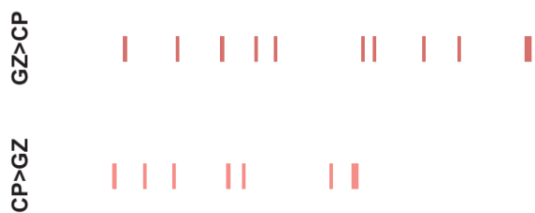
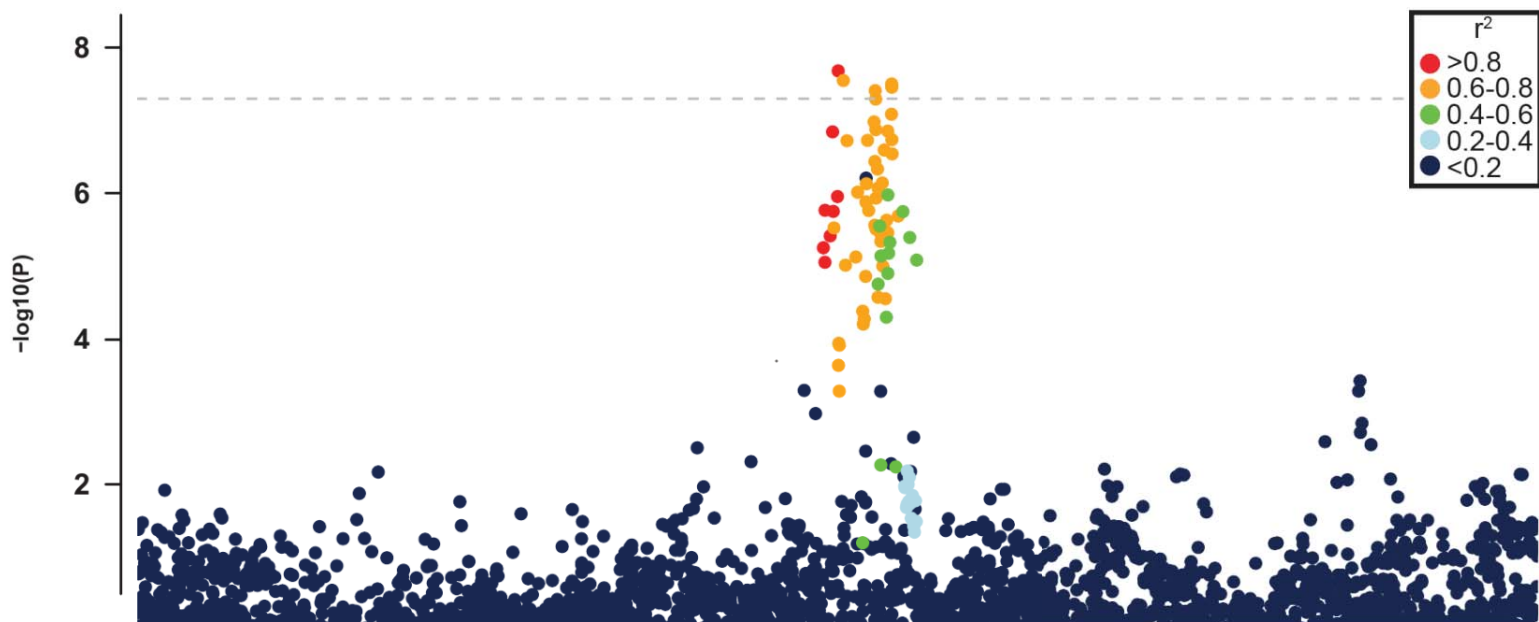
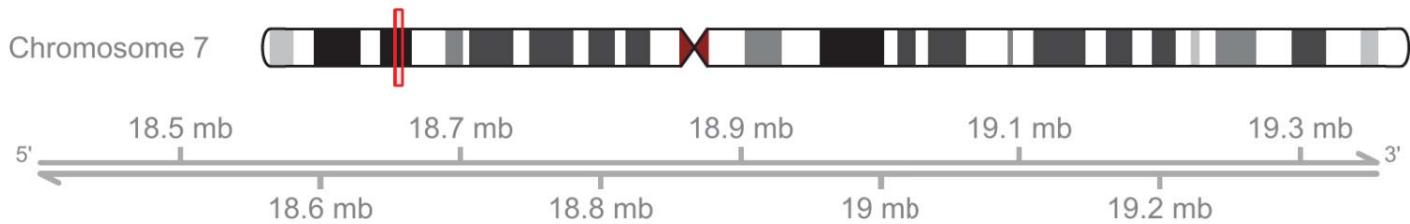
Corrected for total SA

Uncorrected



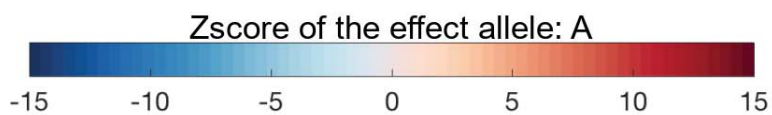
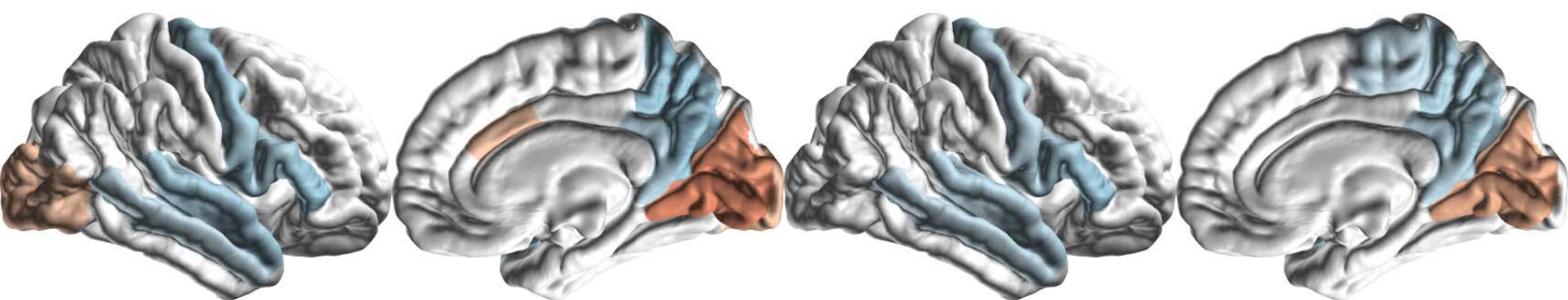
-15 -10 -5 0 5 10 15

Cuneus SA: rs957960

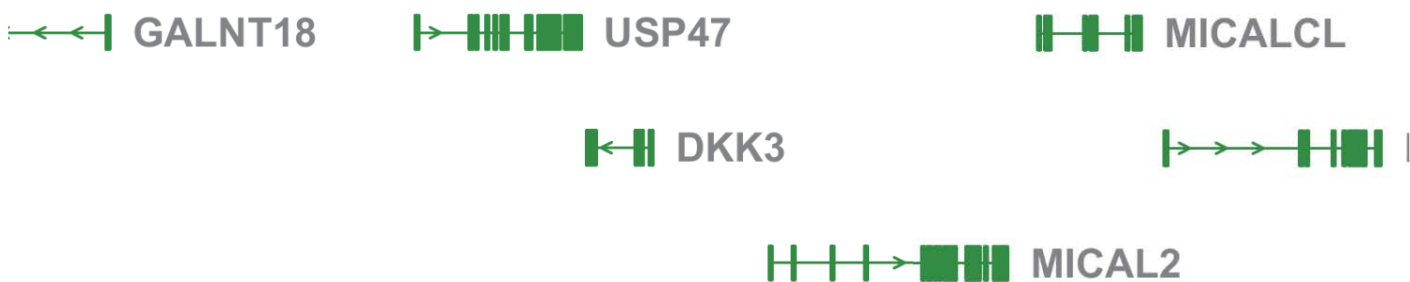
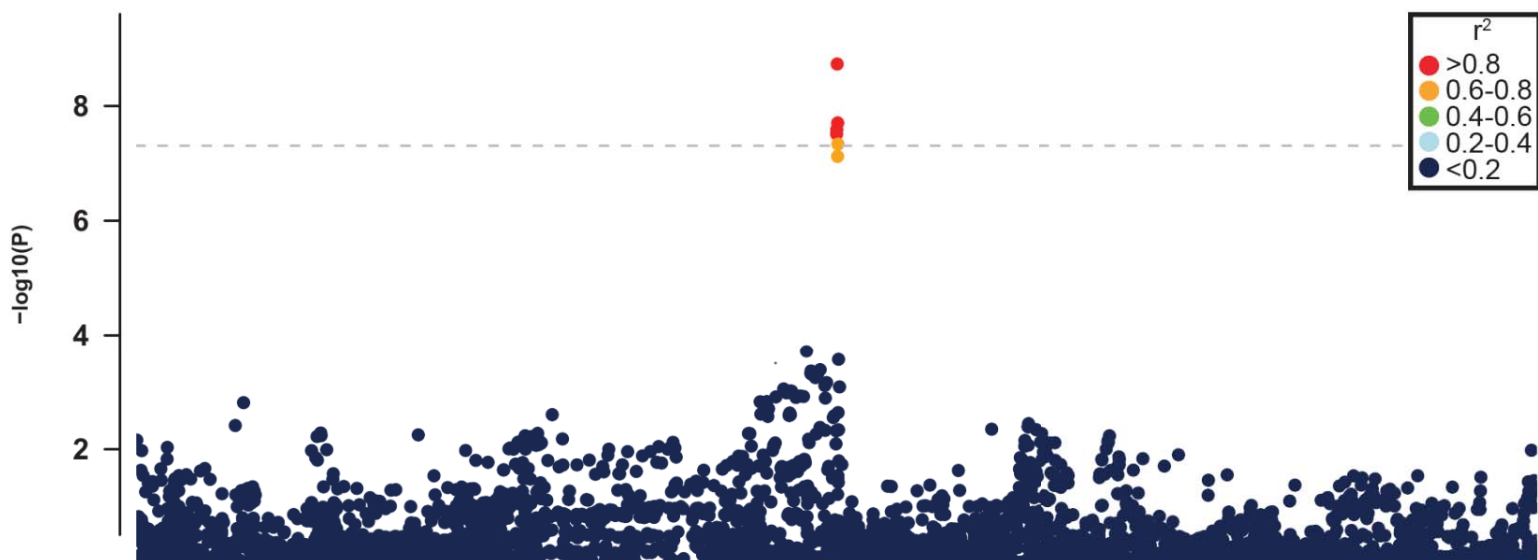
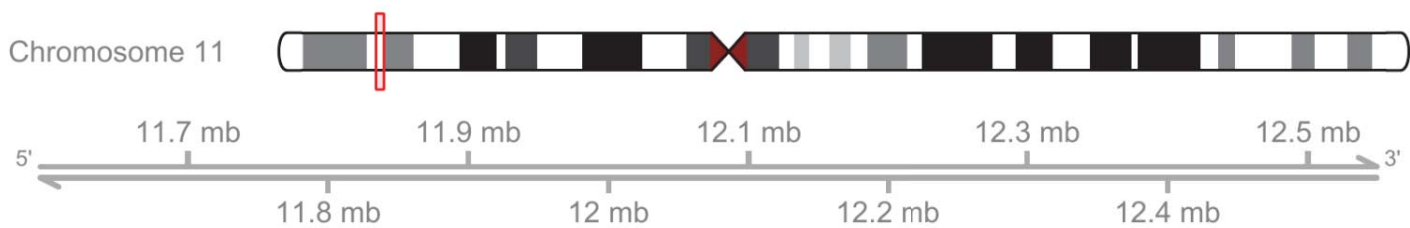


Corrected for total SA

Uncorrected

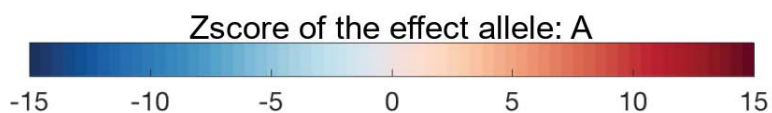
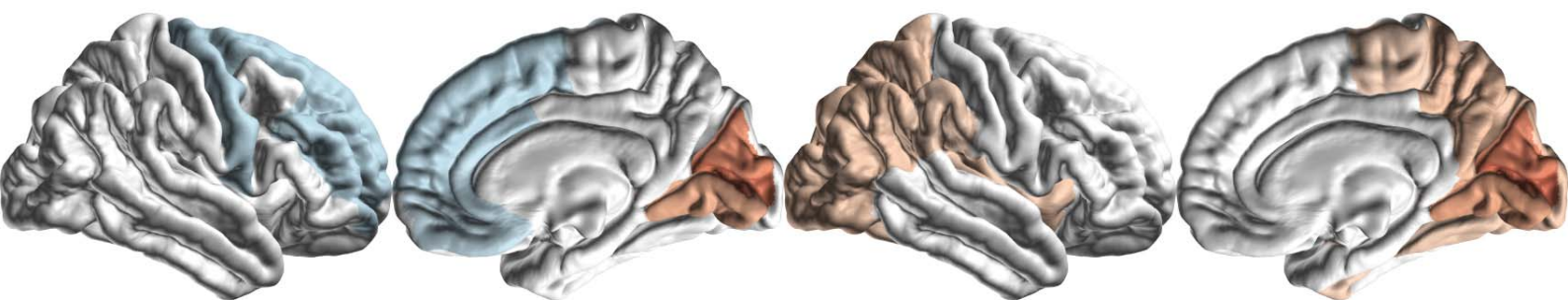


Cuneus SA: rs10765918

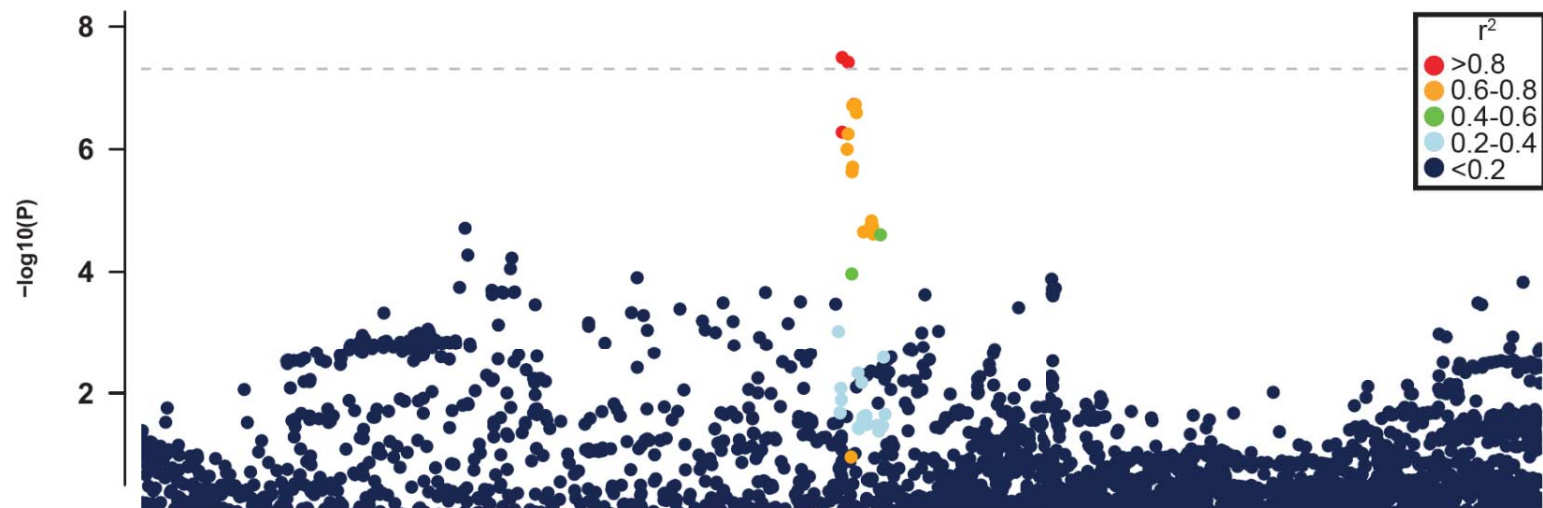
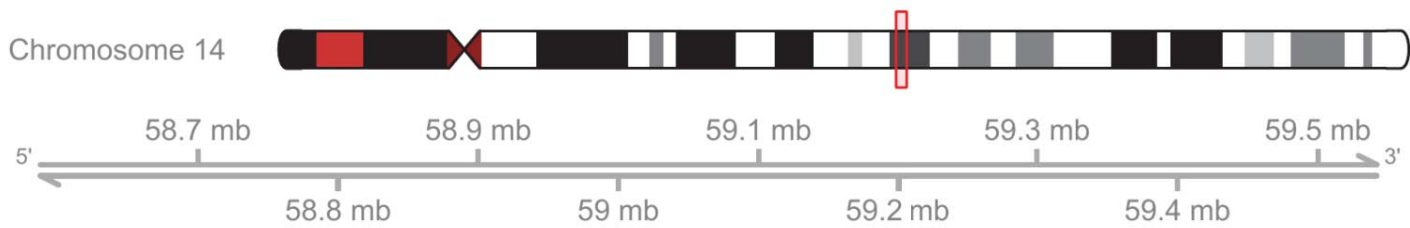


Corrected for total SA

Uncorrected



Cuneus SA: rs170239



C14orf37 ARID4A DACT1

ACTR10 TOMM20L

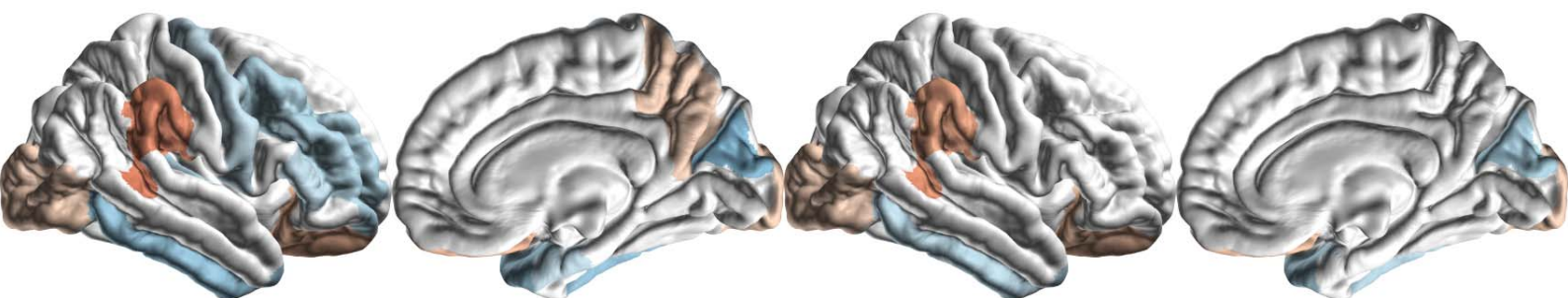
PSMA3 TIMM9

KIAA0586

CP>GZ GZ>CP

Corrected for total SA

Uncorrected

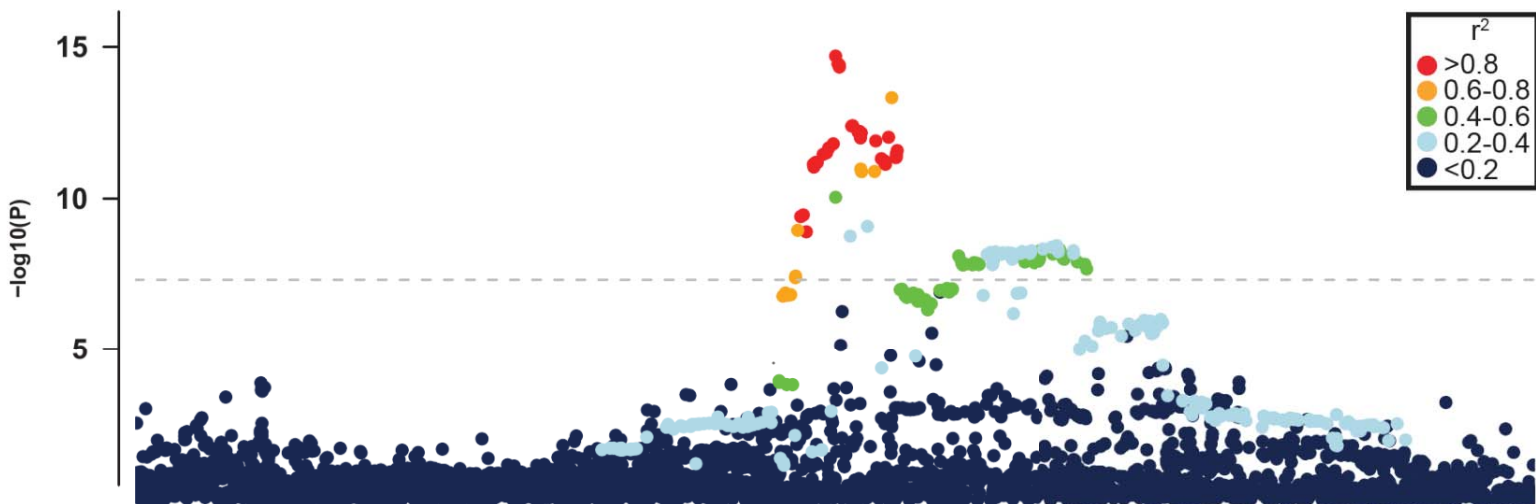
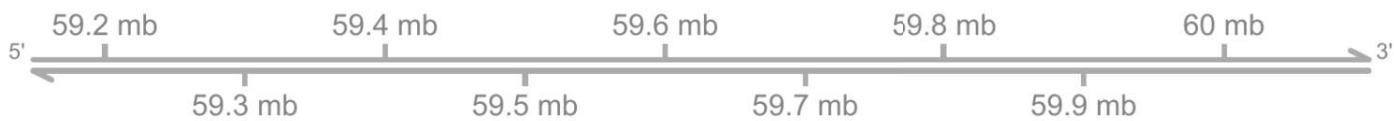


Zscore of the effect allele: T

-15 -10 -5 0 5 10 15

Cuneus SA: rs73313052

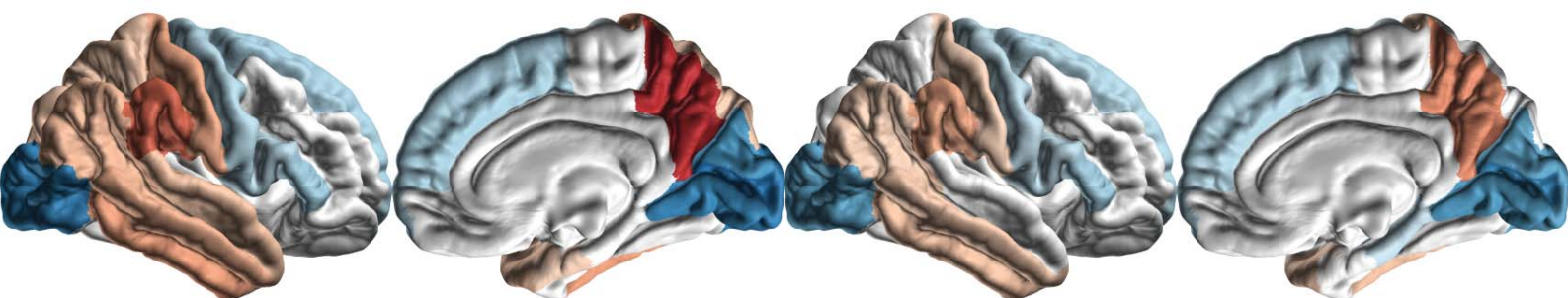
Chromosome 14



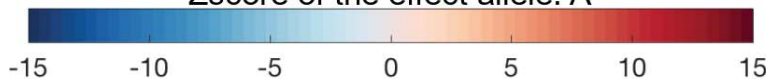
CP>GZ
GZ>CP

Corrected for total SA

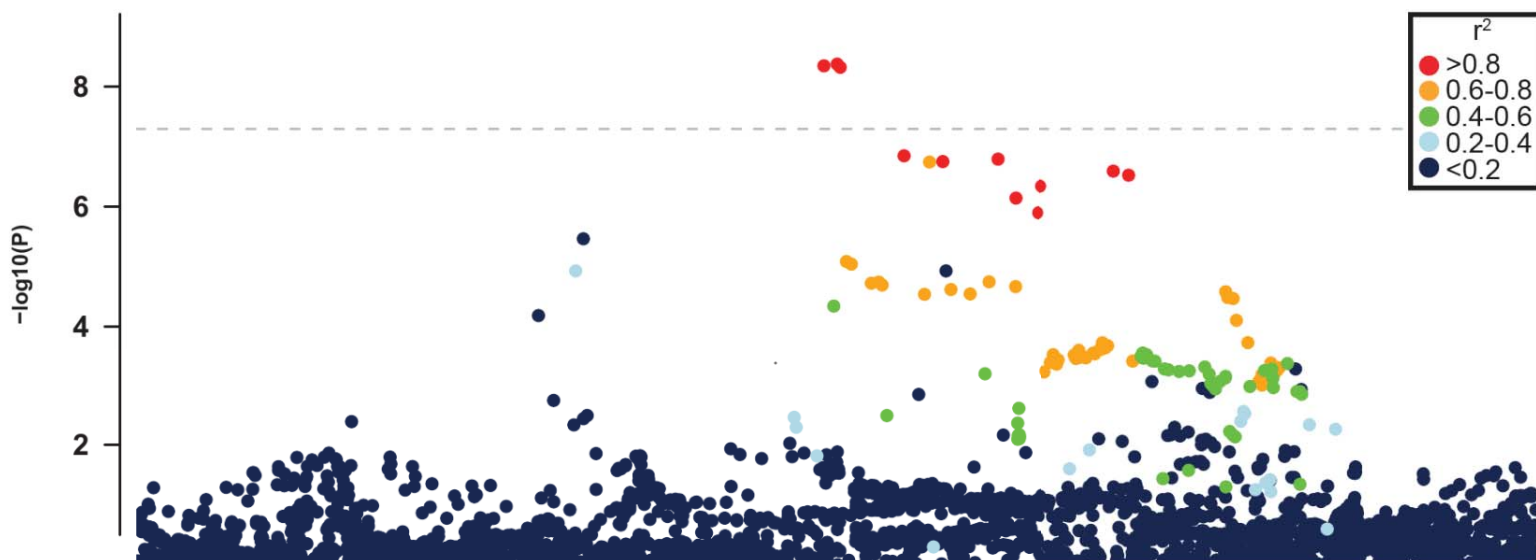
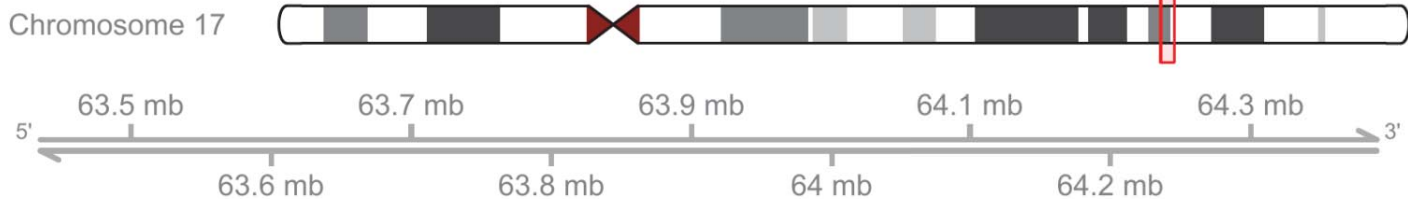
Uncorrected



Zscore of the effect allele: A



Cuneus SA: rs11652557



AXIN2

APOH

CEP112

GZ>CP

||

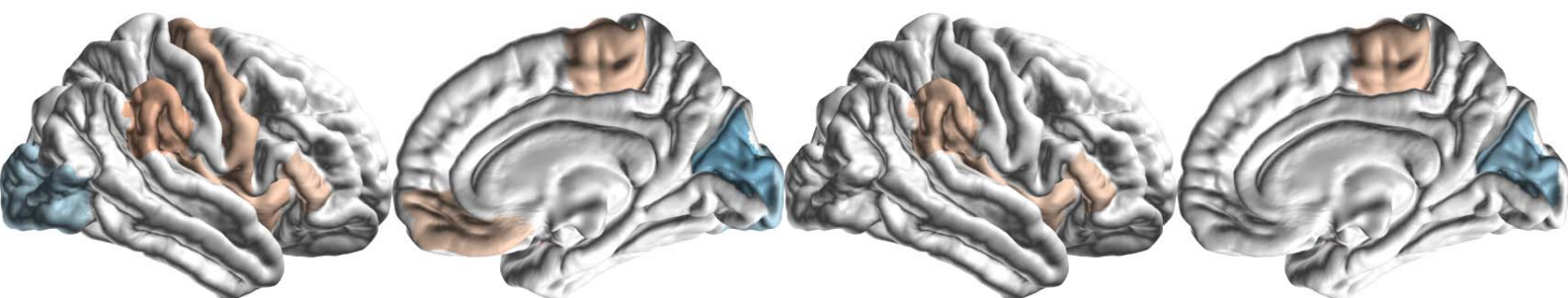
||

CP>GZ

|

Corrected for total SA

Uncorrected

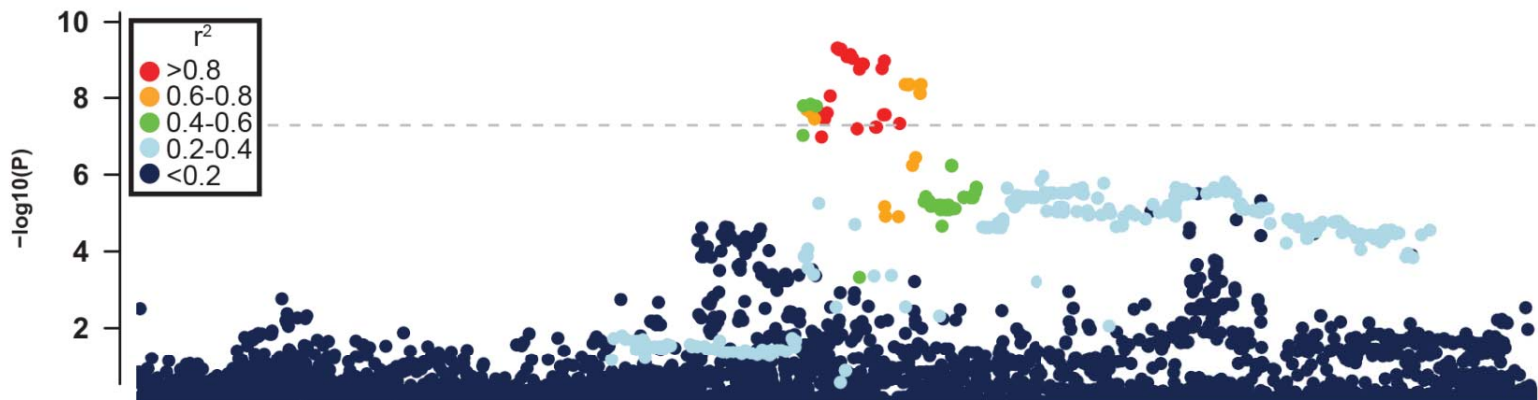
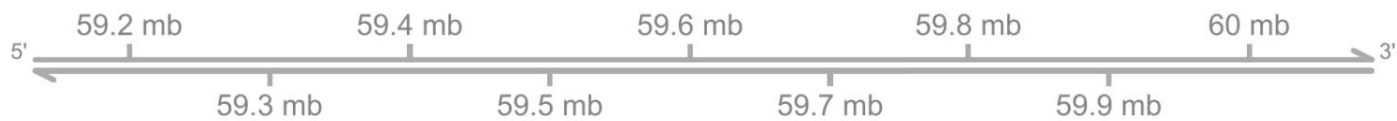


Zscore of the effect allele: A

-15 -10 -5 0 5 10 15

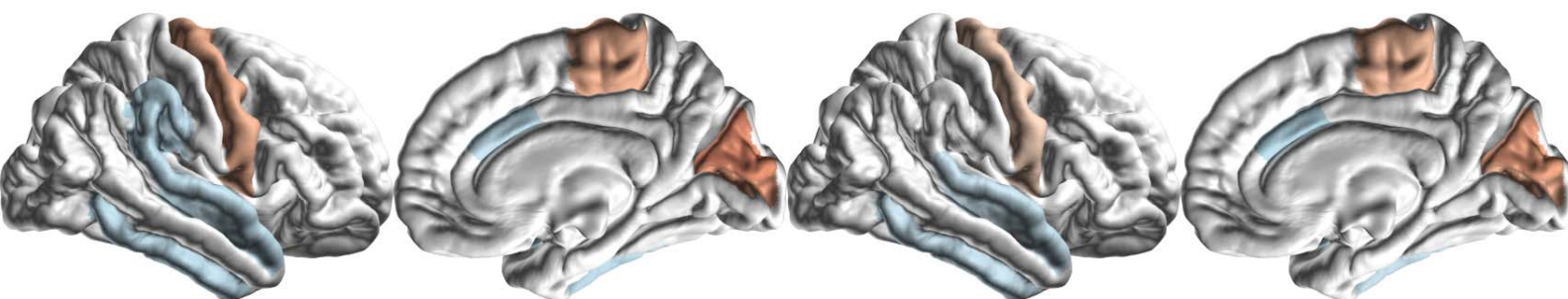
Cuneus TH: rs78580207

Chromosome 14

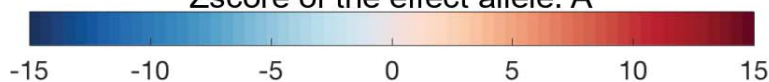


Corrected for average TH

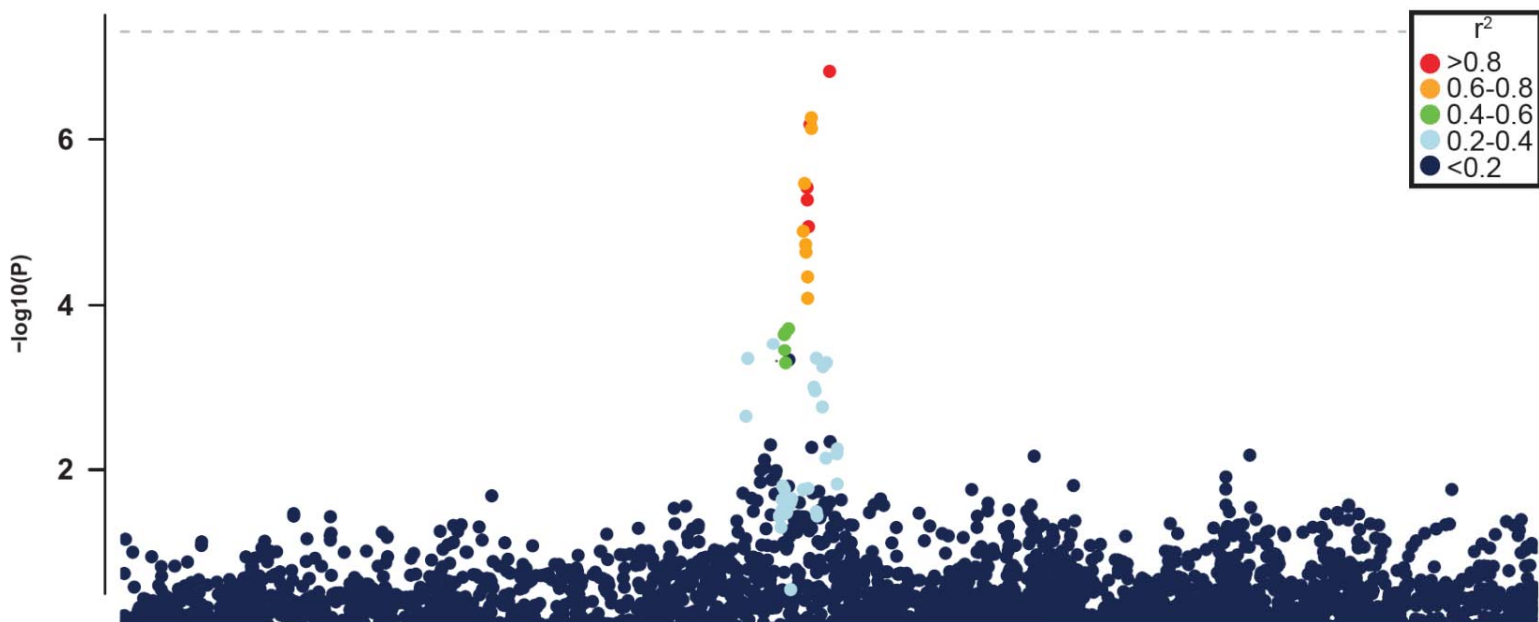
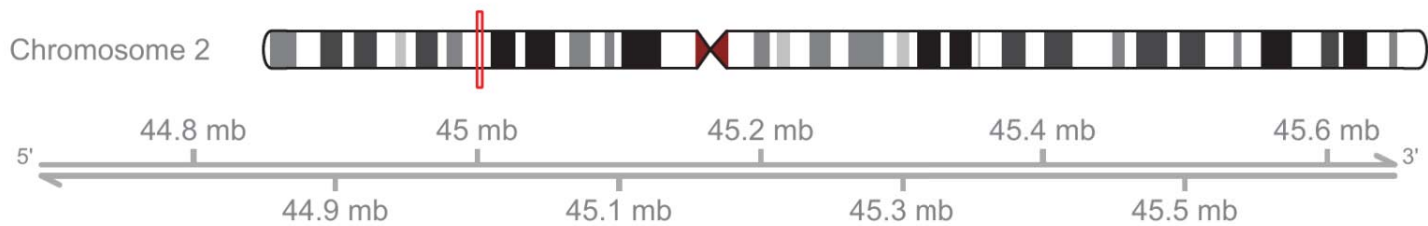
Uncorrected



Zscore of the effect allele: A



Lateral Occipital SA: rs4953152



GZ>CP

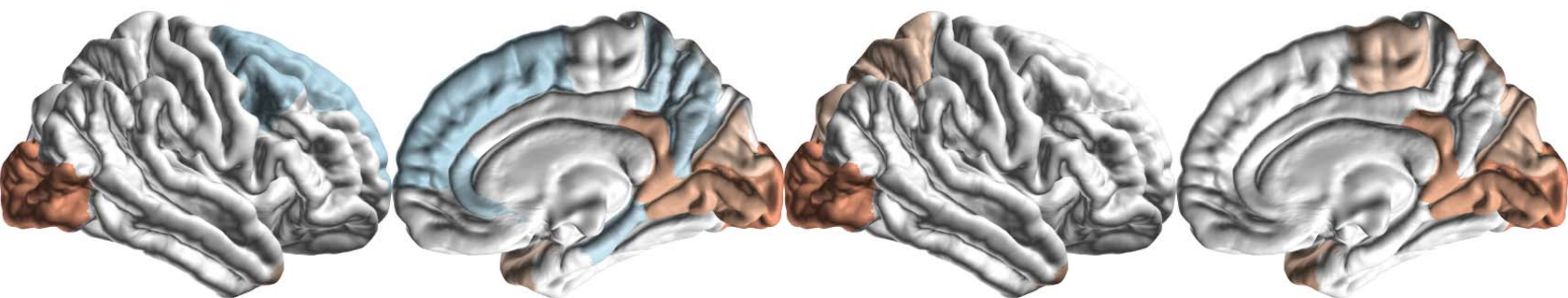


CP>GZ



Corrected for total SA

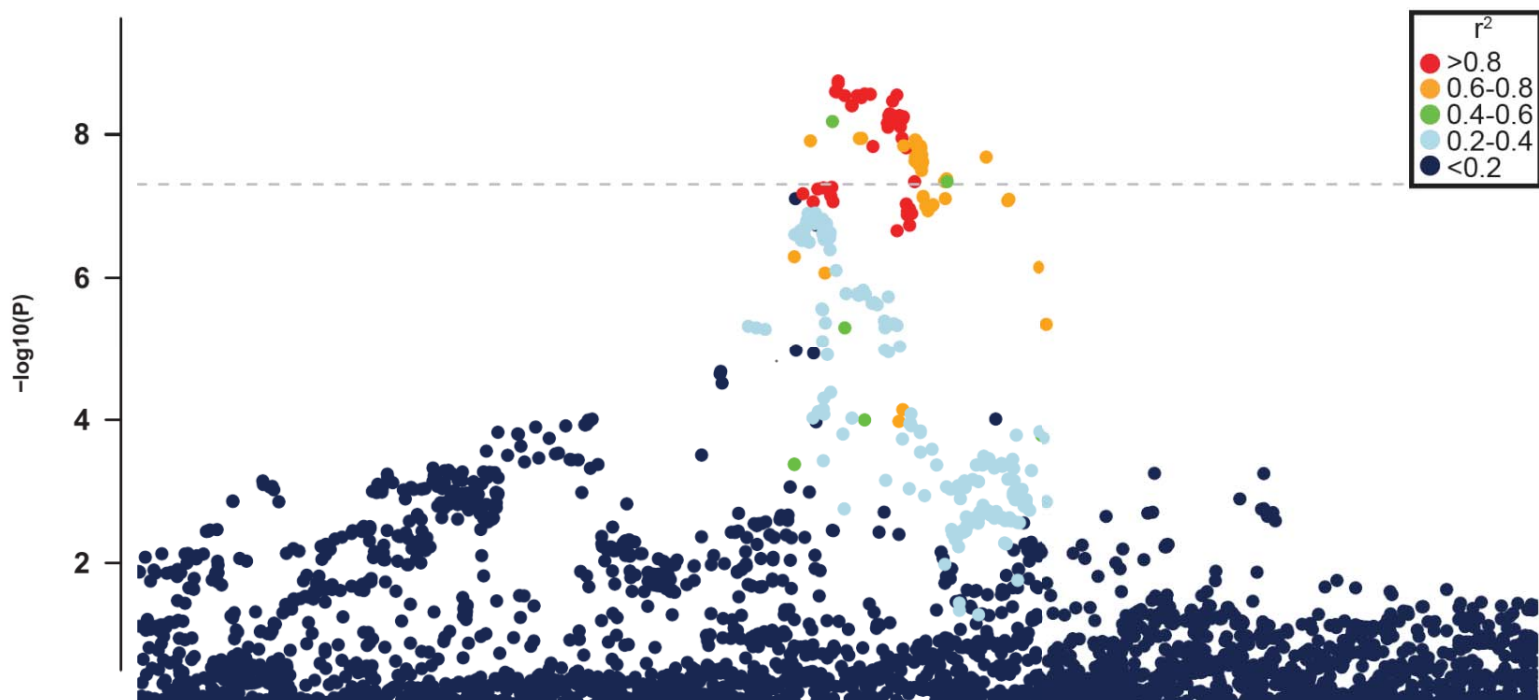
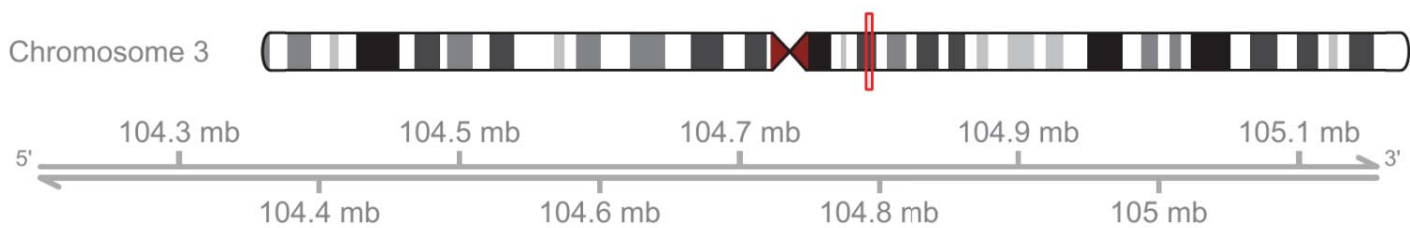
Uncorrected



Zscore of the effect allele: A

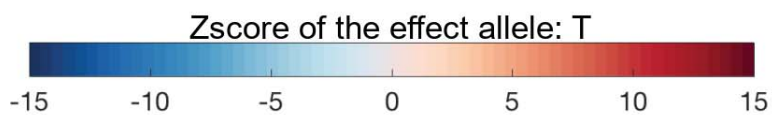
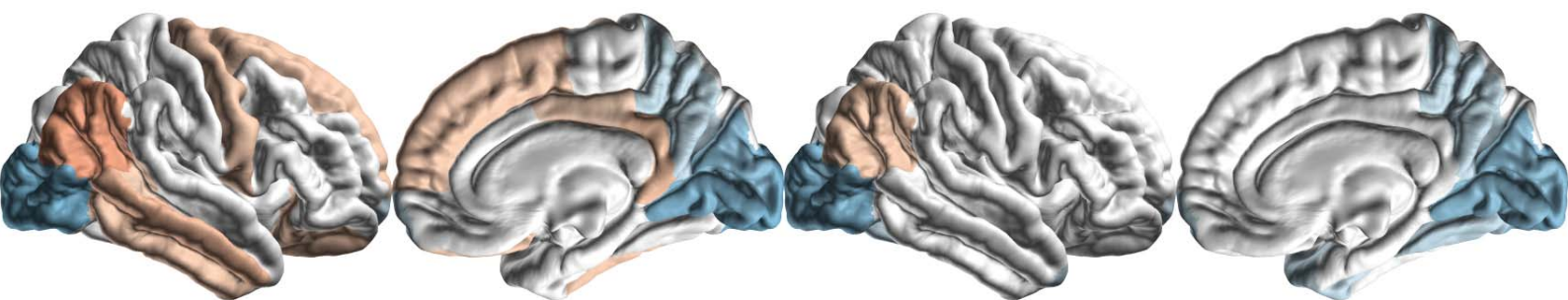


Lateral Occipital SA: rs9863836



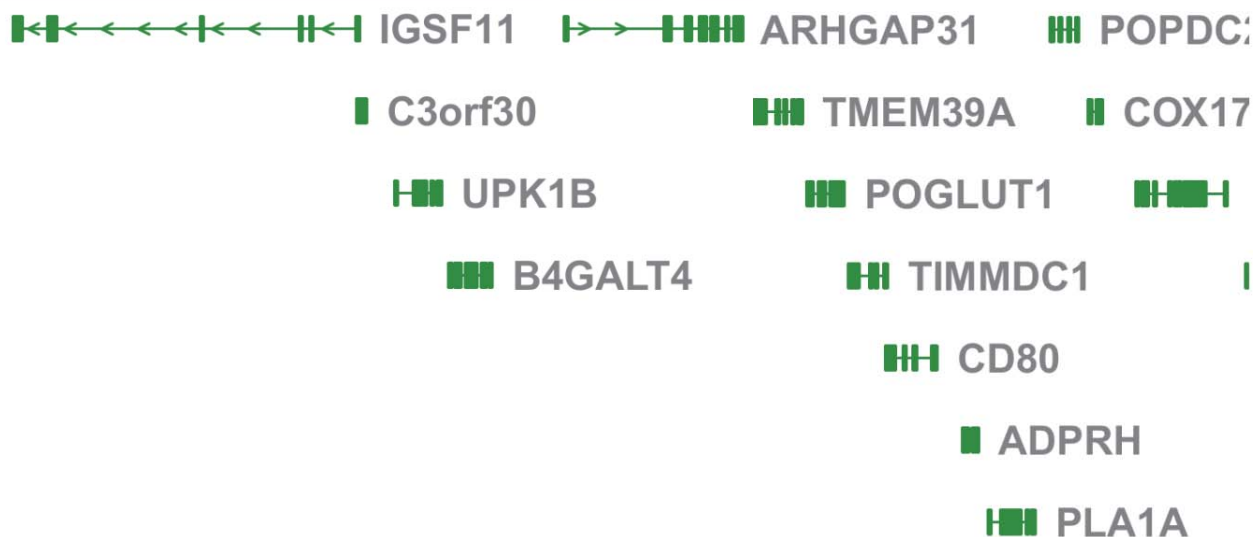
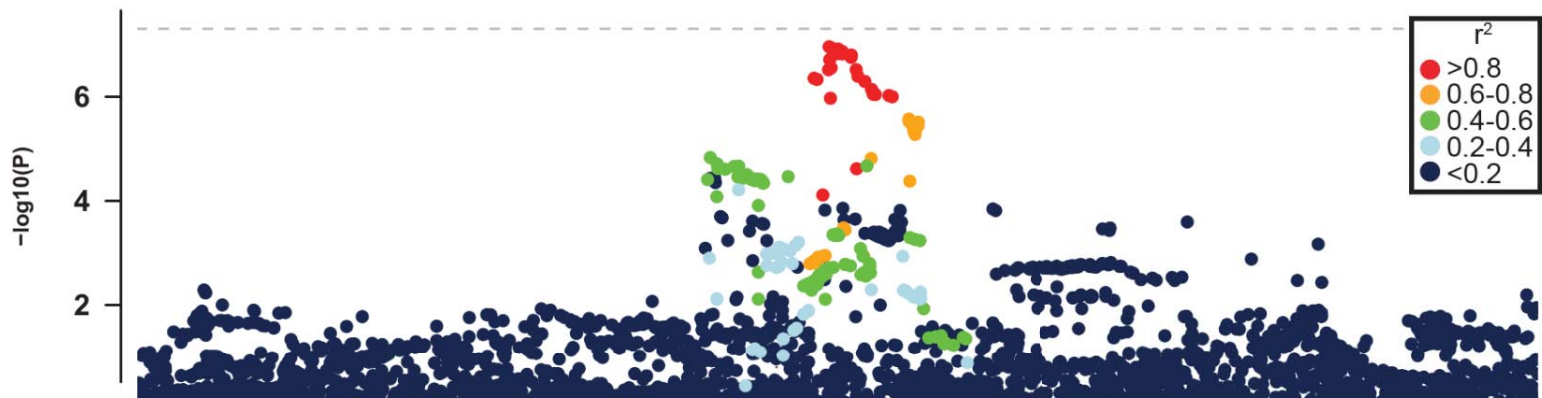
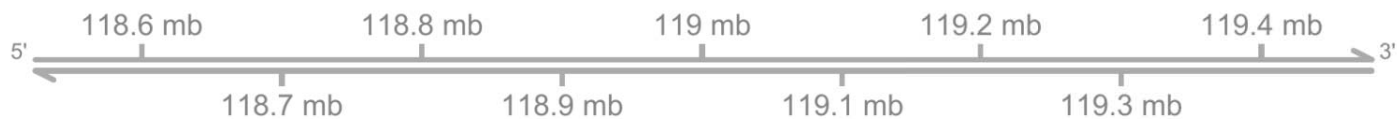
Corrected for total SA

Uncorrected



Lateral Occipital SA: rs9813454

Chromosome 3

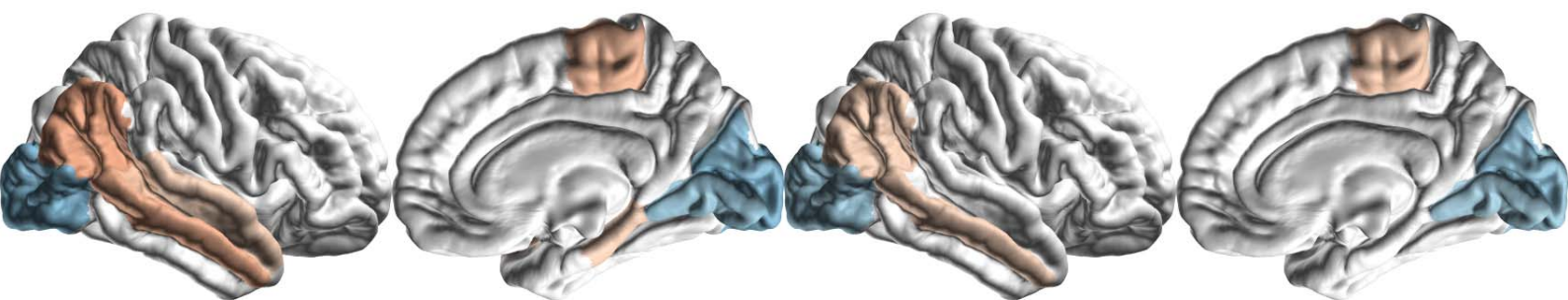


CP>GZ GZ>CP

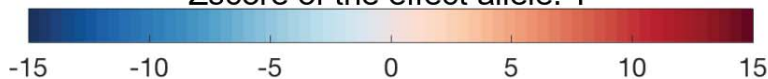


Corrected for total SA

Uncorrected

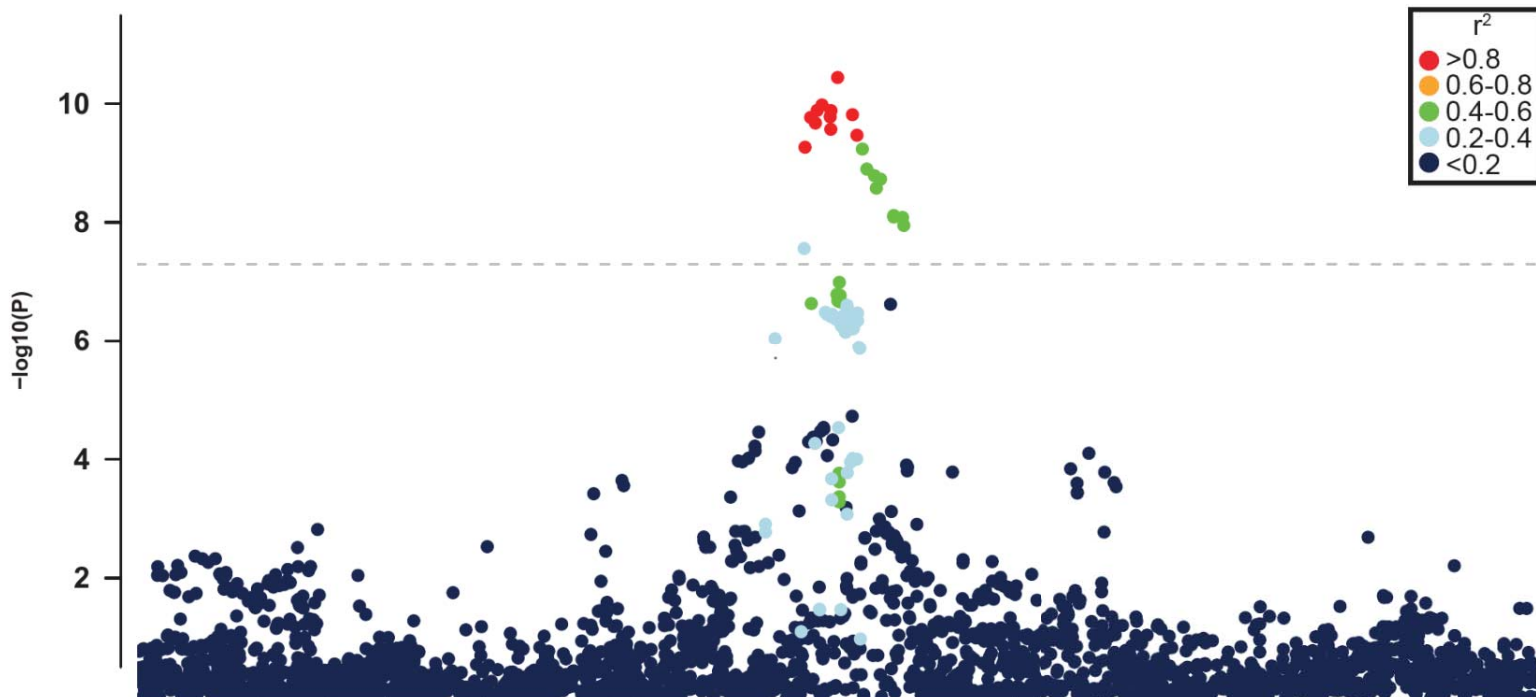
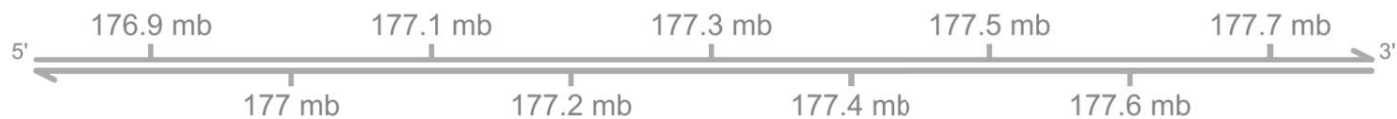


Zscore of the effect allele: T



Lateral Occipital SA: rs552305

Chromosome 3



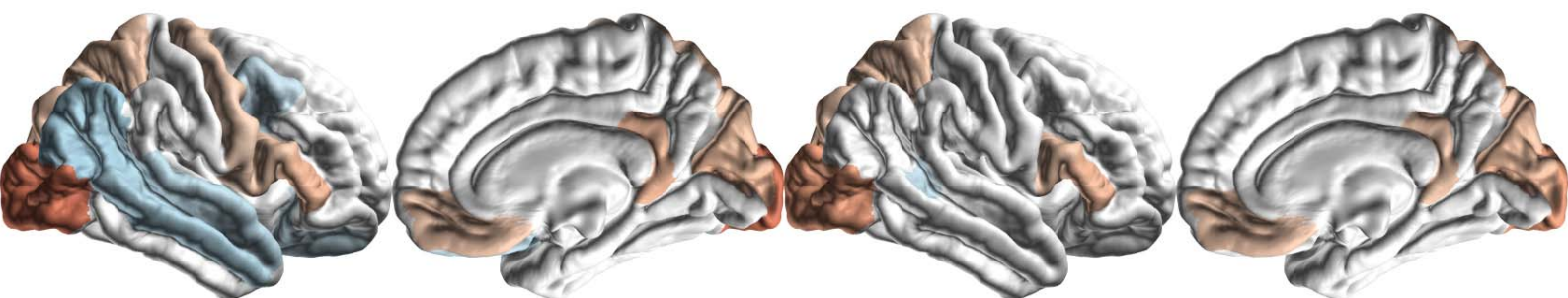
TBL1XR1

GZ>CP

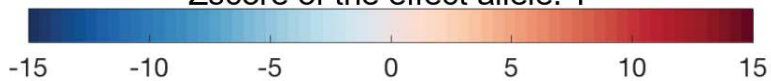
CP>GZ

Corrected for total SA

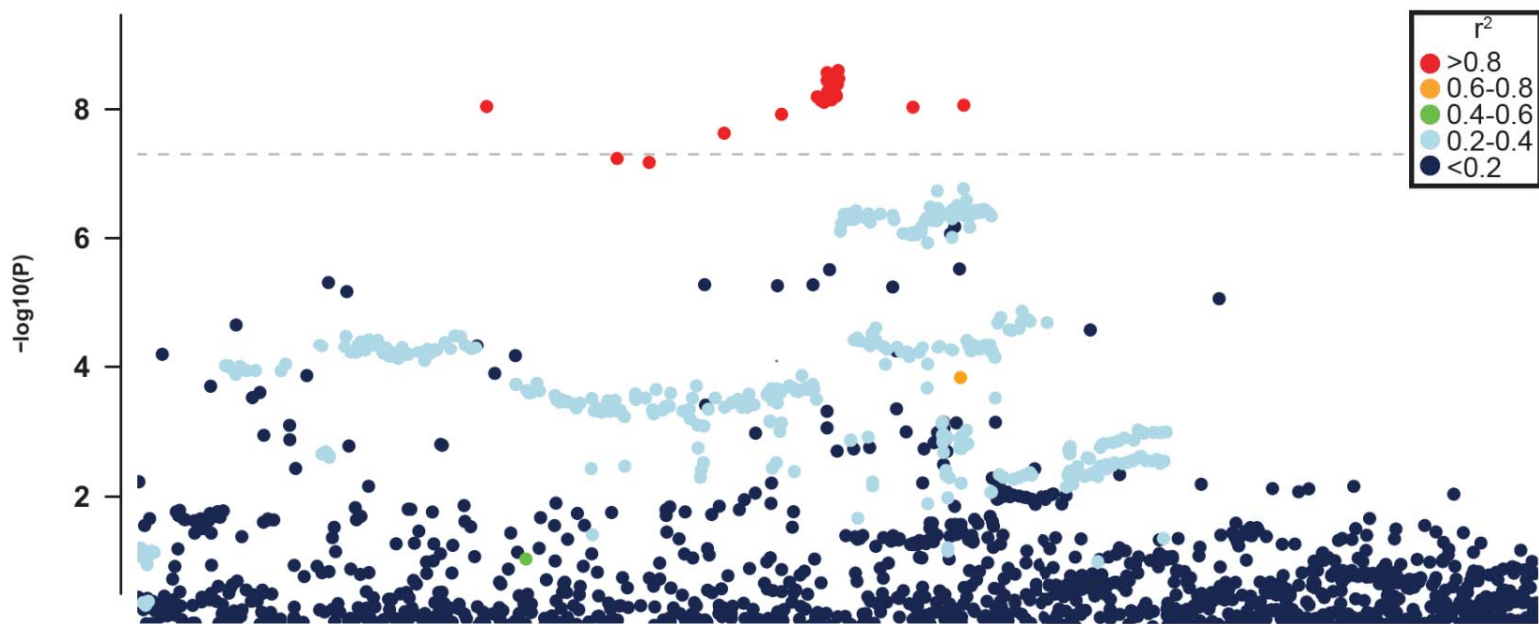
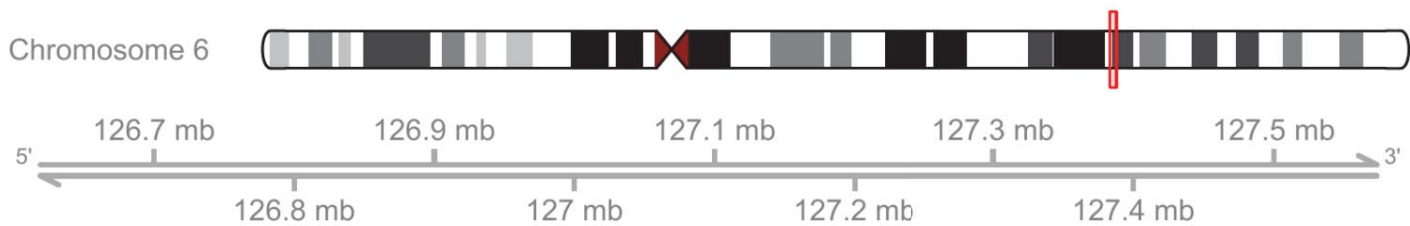
Uncorrected



Zscore of the effect allele: T



Lateral Occipital SA: rs9401907



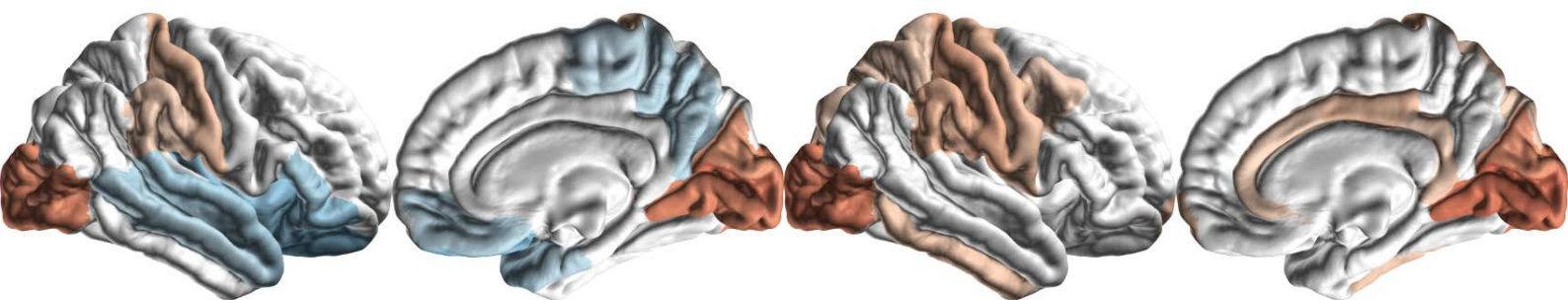
CENPW

RSP

GZ>CP
CP>GZ

Corrected for total SA

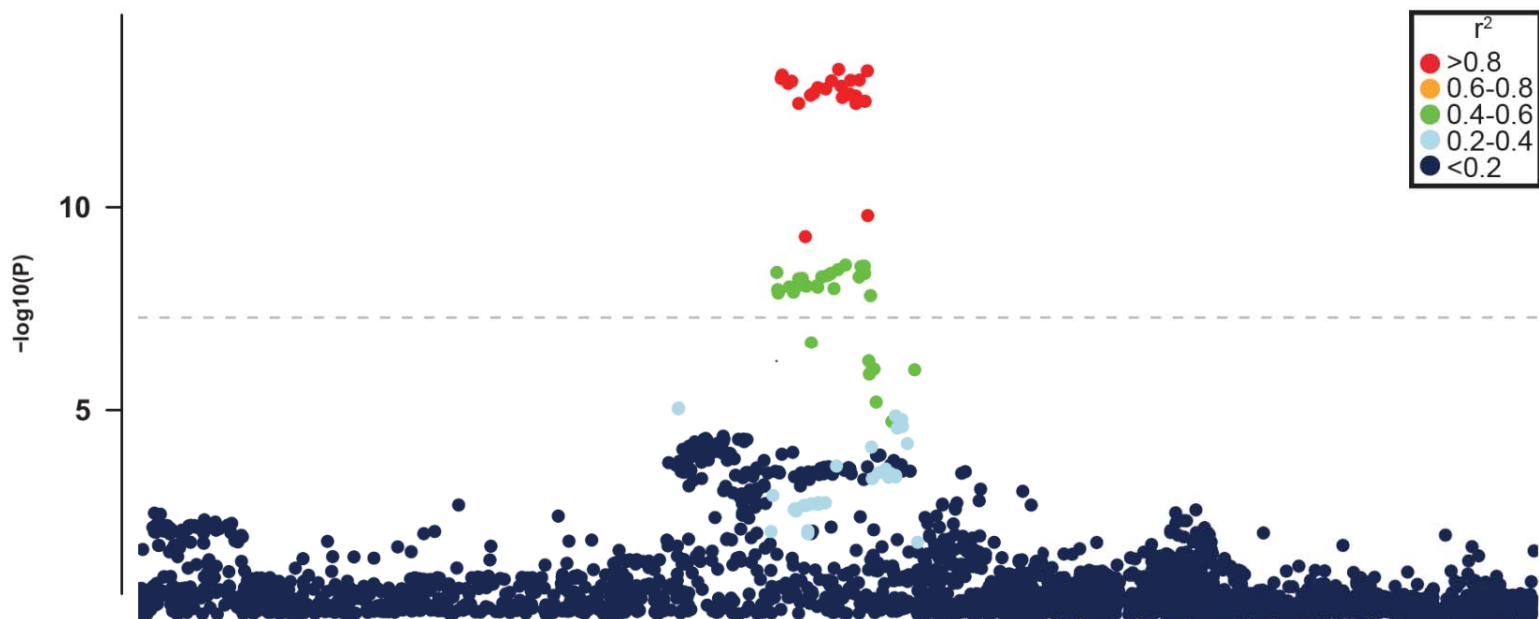
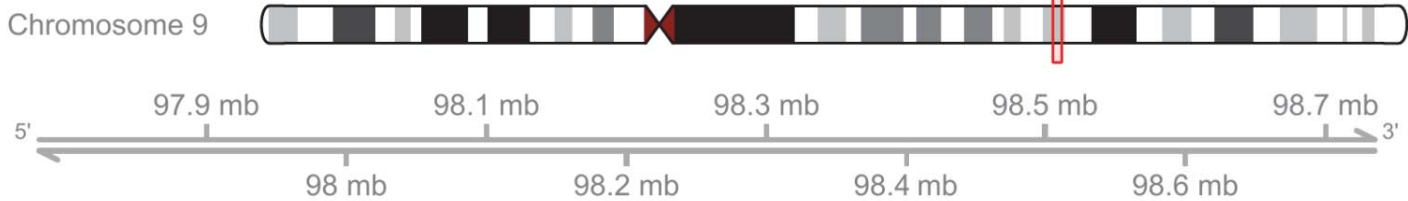
Uncorrected



Zscore of the effect allele: T



Lateral Occipital SA: rs28504650



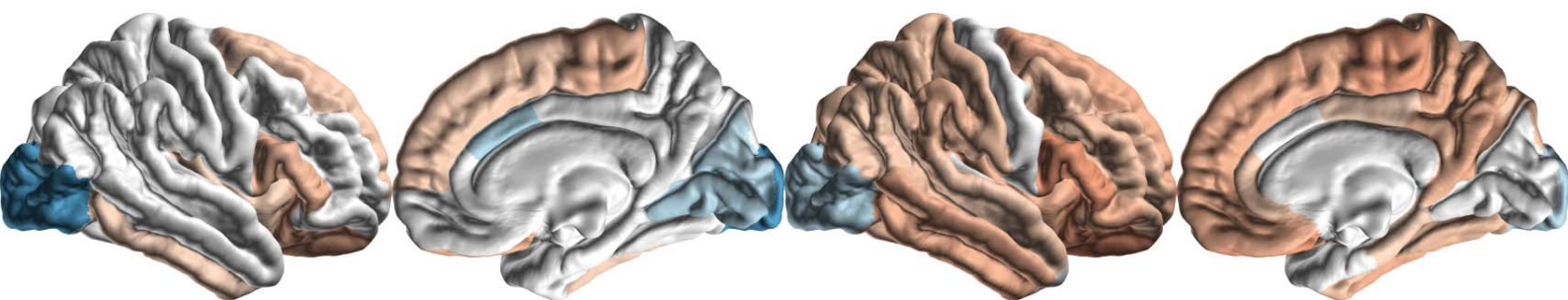
GZ>CP

CP>GZ

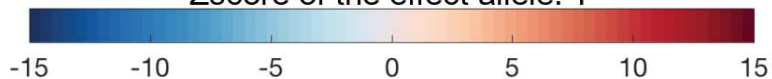


Corrected for total SA

Uncorrected

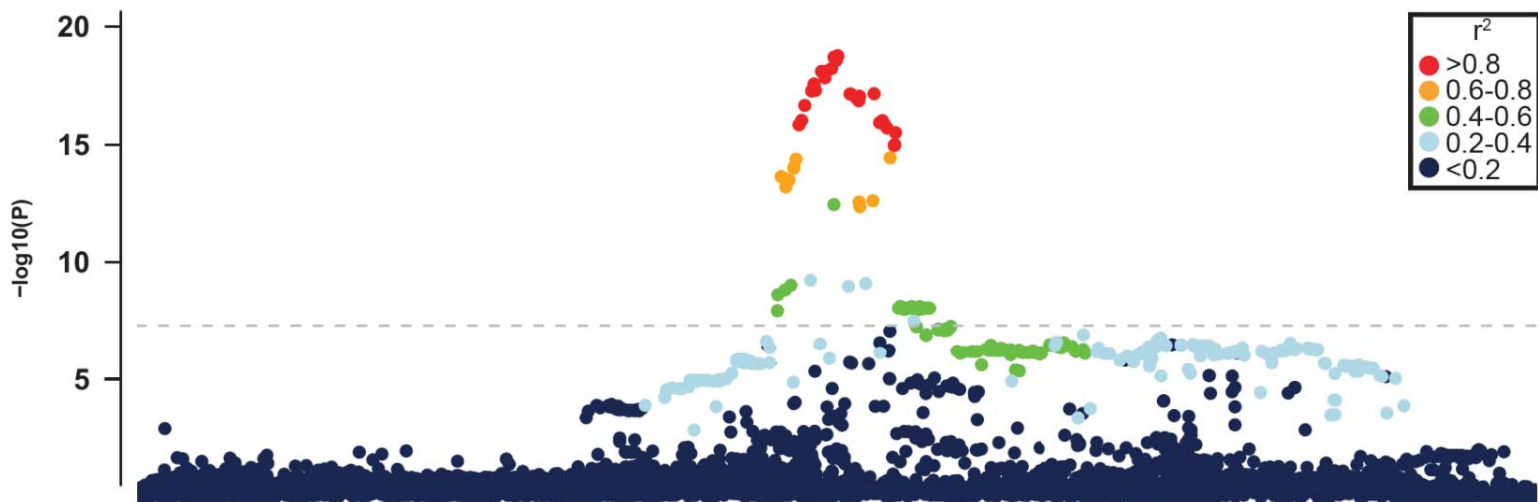
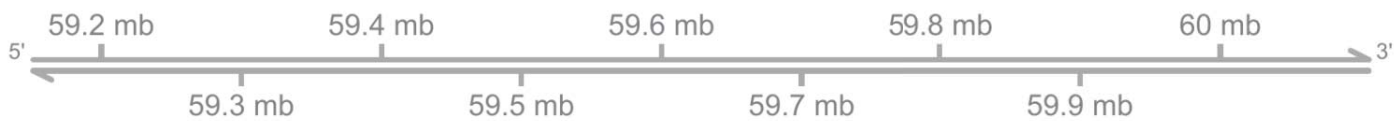


Zscore of the effect allele: T



Lateral Occipital SA: rs74826997

Chromosome 14



GPR135

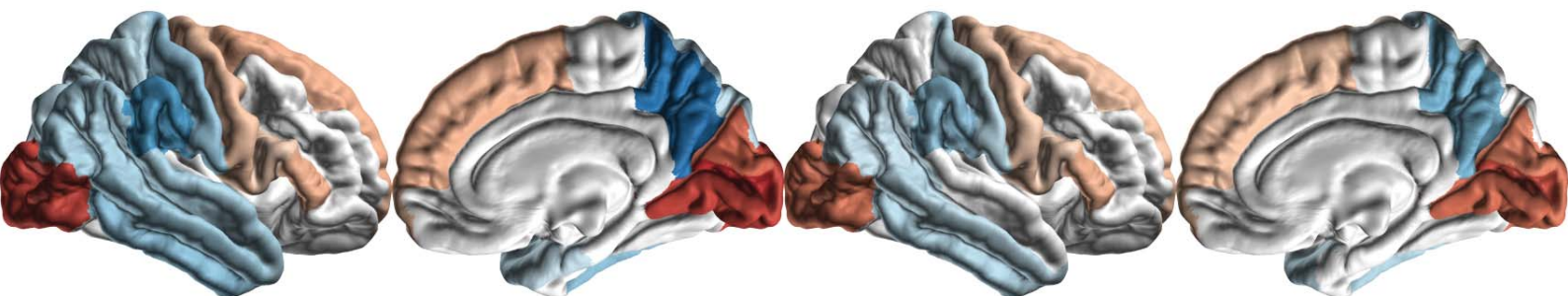
JKAMP

CCDC

CP>GZ
GZ>CP

Corrected for total SA

Uncorrected



Zscore of the effect allele: T

