

S2 Table. Gene-Wide Significant Genes from POLARIS Gene-based Analysis Conditioned on *APOE* and *BCAM* and best IGAP SNP and p-value

Chr	Gene	No. of SNPs	POLARIS, conditioned on <i>APOE</i> and <i>BCAM</i>			IGAP	
			Beta	SE	P-value	SNP	P-value
4	<i>PPARGC1A</i>	480	0.896	0.1896	2.3×10^{-6}	rs11734408	0.001345
8	<i>SCARA3 (CLU)</i>	240	0.549	0.1097	5.6×10^{-7}	rs7982	2.48×10^{-7}
15	<i>RORA</i>	1813	0.347	0.0693	5.6×10^{-7}	15-61211520	5.26×10^{-6}
16	<i>ZNF423</i>	1056	0.549	0.1194	4.3×10^{-6}	16-49888847	0.0003226
19	<i>BCL3</i>	88	0.167	0.0676	0.0133	rs12459810	6.77×10^{-43}
19	<i>CBL3</i>	50	0.151	0.1243	0.2237	rs2889414	2.49×10^{-43}
19	<i>BCAM</i>	71	NA	NA	NA	rs28399637	4.66×10^{-69}
19	<i>PVRL2</i>	160	0.255	0.0644	7.4×10^{-5}	rs6857	2.50×10^{-575}
19	<i>TOMM40</i>	108	0.335	0.0897	0.0002	rs6857	2.50×10^{-575}
19	<i>APOE</i>	55	NA	NA	NA	rs6857	2.50×10^{-575}
19	<i>APOC1</i>	34	-0.146	0.1045	0.1621	rs6857	2.50×10^{-575}
19	<i>APOC4-APOC2</i>	62	0.435	0.0897	1.2×10^{-6}	rs429358	6.70×10^{-536}