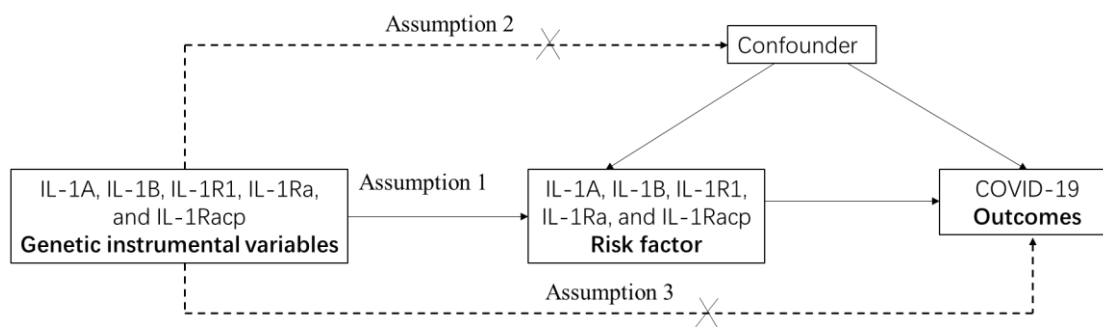


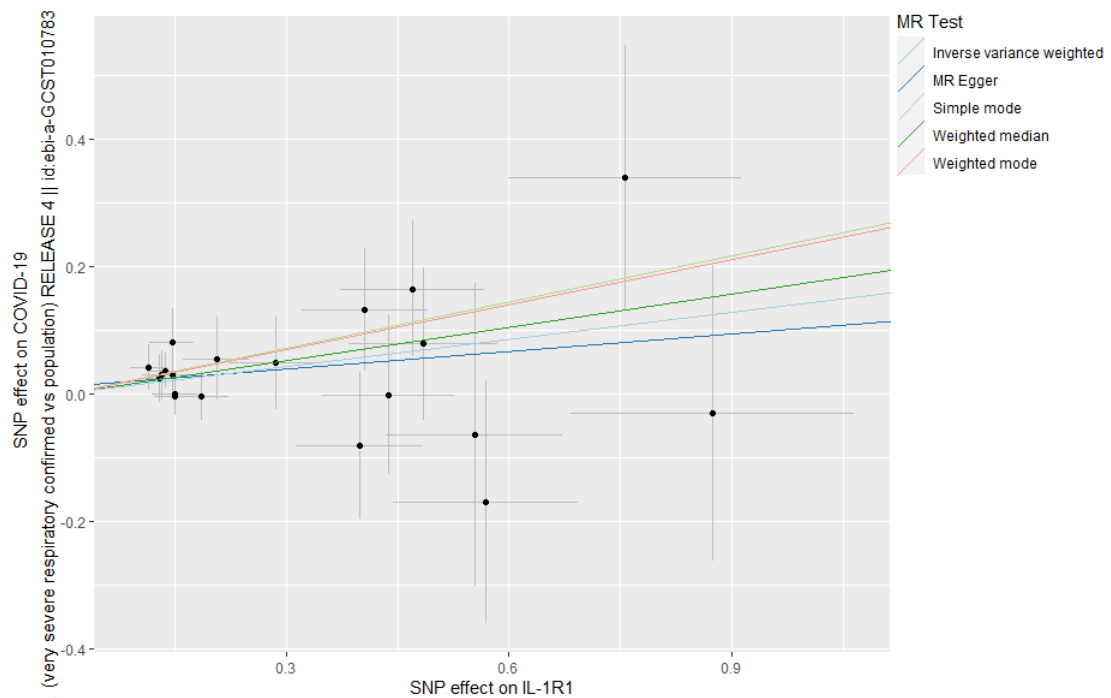
Supplementary figure 1.



Supplementary figure 1. The design flow chart for the Mendelian randomization

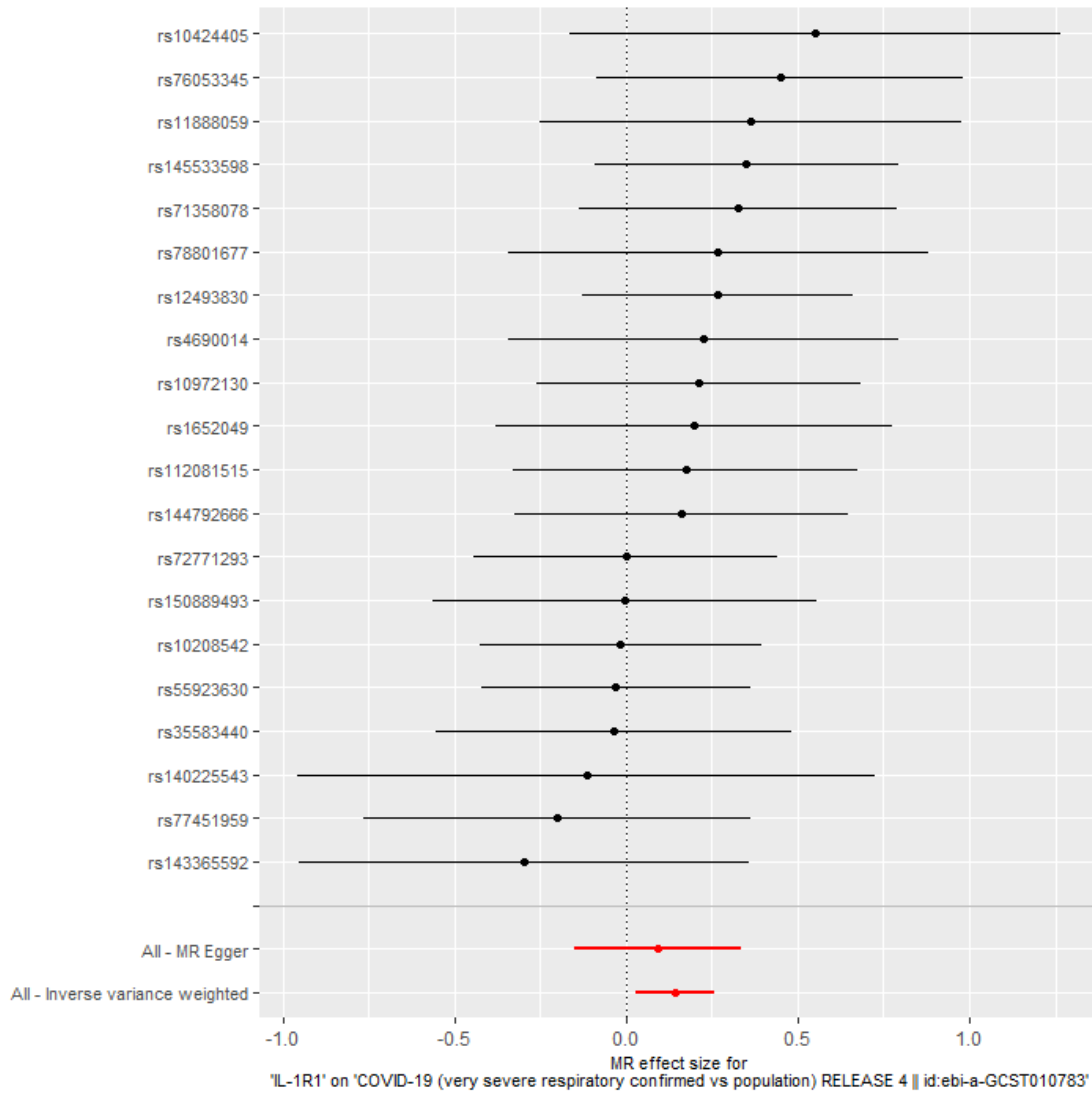
(MR) study. MR assumptions: assumption 1, 2 and 3. Solid line represents direct causal effects that IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp genetic instrumental variants are reliably associated with IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp levels, respectively and influence the risk of Corona Virus Disease 2019 (COVID-19) through the IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp, respectively, in assumption 1. Dotted line represents that IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp genetic instrumental variants are not associated with any measured and unmeasured confounders and do not influence the risk of COVID-19 through other pathways in assumption 2 and 3, respectively.

Supplementary figure 2.



Supplementary figure 2. Individual estimates about the causal effect of IL-1R1 on COVID-19. The x-axis shows the SNP (single nucleotide polymorphism) effect and SE (standard error) on each of IL-1R1. The y-axis shows the SNP effect and SE on COVID-19 by analyzing GWAS ID: ebi-a-GCST010783 (very severe respiratory confirmed vs population). The regression line for MR Egger, Weighted median, IVW, Simple mode, and Weighted mode is shown.

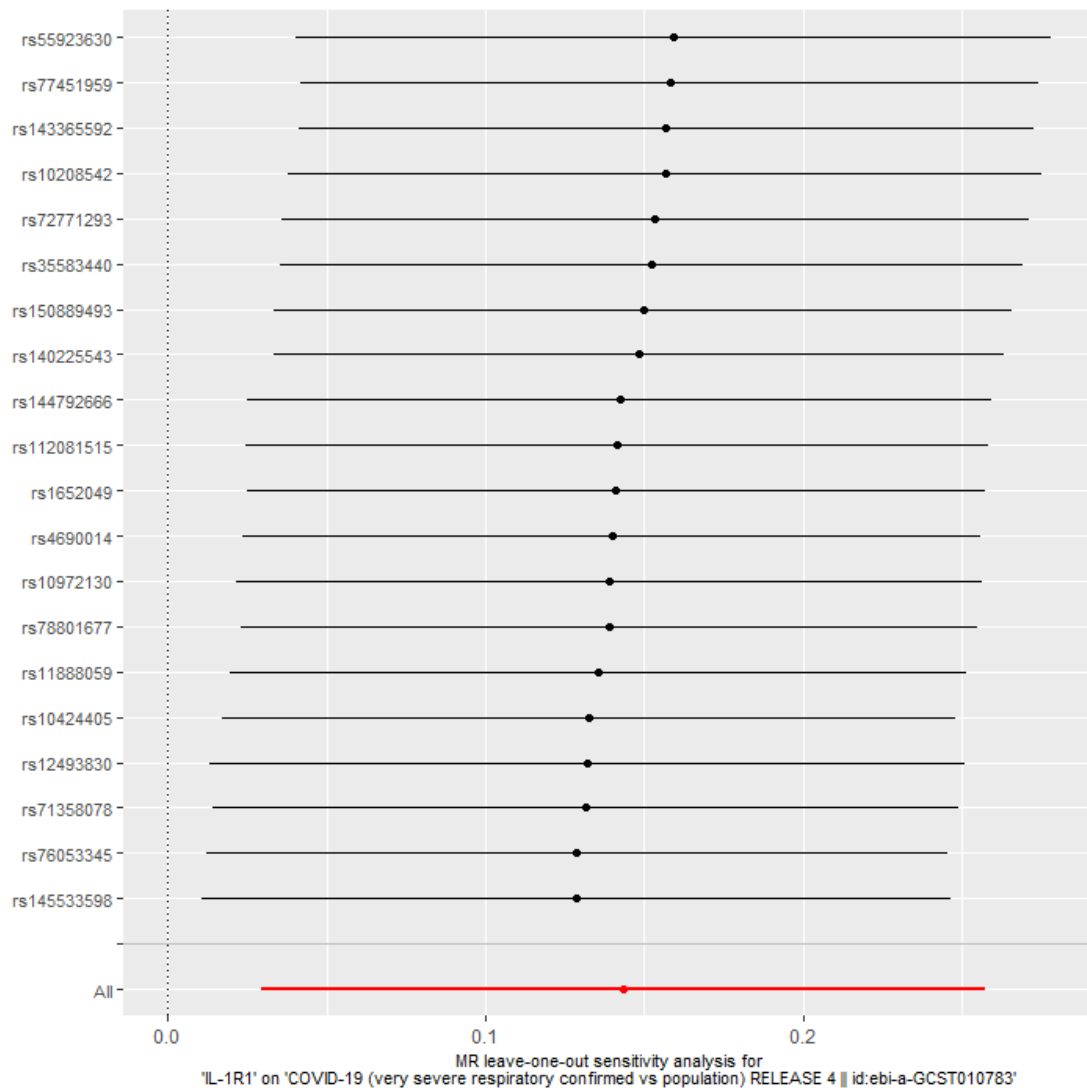
Supplementary figure 3.



Supplementary figure 3. Forest plot of IL-1R1 associated with risk of COVID-19.

The x-axis shows MR effect size for IL-1R1 on COVID-19 by analyzing GWAS ID: ebi-a-GCST010783 (very severe respiratory confirmed vs population). The y-axis shows the analysis for each of SNPs.

Supplementary figure 4.



Supplementary figure 4. MR leave-one-out sensitivity analysis for the effect of IL-1R1 SNPs on COVID-19. The x-axis shows MR leave-one-out sensitivity analysis for IL-1R1 on COVID-19 by analyzing GWAS ID: ebi-a-GCST010783 (very severe respiratory confirmed vs population). The y-axis shows the analysis for the effect of leave-one-out of SNPs on COVID-19.

Supplementary table 1. IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp genetic instrumental variants (IVs).

Exposure	SNP	Beta	SE	EA	NEA	EAF	p val	Gene
IL-1A	rs13170249	0.353	0.077	T	C	0.03	4.37E-06	PAM
IL-1A	rs78084692	0.874	0.182	G	A	0.01	1.51E-06	AC004540.5
IL-1A	rs7148359	-0.135	0.029	T	C	0.27	2.14E-06	AL445384.1
IL-1A	rs4867603	-0.131	0.027	C	T	0.55	1.05E-06	KCNIP1
IL-1A	rs10418046	0.153	0.030	G	T	0.22	3.09E-07	NLRP12
IL-1A	rs6549977	-0.120	0.026	A	G	0.38	3.89E-06	AC137674.2
IL-1A	rs8107140	0.115	0.025	A	C	0.59	3.63E-06	ZIM2
IL-1A	rs28895307	0.365	0.072	G	T	0.03	4.27E-07	OTUD7A
IL-1A	rs60010354	0.114	0.025	G	A	0.42	4.17E-06	DDX10
IL-1A	rs41386050	0.318	0.069	T	G	0.03	4.79E-06	LIPG
IL-1A	rs2734376	0.140	0.028	T	C	0.69	4.90E-07	FHIT
IL-1A	rs2749675	-0.172	0.037	G	A	0.87	4.37E-06	ZDHC14
IL-1A	rs77869959	-0.240	0.052	A	C	0.06	4.47E-06	RP11-313L9.1
IL-1A	rs6033085	-0.180	0.039	T	A	0.89	4.90E-06	RP4-734C18.1
IL-1A	rs72647025	0.134	0.028	G	C	0.39	1.86E-06	VWA5B1
IL-1A	rs11922046	0.139	0.029	T	C	0.26	1.20E-06	RN5S142
IL-1A	rs139711468	-0.347	0.073	A	G	0.03	1.86E-06	RP11-405A12.2
IL-1A	rs4757027	-0.143	0.028	C	G	0.68	3.39E-07	NAV2
IL-1A	rs192996211	-0.469	0.103	A	G	0.02	4.90E-06	RP11-428L9.1
IL-1B	rs115242021	0.233	0.041	A	C	0.09	5.07E-07	U1
IL-1B	rs143319329	0.280	0.072	T	C	0.03	2.00E-06	METTL2B
IL-1B	rs61335305	0.297	0.072	A	C	0.01	1.90E-06	MEGF11
IL-1B	rs4786740	0.085	0.020	A	C	0.41	4.67E-06	RP11-420N3.2
IL-1B	rs62015704	0.108	0.028	A	G	0.88	2.09E-06	RBFOX1
IL-1B	rs9898641	0.203	0.045	C	T	0.1	3.59E-06	RP11-567L7.6
IL-1B	rs1942793	0.072	0.019	T	G	0.47	4.98E-06	CTD-2354A18.1
IL-1R1	rs1652049	0.128	0.027	G	T	0.46	1.62E-06	RNASE11
IL-1R1	rs71358078	-0.406	0.085	T	C	0.02	1.86E-06	RP11-172F10.1
IL-1R1	rs112081515	-0.286	0.062	G	A	0.04	3.55E-06	DYM
IL-1R1	rs10208542	-0.185	0.037	C	G	0.13	6.46E-07	IL1R1
IL-1R1	rs77451959	0.400	0.086	T	G	0.02	2.95E-06	MPPED2
IL-1R1	rs12493830	-0.138	0.025	C	T	0.45	6.17E-08	ST3GAL6-AS1
IL-1R1	rs76053345	-0.756	0.157	C	T	0.01	1.38E-06	EPB41L4B
IL-1R1	rs140225543	-0.553	0.119	A	G	0.01	3.47E-06	TMEM86B
IL-1R1	rs72771293	0.150	0.031	G	A	0.22	1.17E-06	HAPLN1
IL-1R1	rs78801677	0.206	0.044	A	G	0.09	3.31E-06	LRRTM3
IL-1R1	rs144792666	0.485	0.101	T	C	0.02	1.45E-06	YIPF7
IL-1R1	rs35583440	0.874	0.191	C	G	0.01	4.79E-06	RCN1
IL-1R1	rs11888059	0.115	0.025	G	A	0.5	3.02E-06	IL1R1
IL-1R1	rs145533598	0.470	0.097	A	G	0.02	1.12E-06	PPARG
IL-1R1	rs10424405	-0.146	0.030	G	A	0.22	1.32E-06	NLRP12
IL-1R1	rs55923630	-0.149	0.026	T	A	0.3	1.35E-08	IGSF11
IL-1R1	rs150889493	-0.438	0.089	C	T	0.02	7.94E-07	RP11-420N3.2
IL-1R1	rs4690014	0.133	0.026	A	G	0.39	2.69E-07	RGS12
IL-1R1	rs143365592	0.568	0.124	G	T	0.01	4.79E-06	KRTAP13-1
IL-1R1	rs10972130	0.147	0.031	A	G	0.2	2.29E-06	DNAI1
IL-1Ra	rs12121840	0.269	0.057	T	C	0.08	2.43E-06	RP11-280O1.2
IL-1Ra	rs56134659	0.112	0.024	G	A	0.4	2.44E-06	H1FX-AS1
IL-1Ra	rs1054402	0.131	0.027	T	C	0.26	1.13E-06	PAPPA
IL-1Ra	rs2809154	0.179	0.039	C	T	0.82	3.74E-06	SLITRK1
IL-1Ra	rs11627423	0.117	0.025	A	C	0.61	2.12E-06	AKAP6
IL-1Ra	rs61335305	0.445	0.091	A	C	0.01	1.00E-06	MEGF11
IL-1Ra	rs11869294	0.227	0.047	G	C	0.03	1.50E-06	CRHR1
IL-1Ra	rs147747784	0.365	0.076	C	G	0.02	1.46E-06	LILRB1

IL-1Ra	rs187166731	0.233	0.051	C	T	0.98	4.70E-06	Z83851.3
IL-1Ra	rs9623661	0.197	0.043	C	T	0.91	3.86E-06	A4GALT
IL-1Racp	rs1988743	0.135	0.028	C	G	0.72	1.51E-06	IL1RAP
IL-1Racp	rs2885370	-0.209	0.031	T	G	0.74	8.91E-12	IL1RAP
IL-1Racp	rs41268633	-0.251	0.050	G	A	0.09	6.03E-07	IL1RAP
IL-1Racp	rs113553332	0.685	0.056	T	C	0.05	4.68E-34	GMNC
IL-1Racp	rs148996090	1.111	0.137	A	C	0.01	3.98E-16	IL1RAP
IL-1Racp	rs67249092	-0.379	0.035	G	A	0.15	5.25E-28	IL1RAP
IL-1Racp	rs6444435	-0.171	0.026	G	A	0.35	5.37E-11	IL1RAP
IL-1Racp	rs1024949	-0.448	0.040	C	T	0.13	2.95E-29	IL1RAP
IL-1Racp	rs3935774	0.741	0.112	T	C	0.02	3.55E-11	IL1RAP
IL-1Racp	rs13084868	0.178	0.039	C	T	0.12	4.27E-06	GMNC
IL-1Racp	rs1607264	-0.146	0.031	C	A	0.8	2.82E-06	ZNF385B
IL-1Racp	rs62196769	-0.205	0.044	C	A	0.09	3.09E-06	AC010983.1
IL-1Racp	rs4815878	0.155	0.030	A	G	0.76	1.95E-07	SRXN1
IL-1Racp	rs6461953	-0.146	0.032	C	T	0.18	4.68E-06	AC073316.1
IL-1Racp	rs4687154	-0.581	0.040	C	G	0.89	7.94E-49	IL1RAP
IL-1Racp	rs6796131	-0.244	0.026	C	G	0.33	1.51E-20	IL1RAP
IL-1Racp	rs4624606	-0.925	0.025	T	A	0.75	1.00E-200	IL1RAP
IL-1Racp	rs6444430	0.154	0.033	A	G	0.79	2.29E-06	IL1RAP
IL-1Racp	rs16969649	-0.140	0.031	T	C	0.21	4.90E-06	MIR4318
IL-1Racp	rs7623802	-0.129	0.027	T	C	0.66	1.41E-06	RP11-6B4.1
IL-1Racp	rs62116908	-0.294	0.064	C	T	0.04	3.72E-06	AC022311.1
IL-1Racp	rs61492644	-0.124	0.025	A	G	0.52	7.41E-07	GFOD1
IL-1Racp	rs6801017	-0.202	0.031	G	A	0.21	3.80E-11	IL1RAP
IL-1Racp	rs78129851	0.793	0.117	A	G	0.01	1.07E-11	IL1RAP
IL-1Racp	rs5753690	0.149	0.029	T	C	0.23	2.95E-07	SFI1
IL-1Racp	rs13059920	0.420	0.038	G	A	0.13	9.33E-29	IL1RAP
IL-1Racp	rs73754360	0.384	0.080	G	T	0.03	1.78E-06	RP11-127B16.1
IL-1Racp	rs75857200	0.199	0.042	A	C	0.1	2.34E-06	CTD-2591A6.2

SNP: single-nucleotide polymorphism. EA: effect allele. NEA: non-effect allele. EAF: effect allele frequency. Beta: the regression coefficient based on the IL-1A, IL-1B, IL-1R1, IL-1Ra, or IL-1Racp raising effect allele. SE: standard error.

Supplementary table 2. Association of IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp genetic IVs in COVID-19 GWAS dataset.

Exposure GWAS ID	SNP	Exposure GWAS			Outcome (COVID-19) GWAS		
		Beta	SE	<i>p</i> val	Beta	SE	<i>p</i> val
ebi-a-GCST010775	rs10418046	0.153	0.030	3.09E-07	-0.114	0.244	0.639
	rs11922046	0.139	0.029	1.20E-06	0.114	0.170	0.503
	rs13170249	0.353	0.077	4.37E-06	-0.043	0.480	0.928
	rs139711468	-0.347	0.073	1.86E-06	-0.790	0.865	0.362
	rs192996211	-0.469	0.103	4.90E-06	-0.414	0.702	0.556
	rs2734376	0.140	0.028	4.90E-07	0.006	0.163	0.970
	rs2749675	-0.172	0.037	4.37E-06	-0.614	0.333	0.065
	rs28895307	0.365	0.072	4.27E-07	0.167	0.405	0.679
	rs41386050	0.318	0.069	4.79E-06	-0.475	0.445	0.286
	rs4757027	-0.143	0.028	3.39E-07	-0.049	0.157	0.756
	rs4867603	-0.131	0.027	1.05E-06	0.125	0.148	0.398
	rs60010354	0.114	0.025	4.17E-06	-0.191	0.148	0.197
	rs6033085	-0.180	0.039	4.90E-06	0.223	0.214	0.298
	rs6549977	-0.120	0.026	3.89E-06	0.019	0.149	0.897
	rs7148359	-0.135	0.029	2.14E-06	-0.097	0.203	0.633
	rs72647025	0.134	0.028	1.86E-06	0.036	0.160	0.820
	rs77869959	-0.240	0.052	4.47E-06	1.317	0.392	0.001
rs8107140	0.115	0.025	3.63E-06	-0.294	0.154	0.056	
ebi-a-GCST010776	rs10418046	0.153	0.030	3.09E-07	0.035	0.022	0.110
	rs11922046	0.139	0.029	1.20E-06	-0.053	0.018	0.003
	rs13170249	0.353	0.077	4.37E-06	-0.061	0.044	0.170
	rs139711468	-0.347	0.073	1.86E-06	0.012	0.046	0.802
	rs192996211	-0.469	0.103	4.90E-06	0.001	0.075	0.985
	rs2734376	0.140	0.028	4.90E-07	0.016	0.017	0.329
	rs2749675	-0.172	0.037	4.37E-06	-0.020	0.031	0.510
	rs28895307	0.365	0.072	4.27E-07	-0.035	0.049	0.473
	rs41386050	0.318	0.069	4.79E-06	-0.010	0.040	0.795
	rs4757027	-0.143	0.028	3.39E-07	-0.013	0.018	0.462
	rs4867603	-0.131	0.027	1.05E-06	-0.011	0.017	0.536
	rs60010354	0.114	0.025	4.17E-06	0.010	0.015	0.494
	rs6033085	-0.180	0.039	4.90E-06	0.066	0.024	0.006
	rs6549977	-0.120	0.026	3.89E-06	-0.022	0.017	0.204
	rs7148359	-0.135	0.029	2.14E-06	-0.035	0.021	0.093
	rs72647025	0.134	0.028	1.86E-06	0.012	0.018	0.493
	rs77869959	-0.240	0.052	4.47E-06	-0.023	0.037	0.530
rs78084692	0.874	0.182	1.51E-06	-0.052	0.138	0.704	
rs8107140	0.115	0.025	3.63E-06	0.002	0.016	0.920	
ebi-a-GCST010777	rs10418046	0.153	0.030	3.09E-07	0.018	0.020	0.349
	rs11922046	0.139	0.029	1.20E-06	-0.044	0.019	0.022
	rs13170249	0.353	0.077	4.37E-06	-0.095	0.049	0.053
	rs139711468	-0.347	0.073	1.86E-06	0.053	0.052	0.310
	rs192996211	-0.469	0.103	4.90E-06	0.053	0.086	0.539
	rs2734376	0.140	0.028	4.90E-07	0.028	0.018	0.124
	rs2749675	-0.172	0.037	4.37E-06	-0.007	0.032	0.826
	rs28895307	0.365	0.072	4.27E-07	-0.069	0.059	0.243
	rs41386050	0.318	0.069	4.79E-06	0.013	0.047	0.786
	rs4757027	-0.143	0.028	3.39E-07	-0.014	0.018	0.452
	rs4867603	-0.131	0.027	1.05E-06	-0.023	0.019	0.222
	rs60010354	0.114	0.025	4.17E-06	0.018	0.016	0.265
	rs6033085	-0.180	0.039	4.90E-06	0.043	0.024	0.072
	rs6549977	-0.120	0.026	3.89E-06	0.007	0.019	0.724

		rs7148359	-0.135	0.029	2.14E-06	-0.017	0.023	0.476
		rs72647025	0.134	0.028	1.86E-06	0.016	0.020	0.421
		rs77869959	-0.240	0.052	4.47E-06	0.012	0.040	0.758
		rs78084692	0.874	0.182	1.51E-06	-0.032	0.197	0.873
		rs8107140	0.115	0.025	3.63E-06	0.014	0.016	0.388
		rs10418046	0.153	0.030	3.09E-07	0.018	0.020	0.349
		rs11922046	0.139	0.029	1.20E-06	-0.044	0.019	0.022
		rs13170249	0.353	0.077	4.37E-06	-0.095	0.049	0.053
		rs139711468	-0.347	0.073	1.86E-06	0.053	0.052	0.310
		rs192996211	-0.469	0.103	4.90E-06	0.053	0.086	0.539
		rs2734376	0.140	0.028	4.90E-07	0.028	0.018	0.124
		rs2749675	-0.172	0.037	4.37E-06	-0.007	0.032	0.826
		rs28895307	0.365	0.072	4.27E-07	-0.069	0.059	0.243
		rs41386050	0.318	0.069	4.79E-06	0.013	0.047	0.786
	ebi-a-GCST010778	rs4757027	-0.143	0.028	3.39E-07	-0.014	0.018	0.452
		rs4867603	-0.131	0.027	1.05E-06	-0.023	0.019	0.222
		rs60010354	0.114	0.025	4.17E-06	0.018	0.016	0.265
		rs6033085	-0.180	0.039	4.90E-06	0.043	0.024	0.072
		rs6549977	-0.120	0.026	3.89E-06	0.007	0.019	0.724
		rs7148359	-0.135	0.029	2.14E-06	-0.017	0.023	0.476
		rs72647025	0.134	0.028	1.86E-06	0.016	0.020	0.421
		rs77869959	-0.240	0.052	4.47E-06	0.012	0.040	0.758
		rs78084692	0.874	0.182	1.51E-06	-0.032	0.197	0.873
		rs8107140	0.115	0.025	3.63E-06	0.014	0.016	0.388
		rs10418046	0.153	0.030	3.09E-07	0.040	0.031	0.202
		rs11922046	0.139	0.029	1.20E-06	-0.078	0.029	0.007
		rs13170249	0.353	0.077	4.37E-06	-0.007	0.064	0.913
		rs139711468	-0.347	0.073	1.86E-06	0.008	0.076	0.917
		rs192996211	-0.469	0.103	4.90E-06	0.088	0.112	0.432
		rs2734376	0.140	0.028	4.90E-07	0.018	0.023	0.430
		rs2749675	-0.172	0.037	4.37E-06	-0.010	0.039	0.798
		rs28895307	0.365	0.072	4.27E-07	0.065	0.074	0.380
		rs41386050	0.318	0.069	4.79E-06	-0.129	0.066	0.050
	ebi-a-GCST010779	rs4757027	-0.143	0.028	3.39E-07	-0.025	0.027	0.358
		rs4867603	-0.131	0.027	1.05E-06	-0.007	0.028	0.802
		rs60010354	0.114	0.025	4.17E-06	0.020	0.021	0.354
		rs6033085	-0.180	0.039	4.90E-06	0.054	0.038	0.158
		rs6549977	-0.120	0.026	3.89E-06	-0.047	0.027	0.085
		rs7148359	-0.135	0.029	2.14E-06	-0.033	0.030	0.270
		rs72647025	0.134	0.028	1.86E-06	-0.021	0.029	0.479
		rs77869959	-0.240	0.052	4.47E-06	0.036	0.059	0.550
		rs78084692	0.874	0.182	1.51E-06	-0.046	0.174	0.793
		rs8107140	0.115	0.025	3.63E-06	-0.015	0.024	0.518
		rs10418046	0.153	0.030	3.09E-07	0.026	0.018	0.139
		rs11922046	0.139	0.029	1.20E-06	-0.055	0.016	0.001
		rs13170249	0.353	0.077	4.37E-06	-0.024	0.040	0.545
		rs139711468	-0.347	0.073	1.86E-06	0.025	0.045	0.585
		rs192996211	-0.469	0.103	4.90E-06	0.012	0.065	0.859
		rs2734376	0.140	0.028	4.90E-07	0.023	0.014	0.104
		rs2749675	-0.172	0.037	4.37E-06	-0.012	0.026	0.638
		rs28895307	0.365	0.072	4.27E-07	-0.020	0.048	0.671
		rs41386050	0.318	0.069	4.79E-06	-0.012	0.039	0.754
	ebi-a-GCST010780	rs4757027	-0.143	0.028	3.39E-07	-0.012	0.015	0.439
		rs4867603	-0.131	0.027	1.05E-06	-0.026	0.016	0.091
		rs60010354	0.114	0.025	4.17E-06	0.012	0.013	0.354
		rs6033085	-0.180	0.039	4.90E-06	0.054	0.021	0.009
		rs6549977	-0.120	0.026	3.89E-06	-0.014	0.015	0.375
		rs7148359	-0.135	0.029	2.14E-06	-0.023	0.018	0.210

	rs72647025	0.134	0.028	1.86E-06	-0.001	0.017	0.940
	rs77869959	-0.240	0.052	4.47E-06	-0.019	0.033	0.572
	rs78084692	0.874	0.182	1.51E-06	0.046	0.127	0.714
	rs8107140	0.115	0.025	3.63E-06	0.002	0.013	0.886
	rs10418046	0.153	0.030	3.09E-07	-0.001	0.039	0.979
	rs11922046	0.139	0.029	1.20E-06	-0.017	0.031	0.586
	rs13170249	0.353	0.077	4.37E-06	0.080	0.080	0.321
	rs139711468	-0.347	0.073	1.86E-06	0.090	0.087	0.301
	rs192996211	-0.469	0.103	4.90E-06	-0.138	0.156	0.375
	rs2734376	0.140	0.028	4.90E-07	0.027	0.030	0.358
	rs2749675	-0.172	0.037	4.37E-06	0.075	0.044	0.088
	rs28895307	0.365	0.072	4.27E-07	0.039	0.088	0.656
	rs41386050	0.318	0.069	4.79E-06	-0.068	0.071	0.343
ebi-a-GCST010781	rs4757027	-0.143	0.028	3.39E-07	0.033	0.033	0.321
	rs4867603	-0.131	0.027	1.05E-06	-0.043	0.030	0.151
	rs60010354	0.114	0.025	4.17E-06	-0.034	0.030	0.265
	rs6033085	-0.180	0.039	4.90E-06	-0.006	0.044	0.891
	rs6549977	-0.120	0.026	3.89E-06	-0.016	0.028	0.574
	rs7148359	-0.135	0.029	2.14E-06	0.012	0.031	0.687
	rs72647025	0.134	0.028	1.86E-06	0.026	0.029	0.361
	rs77869959	-0.240	0.052	4.47E-06	0.012	0.069	0.857
	rs78084692	0.874	0.182	1.51E-06	-0.350	0.258	0.175
	rs8107140	0.115	0.025	3.63E-06	0.001	0.027	0.971
	rs10418046	0.153	0.030	3.09E-07	0.037	0.049	0.449
	rs11922046	0.139	0.029	1.20E-06	-0.030	0.050	0.546
	rs13170249	0.353	0.077	4.37E-06	-0.139	0.133	0.298
	rs139711468	-0.347	0.073	1.86E-06	0.166	0.142	0.243
	rs192996211	-0.469	0.103	4.90E-06	0.186	0.221	0.400
	rs2734376	0.140	0.028	4.90E-07	-0.015	0.047	0.757
	rs2749675	-0.172	0.037	4.37E-06	0.131	0.077	0.091
	rs28895307	0.365	0.072	4.27E-07	-0.191	0.159	0.232
	rs41386050	0.318	0.069	4.79E-06	-0.345	0.138	0.012
ebi-a-GCST010782	rs4757027	-0.143	0.028	3.39E-07	0.032	0.047	0.493
	rs4867603	-0.131	0.027	1.05E-06	-0.011	0.052	0.838
	rs60010354	0.114	0.025	4.17E-06	0.062	0.052	0.236
	rs6033085	-0.180	0.039	4.90E-06	-0.036	0.060	0.553
	rs6549977	-0.120	0.026	3.89E-06	-0.039	0.054	0.464
	rs7148359	-0.135	0.029	2.14E-06	-0.036	0.062	0.563
	rs72647025	0.134	0.028	1.86E-06	0.006	0.056	0.914
	rs77869959	-0.240	0.052	4.47E-06	0.152	0.102	0.136
	rs78084692	0.874	0.182	1.51E-06	0.160	0.551	0.772
	rs8107140	0.115	0.025	3.63E-06	-0.051	0.041	0.212
	rs10418046	0.153	0.030	3.09E-07	-0.078	0.053	0.139
	rs11922046	0.139	0.029	1.20E-06	-0.091	0.043	0.036
	rs13170249	0.353	0.077	4.37E-06	0.117	0.083	0.162
	rs139711468	-0.347	0.073	1.86E-06	0.052	0.095	0.584
	rs192996211	-0.469	0.103	4.90E-06	0.272	0.150	0.070
	rs2734376	0.140	0.028	4.90E-07	-0.001	0.030	0.981
	rs2749675	-0.172	0.037	4.37E-06	-0.039	0.056	0.481
	rs28895307	0.365	0.072	4.27E-07	-0.013	0.091	0.889
	rs41386050	0.318	0.069	4.79E-06	-0.185	0.083	0.025
ebi-a-GCST010783	rs4757027	-0.143	0.028	3.39E-07	-0.037	0.041	0.374
	rs4867603	-0.131	0.027	1.05E-06	-0.011	0.039	0.787
	rs60010354	0.114	0.025	4.17E-06	0.017	0.028	0.543
	rs6033085	-0.180	0.039	4.90E-06	0.056	0.060	0.349
	rs6549977	-0.120	0.026	3.89E-06	-0.060	0.038	0.117
	rs7148359	-0.135	0.029	2.14E-06	0.014	0.040	0.737
	rs72647025	0.134	0.028	1.86E-06	-0.001	0.041	0.989

	rs77869959	-0.240	0.052	4.47E-06	-0.028	0.089	0.754
	rs78084692	0.874	0.182	1.51E-06	0.095	0.213	0.657
	rs8107140	0.115	0.025	3.63E-06	0.018	0.037	0.629
	rs143319329	0.280	0.072	2.00E-06	-1.066	0.437	0.015
	rs1942793	0.072	0.019	4.98E-06	-0.190	0.158	0.230
ebi-a-GCST010775	rs4786740	0.085	0.020	4.67E-06	0.007	0.160	0.964
	rs61335305	0.297	0.072	1.90E-06	-0.369	0.451	0.413
	rs62015704	0.108	0.028	2.09E-06	-0.264	0.232	0.255
	rs9898641	0.203	0.045	3.59E-06	0.032	0.237	0.891
	rs143319329	0.280	0.072	2.00E-06	0.002	0.048	0.960
	rs1942793	0.072	0.019	4.98E-06	-0.013	0.014	0.382
ebi-a-GCST010776	rs4786740	0.085	0.020	4.67E-06	-0.005	0.016	0.767
	rs61335305	0.297	0.072	1.90E-06	0.059	0.060	0.332
	rs62015704	0.108	0.028	2.09E-06	-0.004	0.022	0.863
	rs9898641	0.203	0.045	3.59E-06	-0.023	0.021	0.271
	rs143319329	0.280	0.072	2.00E-06	0.055	0.048	0.252
	rs1942793	0.072	0.019	4.98E-06	0.009	0.015	0.555
ebi-a-GCST010777	rs4786740	0.085	0.020	4.67E-06	0.011	0.017	0.537
	rs61335305	0.297	0.072	1.90E-06	0.028	0.042	0.504
	rs62015704	0.108	0.028	2.09E-06	-0.025	0.024	0.304
	rs9898641	0.203	0.045	3.59E-06	0.012	0.022	0.595
	rs143319329	0.280	0.072	2.00E-06	0.055	0.048	0.252
	rs1942793	0.072	0.019	4.98E-06	0.009	0.015	0.555
ebi-a-GCST010778	rs4786740	0.085	0.020	4.67E-06	0.011	0.017	0.537
	rs61335305	0.297	0.072	1.90E-06	0.028	0.042	0.504
	rs62015704	0.108	0.028	2.09E-06	-0.025	0.024	0.304
	rs9898641	0.203	0.045	3.59E-06	0.012	0.022	0.595
	rs143319329	0.280	0.072	2.00E-06	0.087	0.065	0.180
	rs1942793	0.072	0.019	4.98E-06	-0.039	0.021	0.062
IL-1B ebi-a-GCST010779	rs4786740	0.085	0.020	4.67E-06	-0.018	0.026	0.493
	rs61335305	0.297	0.072	1.90E-06	0.074	0.067	0.268
	rs62015704	0.108	0.028	2.09E-06	0.045	0.032	0.158
	rs9898641	0.203	0.045	3.59E-06	0.024	0.037	0.521
	rs143319329	0.280	0.072	2.00E-06	0.039	0.038	0.305
	rs1942793	0.072	0.019	4.98E-06	-0.019	0.013	0.134
ebi-a-GCST010780	rs4786740	0.085	0.020	4.67E-06	0.001	0.015	0.941
	rs61335305	0.297	0.072	1.90E-06	0.017	0.038	0.653
	rs62015704	0.108	0.028	2.09E-06	0.010	0.019	0.608
	rs9898641	0.203	0.045	3.59E-06	0.003	0.020	0.894
	rs143319329	0.280	0.072	2.00E-06	0.036	0.093	0.698
	rs1942793	0.072	0.019	4.98E-06	-0.022	0.026	0.407
ebi-a-GCST010781	rs4786740	0.085	0.020	4.67E-06	-0.021	0.031	0.495
	rs61335305	0.297	0.072	1.90E-06	0.124	0.170	0.466
	rs62015704	0.108	0.028	2.09E-06	0.010	0.044	0.816
	rs9898641	0.203	0.045	3.59E-06	-0.049	0.028	0.087
	rs143319329	0.280	0.072	2.00E-06	-0.099	0.116	0.391
	rs1942793	0.072	0.019	4.98E-06	-0.015	0.040	0.707
ebi-a-GCST010782	rs4786740	0.085	0.020	4.67E-06	-0.094	0.044	0.033
	rs61335305	0.297	0.072	1.90E-06	-0.037	0.101	0.715
	rs62015704	0.108	0.028	2.09E-06	0.009	0.062	0.888
	rs9898641	0.203	0.045	3.59E-06	0.047	0.050	0.343
	rs143319329	0.280	0.072	2.00E-06	0.028	0.101	0.784
	rs1942793	0.072	0.019	4.98E-06	-0.032	0.027	0.243
ebi-a-GCST010783	rs4786740	0.085	0.020	4.67E-06	-0.012	0.040	0.756
	rs61335305	0.297	0.072	1.90E-06	0.114	0.114	0.319
	rs62015704	0.108	0.028	2.09E-06	0.054	0.042	0.202
	rs9898641	0.203	0.045	3.59E-06	-0.036	0.038	0.343
	rs10208542	-0.185	0.037	6.46E-07	-0.095	0.229	0.678

	rs10424405	-0.146	0.030	1.32E-06	-0.135	0.236	0.567
	rs10972130	0.147	0.031	2.29E-06	-0.085	0.180	0.638
	rs112081515	-0.286	0.062	3.55E-06	-0.371	0.366	0.311
	rs11888059	0.115	0.025	3.02E-06	-0.115	0.152	0.450
	rs12493830	-0.138	0.025	6.17E-08	0.166	0.162	0.305
	rs140225543	-0.553	0.119	3.47E-06	0.942	1.019	0.355
	rs144792666	0.485	0.101	1.45E-06	-0.443	0.817	0.588
ebi-a-GCST010775	rs145533598	0.470	0.097	1.12E-06	0.172	0.614	0.780
	rs150889493	-0.438	0.089	7.94E-07	0.437	0.448	0.329
	rs1652049	0.128	0.027	1.62E-06	0.140	0.152	0.357
	rs35583440	0.874	0.191	4.79E-06	-0.337	0.945	0.721
	rs4690014	0.133	0.026	2.69E-07	0.167	0.158	0.289
	rs55923630	-0.149	0.026	1.35E-08	-0.017	0.171	0.922
	rs71358078	-0.406	0.085	1.86E-06	0.001	0.746	0.999
	rs72771293	0.150	0.031	1.17E-06	0.046	0.172	0.787
	rs77451959	0.400	0.086	2.95E-06	0.546	0.502	0.277
	rs78801677	0.206	0.044	3.31E-06	0.023	0.302	0.940
	rs10208542	-0.185	0.037	6.46E-07	-0.003	0.022	0.899
	rs10424405	-0.146	0.030	1.32E-06	0.025	0.022	0.263
	rs10972130	0.147	0.031	2.29E-06	0.027	0.018	0.135
	rs112081515	-0.286	0.062	3.55E-06	0.054	0.036	0.132
	rs11888059	0.115	0.025	3.02E-06	0.005	0.015	0.766
	rs12493830	-0.138	0.025	6.17E-08	-0.013	0.014	0.371
	rs140225543	-0.553	0.119	3.47E-06	0.009	0.089	0.918
	rs143365592	0.568	0.124	4.79E-06	-0.150	0.087	0.086
	rs144792666	0.485	0.101	1.45E-06	-0.010	0.064	0.879
ebi-a-GCST010776	rs145533598	0.470	0.097	1.12E-06	-0.013	0.059	0.819
	rs150889493	-0.438	0.089	7.94E-07	0.090	0.053	0.093
	rs1652049	0.128	0.027	1.62E-06	-0.003	0.016	0.830
	rs35583440	0.874	0.191	4.79E-06	-0.121	0.108	0.263
	rs4690014	0.133	0.026	2.69E-07	0.011	0.016	0.490
	rs55923630	-0.149	0.026	1.35E-08	0.013	0.016	0.406
	rs71358078	-0.406	0.085	1.86E-06	-0.006	0.053	0.908
	rs72771293	0.150	0.031	1.17E-06	-0.014	0.018	0.415
	rs76053345	-0.756	0.157	1.38E-06	-0.078	0.112	0.489
	rs77451959	0.400	0.086	2.95E-06	-0.071	0.052	0.174
	rs78801677	0.206	0.044	3.31E-06	-0.010	0.030	0.751
	rs10208542	-0.185	0.037	6.46E-07	-0.001	0.021	0.954
	rs10424405	-0.146	0.030	1.32E-06	0.017	0.020	0.399
	rs10972130	0.147	0.031	2.29E-06	0.029	0.020	0.152
	rs112081515	-0.286	0.062	3.55E-06	0.034	0.042	0.419
	rs11888059	0.115	0.025	3.02E-06	-0.002	0.016	0.884
	rs12493830	-0.138	0.025	6.17E-08	0.002	0.016	0.922
	rs140225543	-0.553	0.119	3.47E-06	-0.019	0.097	0.848
	rs143365592	0.568	0.124	4.79E-06	-0.096	0.092	0.298
	rs144792666	0.485	0.101	1.45E-06	0.019	0.076	0.805
	rs145533598	0.470	0.097	1.12E-06	-0.092	0.072	0.201
ebi-a-GCST010777	rs150889493	-0.438	0.089	7.94E-07	0.054	0.059	0.359
	rs1652049	0.128	0.027	1.62E-06	-0.005	0.016	0.759
	rs35583440	0.874	0.191	4.79E-06	-0.187	0.124	0.133
	rs4690014	0.133	0.026	2.69E-07	0.018	0.017	0.285
	rs55923630	-0.149	0.026	1.35E-08	0.023	0.018	0.190
	rs71358078	-0.406	0.085	1.86E-06	-0.018	0.061	0.774
	rs72771293	0.150	0.031	1.17E-06	-0.007	0.020	0.748
	rs76053345	-0.756	0.157	1.38E-06	-0.133	0.116	0.252
	rs77451959	0.400	0.086	2.95E-06	-0.095	0.058	0.101
	rs78801677	0.206	0.044	3.31E-06	-0.031	0.033	0.349
	rs10208542	-0.185	0.037	6.46E-07	-0.001	0.021	0.954

	rs10424405	-0.146	0.030	1.32E-06	0.017	0.020	0.399
	rs10972130	0.147	0.031	2.29E-06	0.029	0.020	0.152
	rs112081515	-0.286	0.062	3.55E-06	0.034	0.042	0.419
	rs11888059	0.115	0.025	3.02E-06	-0.002	0.016	0.884
	rs12493830	-0.138	0.025	6.17E-08	0.002	0.016	0.922
	rs140225543	-0.553	0.119	3.47E-06	-0.019	0.097	0.848
	rs143365592	0.568	0.124	4.79E-06	-0.096	0.092	0.298
	rs144792666	0.485	0.101	1.45E-06	0.019	0.076	0.805
ebi-a-GCST010778	rs145533598	0.470	0.097	1.12E-06	-0.092	0.072	0.201
	rs150889493	-0.438	0.089	7.94E-07	0.054	0.059	0.359
	rs1652049	0.128	0.027	1.62E-06	-0.005	0.016	0.759
	rs35583440	0.874	0.191	4.79E-06	-0.187	0.124	0.133
	rs4690014	0.133	0.026	2.69E-07	0.018	0.017	0.285
	rs55923630	-0.149	0.026	1.35E-08	0.023	0.018	0.190
	rs71358078	-0.406	0.085	1.86E-06	-0.018	0.061	0.774
	rs72771293	0.150	0.031	1.17E-06	-0.007	0.020	0.748
	rs76053345	-0.756	0.157	1.38E-06	-0.133	0.116	0.252
	rs77451959	0.400	0.086	2.95E-06	-0.095	0.058	0.101
	rs78801677	0.206	0.044	3.31E-06	-0.031	0.033	0.349
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	rs10208542	-0.185	0.037	6.46E-07	0.008	0.028	0.771
	rs10424405	-0.146	0.030	1.32E-06	0.047	0.033	0.156
	rs10972130	0.147	0.031	2.29E-06	0.018	0.027	0.494
	rs112081515	-0.286	0.062	3.55E-06	-0.016	0.058	0.777
	rs11888059	0.115	0.025	3.02E-06	0.000	0.024	0.987
	rs12493830	-0.138	0.025	6.17E-08	-0.037	0.021	0.078
	rs140225543	-0.553	0.119	3.47E-06	0.053	0.159	0.737
	rs143365592	0.568	0.124	4.79E-06	-0.161	0.148	0.276
	rs144792666	0.485	0.101	1.45E-06	-0.002	0.098	0.980
ebi-a-GCST010779	rs145533598	0.470	0.097	1.12E-06	0.112	0.086	0.192
IL-1R1	rs150889493	-0.438	0.089	7.94E-07	-0.092	0.089	0.303
	rs1652049	0.128	0.027	1.62E-06	0.010	0.025	0.679
	rs35583440	0.874	0.191	4.79E-06	-0.066	0.189	0.728
	rs4690014	0.133	0.026	2.69E-07	0.008	0.026	0.762
	rs55923630	-0.149	0.026	1.35E-08	-0.002	0.023	0.936
	rs71358078	-0.406	0.085	1.86E-06	-0.074	0.076	0.332
	rs72771293	0.150	0.031	1.17E-06	-0.039	0.026	0.140
	rs76053345	-0.756	0.157	1.38E-06	-0.174	0.173	0.314
	rs77451959	0.400	0.086	2.95E-06	-0.094	0.084	0.264
	rs78801677	0.206	0.044	3.31E-06	0.006	0.047	0.898
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	rs10208542	-0.185	0.037	6.46E-07	-0.003	0.017	0.885
	rs10424405	-0.146	0.030	1.32E-06	0.017	0.018	0.353
	rs10972130	0.147	0.031	2.29E-06	0.024	0.016	0.147
	rs112081515	-0.286	0.062	3.55E-06	0.045	0.034	0.193
	rs11888059	0.115	0.025	3.02E-06	0.005	0.014	0.707
	rs12493830	-0.138	0.025	6.17E-08	-0.013	0.013	0.305
	rs140225543	-0.553	0.119	3.47E-06	0.000	0.087	0.996
	rs143365592	0.568	0.124	4.79E-06	-0.172	0.081	0.034
	rs144792666	0.485	0.101	1.45E-06	0.002	0.062	0.977
ebi-a-GCST010780	rs145533598	0.470	0.097	1.12E-06	0.003	0.055	0.958
	rs150889493	-0.438	0.089	7.94E-07	0.044	0.049	0.375
	rs1652049	0.128	0.027	1.62E-06	-0.001	0.014	0.953
	rs35583440	0.874	0.191	4.79E-06	-0.121	0.104	0.244
	rs4690014	0.133	0.026	2.69E-07	0.015	0.014	0.291
	rs55923630	-0.149	0.026	1.35E-08	0.008	0.014	0.562
	rs71358078	-0.406	0.085	1.86E-06	-0.003	0.049	0.946
	rs72771293	0.150	0.031	1.17E-06	-0.018	0.016	0.265
	rs76053345	-0.756	0.157	1.38E-06	-0.096	0.105	0.360
	rs77451959	0.400	0.086	2.95E-06	-0.077	0.049	0.117

	rs78801677	0.206	0.044	3.31E-06	-0.007	0.028	0.793
	rs10208542	-0.185	0.037	6.46E-07	-0.035	0.040	0.381
	rs10424405	-0.146	0.030	1.32E-06	-0.010	0.040	0.799
	rs10972130	0.147	0.031	2.29E-06	-0.075	0.033	0.022
	rs112081515	-0.286	0.062	3.55E-06	-0.063	0.065	0.331
	rs11888059	0.115	0.025	3.02E-06	-0.007	0.026	0.785
	rs12493830	-0.138	0.025	6.17E-08	0.005	0.027	0.847
	rs140225543	-0.553	0.119	3.47E-06	0.228	0.168	0.175
	rs143365592	0.568	0.124	4.79E-06	-0.166	0.147	0.259
	rs144792666	0.485	0.101	1.45E-06	0.062	0.118	0.602
ebi-a-GCST010781	rs145533598	0.470	0.097	1.12E-06	-0.050	0.107	0.644
	rs150889493	-0.438	0.089	7.94E-07	0.063	0.109	0.563
	rs1652049	0.128	0.027	1.62E-06	-0.025	0.029	0.379
	rs35583440	0.874	0.191	4.79E-06	0.319	0.157	0.042
	rs4690014	0.133	0.026	2.69E-07	0.022	0.029	0.439
	rs55923630	-0.149	0.026	1.35E-08	-0.011	0.029	0.709
	rs71358078	-0.406	0.085	1.86E-06	0.084	0.107	0.431
	rs72771293	0.150	0.031	1.17E-06	0.050	0.033	0.126
	rs76053345	-0.756	0.157	1.38E-06	-0.057	0.187	0.759
	rs77451959	0.400	0.086	2.95E-06	-0.016	0.103	0.876
	rs78801677	0.206	0.044	3.31E-06	-0.043	0.060	0.469
	rs10208542	-0.185	0.037	6.46E-07	-0.028	0.053	0.602
	rs10424405	-0.146	0.030	1.32E-06	0.041	0.054	0.439
	rs10972130	0.147	0.031	2.29E-06	-0.034	0.053	0.518
	rs112081515	-0.286	0.062	3.55E-06	-0.154	0.120	0.198
	rs11888059	0.115	0.025	3.02E-06	-0.008	0.042	0.853
	rs12493830	-0.138	0.025	6.17E-08	-0.017	0.041	0.678
	rs140225543	-0.553	0.119	3.47E-06	0.032	0.269	0.906
	rs143365592	0.568	0.124	4.79E-06	0.072	0.258	0.779
	rs144792666	0.485	0.101	1.45E-06	0.046	0.228	0.841
ebi-a-GCST010782	rs145533598	0.470	0.097	1.12E-06	-0.124	0.195	0.527
	rs150889493	-0.438	0.089	7.94E-07	-0.079	0.160	0.623
	rs1652049	0.128	0.027	1.62E-06	0.032	0.042	0.449
	rs35583440	0.874	0.191	4.79E-06	0.111	0.360	0.759
	rs4690014	0.133	0.026	2.69E-07	-0.013	0.043	0.756
	rs55923630	-0.149	0.026	1.35E-08	-0.041	0.046	0.376
	rs71358078	-0.406	0.085	1.86E-06	0.023	0.170	0.893
	rs72771293	0.150	0.031	1.17E-06	-0.114	0.054	0.034
	rs76053345	-0.756	0.157	1.38E-06	0.018	0.324	0.955
	rs77451959	0.400	0.086	2.95E-06	0.027	0.157	0.866
	rs78801677	0.206	0.044	3.31E-06	-0.001	0.089	0.992
	rs10208542	-0.185	0.037	6.46E-07	0.003	0.039	0.941
	rs10424405	-0.146	0.030	1.32E-06	-0.081	0.053	0.130
	rs10972130	0.147	0.031	2.29E-06	0.031	0.035	0.377
	rs112081515	-0.286	0.062	3.55E-06	-0.050	0.073	0.499
	rs11888059	0.115	0.025	3.02E-06	0.042	0.036	0.247
	rs12493830	-0.138	0.025	6.17E-08	-0.037	0.028	0.187
	rs140225543	-0.553	0.119	3.47E-06	0.064	0.237	0.787
	rs143365592	0.568	0.124	4.79E-06	-0.169	0.190	0.374
	rs144792666	0.485	0.101	1.45E-06	0.079	0.120	0.513
ebi-a-GCST010783	rs145533598	0.470	0.097	1.12E-06	0.165	0.106	0.120
	rs150889493	-0.438	0.089	7.94E-07	0.001	0.125	0.992
	rs1652049	0.128	0.027	1.62E-06	0.025	0.038	0.501
	rs35583440	0.874	0.191	4.79E-06	-0.030	0.231	0.895
	rs4690014	0.133	0.026	2.69E-07	0.030	0.038	0.434
	rs55923630	-0.149	0.026	1.35E-08	0.004	0.030	0.882
	rs71358078	-0.406	0.085	1.86E-06	-0.132	0.096	0.168
	rs72771293	0.150	0.031	1.17E-06	0.000	0.034	0.995

	rs76053345	-0.756	0.157	1.38E-06	-0.339	0.207	0.101
	rs77451959	0.400	0.086	2.95E-06	-0.081	0.115	0.481
	rs78801677	0.206	0.044	3.31E-06	0.055	0.064	0.390
	rs1054402	0.131	0.027	1.13E-06	-0.049	0.163	0.764
	rs11627423	0.117	0.025	2.12E-06	-0.114	0.151	0.453
	rs11869294	0.227	0.047	1.50E-06	0.285	0.328	0.386
	rs12121840	0.269	0.057	2.43E-06	0.249	0.301	0.408
ebi-a-GCST010775	rs187166731	0.233	0.051	4.70E-06	-0.016	0.552	0.977
	rs2809154	0.179	0.039	3.74E-06	-0.074	0.225	0.742
	rs56134659	0.112	0.024	2.44E-06	0.060	0.181	0.740
	rs61335305	0.445	0.091	1.00E-06	-0.369	0.451	0.413
	rs9623661	0.197	0.043	3.86E-06	0.449	0.243	0.064
	rs1054402	0.131	0.027	1.13E-06	-0.039	0.019	0.040
	rs11627423	0.117	0.025	2.12E-06	-0.008	0.017	0.619
	rs11869294	0.227	0.047	1.50E-06	0.033	0.046	0.477
	rs12121840	0.269	0.057	2.43E-06	-0.020	0.033	0.550
ebi-a-GCST010776	rs187166731	0.233	0.051	4.70E-06	0.022	0.098	0.827
	rs2809154	0.179	0.039	3.74E-06	0.020	0.021	0.325
	rs56134659	0.112	0.024	2.44E-06	-0.015	0.018	0.399
	rs61335305	0.445	0.091	1.00E-06	0.059	0.060	0.332
	rs9623661	0.197	0.043	3.86E-06	-0.017	0.028	0.540
	rs1054402	0.131	0.027	1.13E-06	-0.052	0.021	0.015
	rs11627423	0.117	0.025	2.12E-06	-0.038	0.018	0.033
	rs11869294	0.227	0.047	1.50E-06	-0.013	0.054	0.811
	rs12121840	0.269	0.057	2.43E-06	0.010	0.037	0.785
ebi-a-GCST010777	rs187166731	0.233	0.051	4.70E-06	0.179	0.127	0.159
	rs2809154	0.179	0.039	3.74E-06	0.006	0.024	0.788
	rs56134659	0.112	0.024	2.44E-06	0.033	0.019	0.077
	rs61335305	0.445	0.091	1.00E-06	0.028	0.042	0.504
	rs9623661	0.197	0.043	3.86E-06	-0.027	0.030	0.374
	rs1054402	0.131	0.027	1.13E-06	-0.052	0.021	0.015
	rs11627423	0.117	0.025	2.12E-06	-0.038	0.018	0.033
	rs11869294	0.227	0.047	1.50E-06	-0.013	0.054	0.811
	rs12121840	0.269	0.057	2.43E-06	0.010	0.037	0.785
ebi-a-GCST010778	rs187166731	0.233	0.051	4.70E-06	0.179	0.127	0.159
	rs2809154	0.179	0.039	3.74E-06	0.006	0.024	0.788
	rs56134659	0.112	0.024	2.44E-06	0.033	0.019	0.077
	rs61335305	0.445	0.091	1.00E-06	0.028	0.042	0.504
	rs9623661	0.197	0.043	3.86E-06	-0.027	0.030	0.374
	rs1054402	0.131	0.027	1.13E-06	0.021	0.029	0.474
	rs11627423	0.117	0.025	2.12E-06	-0.014	0.026	0.583
	rs11869294	0.227	0.047	1.50E-06	-0.024	0.073	0.744
	rs12121840	0.269	0.057	2.43E-06	0.015	0.051	0.767
ebi-a-GCST010779	rs187166731	0.233	0.051	4.70E-06	0.042	0.182	0.815
	rs2809154	0.179	0.039	3.74E-06	0.035	0.029	0.235
	rs56134659	0.112	0.024	2.44E-06	0.021	0.037	0.575
	rs61335305	0.445	0.091	1.00E-06	0.074	0.067	0.268
	rs9623661	0.197	0.043	3.86E-06	-0.087	0.045	0.053
IL-1Ra	rs1054402	0.131	0.027	1.13E-06	-0.025	0.017	0.140
	rs11627423	0.117	0.025	2.12E-06	-0.021	0.015	0.173
	rs11869294	0.227	0.047	1.50E-06	-0.022	0.040	0.587
	rs12121840	0.269	0.057	2.43E-06	0.005	0.031	0.883
ebi-a-GCST010780	rs187166731	0.233	0.051	4.70E-06	0.035	0.097	0.718
	rs2809154	0.179	0.039	3.74E-06	0.023	0.018	0.206
	rs56134659	0.112	0.024	2.44E-06	0.011	0.021	0.598
	rs61335305	0.445	0.091	1.00E-06	0.017	0.038	0.653
	rs9623661	0.197	0.043	3.86E-06	-0.025	0.026	0.322
	rs1054402	0.131	0.027	1.13E-06	-0.008	0.030	0.800

	rs11627423	0.117	0.025	2.12E-06	0.005	0.027	0.856
	rs11869294	0.227	0.047	1.50E-06	0.134	0.092	0.146
	rs12121840	0.269	0.057	2.43E-06	-0.014	0.058	0.812
ebi-a-GCST010781	rs147747784	0.365	0.076	1.46E-06	-0.255	0.146	0.079
	rs187166731	0.233	0.051	4.70E-06	0.191	0.151	0.205
	rs2809154	0.179	0.039	3.74E-06	-0.024	0.036	0.503
	rs56134659	0.112	0.024	2.44E-06	0.071	0.038	0.062
	rs61335305	0.445	0.091	1.00E-06	0.124	0.170	0.466
	rs9623661	0.197	0.043	3.86E-06	0.050	0.050	0.317
	rs1054402	0.131	0.027	1.13E-06	0.067	0.057	0.243
	rs11627423	0.117	0.025	2.12E-06	-0.066	0.048	0.170
	rs11869294	0.227	0.047	1.50E-06	0.027	0.146	0.856
	rs12121840	0.269	0.057	2.43E-06	0.023	0.099	0.820
ebi-a-GCST010782	rs187166731	0.233	0.051	4.70E-06	0.117	0.306	0.704
	rs2809154	0.179	0.039	3.74E-06	0.028	0.065	0.671
	rs56134659	0.112	0.024	2.44E-06	0.038	0.051	0.453
	rs61335305	0.445	0.091	1.00E-06	-0.037	0.101	0.715
	rs9623661	0.197	0.043	3.86E-06	-0.074	0.075	0.327
	rs1054402	0.131	0.027	1.13E-06	0.023	0.040	0.569
	rs11627423	0.117	0.025	2.12E-06	-0.016	0.038	0.669
	rs11869294	0.227	0.047	1.50E-06	0.115	0.100	0.250
	rs12121840	0.269	0.057	2.43E-06	0.082	0.070	0.243
ebi-a-GCST010783	rs187166731	0.233	0.051	4.70E-06	-0.089	0.220	0.687
	rs2809154	0.179	0.039	3.74E-06	0.058	0.037	0.118
	rs56134659	0.112	0.024	2.44E-06	-0.040	0.038	0.292
	rs61335305	0.445	0.091	1.00E-06	0.114	0.114	0.319
	rs9623661	0.197	0.043	3.86E-06	-0.040	0.065	0.546
	rs1024949	-0.448	0.040	2.95E-29	-0.257	0.221	0.245
	rs113553332	0.685	0.056	4.68E-34	0.198	0.383	0.606
	rs13059920	0.420	0.038	9.33E-29	0.007	0.242	0.975
	rs13084868	0.178	0.039	4.27E-06	-0.062	0.209	0.768
	rs148996090	1.111	0.137	3.98E-16	1.174	0.650	0.071
	rs1607264	-0.146	0.031	2.82E-06	0.203	0.186	0.275
	rs16969649	-0.140	0.031	4.90E-06	-0.233	0.183	0.202
	rs1988743	0.135	0.028	1.51E-06	-0.091	0.178	0.608
	rs2885370	-0.209	0.031	8.91E-12	-0.081	0.185	0.663
	rs3935774	0.741	0.112	3.55E-11	-0.678	0.518	0.191
	rs41268633	-0.251	0.050	6.03E-07	0.542	0.380	0.153
	rs4624606	-0.925	0.025	1.00E-200	-0.144	0.185	0.437
	rs4687154	-0.581	0.040	7.94E-49	0.144	0.262	0.582
ebi-a-GCST010775	rs4815878	0.155	0.030	1.95E-07	0.355	0.168	0.035
	rs5753690	0.149	0.029	2.95E-07	-0.269	0.174	0.123
	rs61492644	-0.124	0.025	7.41E-07	0.002	0.156	0.990
	rs62116908	-0.294	0.064	3.72E-06	0.039	0.422	0.926
	rs62196769	-0.205	0.044	3.09E-06	-0.259	0.219	0.237
	rs6444430	0.154	0.033	2.29E-06	-0.025	0.206	0.905
	rs6444435	-0.171	0.026	5.37E-11	-0.064	0.155	0.679
	rs6461953	-0.146	0.032	4.68E-06	0.339	0.189	0.073
	rs67249092	-0.379	0.035	5.25E-28	0.035	0.227	0.879
	rs6796131	-0.244	0.026	1.51E-20	-0.228	0.165	0.166
	rs6801017	-0.202	0.031	3.80E-11	0.122	0.186	0.512
	rs73754360	0.384	0.080	1.78E-06	0.187	0.495	0.705
	rs75857200	0.199	0.042	2.34E-06	-0.175	0.285	0.538
	rs7623802	-0.129	0.027	1.41E-06	-0.033	0.156	0.834
	rs78129851	0.793	0.117	1.07E-11	-0.842	0.658	0.200
	rs1024949	-0.448	0.040	2.95E-29	-0.025	0.028	0.371
	rs113553332	0.685	0.056	4.68E-34	0.048	0.037	0.199
	rs13059920	0.420	0.038	9.33E-29	-0.010	0.025	0.669

	rs13084868	0.178	0.039	4.27E-06	-0.025	0.022	0.240
	rs148996090	1.111	0.137	3.98E-16	-0.094	0.116	0.417
	rs1607264	-0.146	0.031	2.82E-06	0.005	0.019	0.801
	rs16969649	-0.140	0.031	4.90E-06	0.030	0.019	0.110
	rs1988743	0.135	0.028	1.51E-06	-0.003	0.021	0.892
	rs2885370	-0.209	0.031	8.91E-12	0.013	0.021	0.521
	rs3935774	0.741	0.112	3.55E-11	0.059	0.059	0.320
	rs41268633	-0.251	0.050	6.03E-07	0.027	0.039	0.496
	rs4624606	-0.925	0.025	1.00E-200	0.007	0.021	0.744
	rs4687154	-0.581	0.040	7.94E-49	-0.015	0.026	0.561
ebi-a-GCST010776	rs4815878	0.155	0.030	1.95E-07	-0.008	0.018	0.663
	rs5753690	0.149	0.029	2.95E-07	0.014	0.017	0.408
	rs61492644	-0.124	0.025	7.41E-07	0.005	0.017	0.768
	rs62116908	-0.294	0.064	3.72E-06	-0.026	0.040	0.511
	rs62196769	-0.205	0.044	3.09E-06	-0.044	0.026	0.089
	rs6444430	0.154	0.033	2.29E-06	0.006	0.024	0.792
	rs6444435	-0.171	0.026	5.37E-11	-0.006	0.018	0.720
	rs6461953	-0.146	0.032	4.68E-06	0.006	0.019	0.751
	rs67249092	-0.379	0.035	5.25E-28	-0.002	0.021	0.917
	rs6796131	-0.244	0.026	1.51E-20	-0.019	0.015	0.202
	rs6801017	-0.202	0.031	3.80E-11	-0.016	0.019	0.413
	rs73754360	0.384	0.080	1.78E-06	-0.012	0.043	0.787
	rs75857200	0.199	0.042	2.34E-06	0.044	0.026	0.095
	rs7623802	-0.129	0.027	1.41E-06	-0.018	0.015	0.236
	rs78129851	0.793	0.117	1.07E-11	0.066	0.091	0.469
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	rs1024949	-0.448	0.040	2.95E-29	-0.030	0.031	0.341
	rs113553332	0.685	0.056	4.68E-34	0.046	0.044	0.296
	rs13059920	0.420	0.038	9.33E-29	-0.011	0.025	0.667
	rs13084868	0.178	0.039	4.27E-06	0.008	0.022	0.719
	rs148996090	1.111	0.137	3.98E-16	-0.139	0.144	0.335
	rs1607264	-0.146	0.031	2.82E-06	0.012	0.019	0.506
	rs16969649	-0.140	0.031	4.90E-06	0.044	0.020	0.030
	rs1988743	0.135	0.028	1.51E-06	-0.007	0.024	0.765
	rs2885370	-0.209	0.031	8.91E-12	0.030	0.023	0.184
	rs3935774	0.741	0.112	3.55E-11	0.047	0.043	0.284
	rs41268633	-0.251	0.050	6.03E-07	0.032	0.043	0.457
	rs4624606	-0.925	0.025	1.00E-200	0.008	0.023	0.735
	rs4687154	-0.581	0.040	7.94E-49	-0.020	0.028	0.470
ebi-a-GCST010777	rs4815878	0.155	0.030	1.95E-07	-0.013	0.019	0.492
	rs5753690	0.149	0.029	2.95E-07	0.023	0.018	0.193
	rs61492644	-0.124	0.025	7.41E-07	0.021	0.018	0.247
	rs62116908	-0.294	0.064	3.72E-06	-0.007	0.046	0.884
	rs62196769	-0.205	0.044	3.09E-06	-0.014	0.027	0.608
	rs6444430	0.154	0.033	2.29E-06	0.003	0.025	0.900
	rs6444435	-0.171	0.026	5.37E-11	0.006	0.020	0.758
	rs6461953	-0.146	0.032	4.68E-06	0.026	0.019	0.179
	rs67249092	-0.379	0.035	5.25E-28	-0.003	0.021	0.890
	rs6796131	-0.244	0.026	1.51E-20	-0.005	0.016	0.776
	rs6801017	-0.202	0.031	3.80E-11	-0.016	0.019	0.399
	rs73754360	0.384	0.080	1.78E-06	0.018	0.041	0.659
	rs75857200	0.199	0.042	2.34E-06	0.026	0.028	0.369
	rs7623802	-0.129	0.027	1.41E-06	-0.015	0.017	0.380
	rs78129851	0.793	0.117	1.07E-11	0.160	0.111	0.149
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	rs1024949	-0.448	0.040	2.95E-29	-0.030	0.031	0.341
	rs113553332	0.685	0.056	4.68E-34	0.046	0.044	0.296
	rs13059920	0.420	0.038	9.33E-29	-0.011	0.025	0.667
	rs13084868	0.178	0.039	4.27E-06	0.008	0.022	0.719
	rs148996090	1.111	0.137	3.98E-16	-0.139	0.144	0.335

	rs1607264	-0.146	0.031	2.82E-06	0.012	0.019	0.506
	rs16969649	-0.140	0.031	4.90E-06	0.044	0.020	0.030
	rs1988743	0.135	0.028	1.51E-06	-0.007	0.024	0.765
	rs2885370	-0.209	0.031	8.91E-12	0.030	0.023	0.184
	rs3935774	0.741	0.112	3.55E-11	0.047	0.043	0.284
	rs41268633	-0.251	0.050	6.03E-07	0.032	0.043	0.457
	rs4624606	-0.925	0.025	1.00E-200	0.008	0.023	0.735
	rs4687154	-0.581	0.040	7.94E-49	-0.020	0.028	0.470
ebi-a-GCST010778	rs4815878	0.155	0.030	1.95E-07	-0.013	0.019	0.492
	rs5753690	0.149	0.029	2.95E-07	0.023	0.018	0.193
	rs61492644	-0.124	0.025	7.41E-07	0.021	0.018	0.247
	rs62116908	-0.294	0.064	3.72E-06	-0.007	0.046	0.884
	rs62196769	-0.205	0.044	3.09E-06	-0.014	0.027	0.608
	rs6444430	0.154	0.033	2.29E-06	0.003	0.025	0.900
	rs6444435	-0.171	0.026	5.37E-11	0.006	0.020	0.758
	rs6461953	-0.146	0.032	4.68E-06	0.026	0.019	0.179
	rs67249092	-0.379	0.035	5.25E-28	-0.003	0.021	0.890
	rs6796131	-0.244	0.026	1.51E-20	-0.005	0.016	0.776
	rs6801017	-0.202	0.031	3.80E-11	-0.016	0.019	0.399
	rs73754360	0.384	0.080	1.78E-06	0.018	0.041	0.659
	rs75857200	0.199	0.042	2.34E-06	0.026	0.028	0.369
	rs7623802	-0.129	0.027	1.41E-06	-0.015	0.017	0.380
	rs78129851	0.793	0.117	1.07E-11	0.160	0.111	0.149
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	rs1024949	-0.448	0.040	2.95E-29	0.006	0.044	0.886
	rs113553332	0.685	0.056	4.68E-34	0.001	0.053	0.983
	rs13059920	0.420	0.038	9.33E-29	0.075	0.039	0.055
	rs13084868	0.178	0.039	4.27E-06	-0.037	0.030	0.212
	rs148996090	1.111	0.137	3.98E-16	0.069	0.223	0.758
	rs1607264	-0.146	0.031	2.82E-06	0.039	0.028	0.170
	rs16969649	-0.140	0.031	4.90E-06	-0.005	0.025	0.844
	rs1988743	0.135	0.028	1.51E-06	-0.036	0.031	0.244
	rs2885370	-0.209	0.031	8.91E-12	0.022	0.034	0.513
	rs3935774	0.741	0.112	3.55E-11	0.031	0.060	0.608
	rs41268633	-0.251	0.050	6.03E-07	0.088	0.079	0.263
	rs4624606	-0.925	0.025	1.00E-200	0.012	0.031	0.702
	rs4687154	-0.581	0.040	7.94E-49	-0.031	0.044	0.485
IL-1Racp ebi-a-GCST010779	rs4815878	0.155	0.030	1.95E-07	0.052	0.029	0.069
	rs5753690	0.149	0.029	2.95E-07	0.027	0.024	0.272
	rs61492644	-0.124	0.025	7.41E-07	0.017	0.028	0.536
	rs62116908	-0.294	0.064	3.72E-06	-0.025	0.062	0.691
	rs62196769	-0.205	0.044	3.09E-06	-0.048	0.037	0.190
	rs6444430	0.154	0.033	2.29E-06	0.043	0.046	0.346
	rs6444435	-0.171	0.026	5.37E-11	-0.048	0.029	0.100
	rs6461953	-0.146	0.032	4.68E-06	0.030	0.026	0.253
	rs67249092	-0.379	0.035	5.25E-28	0.024	0.033	0.461
	rs6796131	-0.244	0.026	1.51E-20	-0.031	0.022	0.153
	rs6801017	-0.202	0.031	3.80E-11	-0.018	0.029	0.537
	rs73754360	0.384	0.080	1.78E-06	-0.009	0.055	0.870
	rs75857200	0.199	0.042	2.34E-06	-0.030	0.043	0.477
	rs7623802	-0.129	0.027	1.41E-06	-0.006	0.022	0.785
	rs78129851	0.793	0.117	1.07E-11	0.029	0.127	0.820
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	rs1024949	-0.448	0.040	2.95E-29	-0.007	0.025	0.774
	rs113553332	0.685	0.056	4.68E-34	0.025	0.035	0.483
	rs13059920	0.420	0.038	9.33E-29	-0.008	0.022	0.727
	rs13084868	0.178	0.039	4.27E-06	-0.004	0.018	0.821
	rs148996090	1.111	0.137	3.98E-16	-0.133	0.111	0.229
	rs1607264	-0.146	0.031	2.82E-06	0.019	0.016	0.233
	rs16969649	-0.140	0.031	4.90E-06	0.028	0.017	0.097

	rs1988743	0.135	0.028	1.51E-06	0.001	0.018	0.946
	rs2885370	-0.209	0.031	8.91E-12	0.002	0.019	0.897
	rs3935774	0.741	0.112	3.55E-11	0.041	0.035	0.245
	rs41268633	-0.251	0.050	6.03E-07	0.036	0.037	0.325
	rs4624606	-0.925	0.025	1.00E-200	0.009	0.018	0.610
	rs4687154	-0.581	0.040	7.94E-49	-0.018	0.024	0.471
ebi-a-GCST010780	rs4815878	0.155	0.030	1.95E-07	-0.005	0.016	0.765
	rs5753690	0.149	0.029	2.95E-07	0.022	0.015	0.134
	rs61492644	-0.124	0.025	7.41E-07	0.008	0.015	0.586
	rs62116908	-0.294	0.064	3.72E-06	-0.026	0.038	0.496
	rs62196769	-0.205	0.044	3.09E-06	-0.038	0.022	0.087
	rs6444430	0.154	0.033	2.29E-06	0.016	0.023	0.480
	rs6444435	-0.171	0.026	5.37E-11	-0.015	0.016	0.350
	rs6461953	-0.146	0.032	4.68E-06	0.017	0.016	0.296
	rs67249092	-0.379	0.035	5.25E-28	0.003	0.018	0.873
	rs6796131	-0.244	0.026	1.51E-20	-0.007	0.013	0.627
	rs6801017	-0.202	0.031	3.80E-11	-0.020	0.016	0.206
	rs73754360	0.384	0.080	1.78E-06	-0.009	0.035	0.806
	rs75857200	0.199	0.042	2.34E-06	0.037	0.024	0.133
	rs7623802	-0.129	0.027	1.41E-06	-0.012	0.014	0.396
	rs78129851	0.793	0.117	1.07E-11	0.049	0.083	0.556
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	rs1024949	-0.448	0.040	2.95E-29	-0.002	0.048	0.967
	rs113553332	0.685	0.056	4.68E-34	-0.019	0.065	0.773
	rs13059920	0.420	0.038	9.33E-29	0.009	0.043	0.835
	rs13084868	0.178	0.039	4.27E-06	-0.041	0.039	0.287
	rs148996090	1.111	0.137	3.98E-16	-0.135	0.205	0.512
	rs1607264	-0.146	0.031	2.82E-06	-0.033	0.035	0.345
	rs16969649	-0.140	0.031	4.90E-06	-0.004	0.032	0.898
	rs1988743	0.135	0.028	1.51E-06	0.077	0.033	0.018
	rs2885370	-0.209	0.031	8.91E-12	-0.096	0.036	0.009
	rs3935774	0.741	0.112	3.55E-11	0.246	0.137	0.073
	rs41268633	-0.251	0.050	6.03E-07	-0.007	0.065	0.914
	rs4624606	-0.925	0.025	1.00E-200	-0.043	0.035	0.214
	rs4687154	-0.581	0.040	7.94E-49	-0.024	0.054	0.657
ebi-a-GCST010781	rs4815878	0.155	0.030	1.95E-07	-0.015	0.038	0.696
	rs5753690	0.149	0.029	2.95E-07	0.075	0.031	0.016
	rs61492644	-0.124	0.025	7.41E-07	-0.006	0.027	0.822
	rs62116908	-0.294	0.064	3.72E-06	-0.022	0.076	0.775
	rs62196769	-0.205	0.044	3.09E-06	-0.026	0.046	0.569
	rs6444430	0.154	0.033	2.29E-06	0.066	0.044	0.134
	rs6444435	-0.171	0.026	5.37E-11	-0.008	0.031	0.804
	rs6461953	-0.146	0.032	4.68E-06	-0.061	0.037	0.096
	rs67249092	-0.379	0.035	5.25E-28	0.051	0.042	0.218
	rs6796131	-0.244	0.026	1.51E-20	-0.058	0.029	0.042
	rs6801017	-0.202	0.031	3.80E-11	-0.055	0.033	0.095
	rs73754360	0.384	0.080	1.78E-06	-0.079	0.071	0.267
	rs75857200	0.199	0.042	2.34E-06	-0.002	0.047	0.964
	rs7623802	-0.129	0.027	1.41E-06	0.001	0.028	0.977
	rs78129851	0.793	0.117	1.07E-11	0.304	0.157	0.053
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	rs1024949	-0.448	0.040	2.95E-29	-0.013	0.082	0.872
	rs113553332	0.685	0.056	4.68E-34	-0.055	0.110	0.619
	rs13059920	0.420	0.038	9.33E-29	0.087	0.064	0.177
	rs13084868	0.178	0.039	4.27E-06	-0.018	0.055	0.750
	rs148996090	1.111	0.137	3.98E-16	0.378	0.363	0.297
	rs1607264	-0.146	0.031	2.82E-06	0.103	0.046	0.024
	rs16969649	-0.140	0.031	4.90E-06	-0.004	0.049	0.934
	rs1988743	0.135	0.028	1.51E-06	-0.055	0.063	0.380
	rs2885370	-0.209	0.031	8.91E-12	0.056	0.061	0.360

	rs3935774	0.741	0.112	3.55E-11	0.040	0.102	0.691
	rs41268633	-0.251	0.050	6.03E-07	-0.011	0.129	0.933
	rs4624606	-0.925	0.025	1.00E-200	0.009	0.062	0.883
	rs4687154	-0.581	0.040	7.94E-49	-0.008	0.073	0.913
ebi-a-GCST010782	rs4815878	0.155	0.030	1.95E-07	0.083	0.047	0.077
	rs5753690	0.149	0.029	2.95E-07	0.012	0.045	0.793
	rs61492644	-0.124	0.025	7.41E-07	-0.025	0.049	0.613
	rs62116908	-0.294	0.064	3.72E-06	0.002	0.121	0.989
	rs62196769	-0.205	0.044	3.09E-06	-0.083	0.069	0.231
	rs6444430	0.154	0.033	2.29E-06	0.113	0.073	0.120
	rs6444435	-0.171	0.026	5.37E-11	-0.037	0.056	0.508
	rs6461953	-0.146	0.032	4.68E-06	-0.028	0.049	0.565
	rs67249092	-0.379	0.035	5.25E-28	0.055	0.054	0.312
	rs6796131	-0.244	0.026	1.51E-20	-0.041	0.042	0.332
	rs6801017	-0.202	0.031	3.80E-11	-0.040	0.049	0.412
	rs73754360	0.384	0.080	1.78E-06	-0.026	0.104	0.800
	rs75857200	0.199	0.042	2.34E-06	-0.003	0.075	0.973
	rs7623802	-0.129	0.027	1.41E-06	0.080	0.045	0.074
	rs78129851	0.793	0.117	1.07E-11	-0.219	0.267	0.411
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	rs1024949	-0.448	0.040	2.95E-29	-0.010	0.063	0.872
	rs113553332	0.685	0.056	4.68E-34	0.024	0.067	0.718
	rs13059920	0.420	0.038	9.33E-29	0.056	0.061	0.355
	rs13084868	0.178	0.039	4.27E-06	-0.031	0.041	0.453
	rs148996090	1.111	0.137	3.98E-16	0.240	0.262	0.360
	rs1607264	-0.146	0.031	2.82E-06	0.072	0.044	0.101
	rs16969649	-0.140	0.031	4.90E-06	-0.015	0.034	0.666
	rs1988743	0.135	0.028	1.51E-06	-0.026	0.041	0.529
	rs2885370	-0.209	0.031	8.91E-12	0.037	0.048	0.442
	rs3935774	0.741	0.112	3.55E-11	-0.035	0.094	0.711
	rs41268633	-0.251	0.050	6.03E-07	0.224	0.113	0.047
	rs4624606	-0.925	0.025	1.00E-200	0.001	0.043	0.975
	rs4687154	-0.581	0.040	7.94E-49	0.019	0.066	0.776
ebi-a-GCST010783	rs4815878	0.155	0.030	1.95E-07	0.107	0.045	0.017
	rs5753690	0.149	0.029	2.95E-07	0.047	0.033	0.150
	rs61492644	-0.124	0.025	7.41E-07	0.011	0.042	0.784
	rs62116908	-0.294	0.064	3.72E-06	0.018	0.080	0.818
	rs62196769	-0.205	0.044	3.09E-06	-0.021	0.049	0.668
	rs6444430	0.154	0.033	2.29E-06	0.022	0.070	0.752
	rs6444435	-0.171	0.026	5.37E-11	-0.049	0.041	0.227
	rs6461953	-0.146	0.032	4.68E-06	0.043	0.035	0.215
	rs67249092	-0.379	0.035	5.25E-28	0.000	0.051	0.995
	rs6796131	-0.244	0.026	1.51E-20	-0.040	0.029	0.175
	rs6801017	-0.202	0.031	3.80E-11	0.037	0.044	0.398
	rs73754360	0.384	0.080	1.78E-06	0.074	0.075	0.324
	rs75857200	0.199	0.042	2.34E-06	0.065	0.062	0.295
	rs7623802	-0.129	0.027	1.41E-06	-0.042	0.029	0.148
	rs78129851	0.793	0.117	1.07E-11	-0.025	0.143	0.860

COVID-19: Corona Virus Disease 2019; GWAS ID: Genome wide association study identity; SNP: single-nucleotide polymorphism; Beta is the regression coefficient based on the IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp raising allele (effect allele); SE: standard error.

Supplementary table 3. Pleiotropy and heterogeneity test of IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp genetic IVs in COVID-19 GWAS dataset.

Exposure	GWAS ID	Pleiotropy test				Heterogeneity test					
		MR_Egger			PRESSO	MR Egger			IVW		
		Intercept	SE	<i>p</i> val	<i>p</i> val	Q	Q_df	<i>p</i> val	Q	Q_df	<i>p</i> val
IL-1A	ebi-a-GCST010775	-0.08	0.18	0.67	0.16	23.63	16	0.10	23.90	17	0.12
	ebi-a-GCST010776	0.02	0.02	0.16	0.04	26.34	17	0.07	29.76	18	0.04
	ebi-a-GCST010777	0.04	0.02	0.02	0.16	16.76	17	0.47	23.79	18	0.16
	ebi-a-GCST010778	0.04	0.02	0.02	0.15	16.76	17	0.47	23.79	18	0.16
	ebi-a-GCST010779	0.03	0.02	0.26	0.17	22.26	17	0.18	24.00	18	0.16
	ebi-a-GCST010780	0.02	0.02	0.27	0.03	28.90	17	0.04	31.06	18	0.03
	ebi-a-GCST010781	0.01	0.02	0.56	0.65	15.01	17	0.59	15.36	18	0.64
	ebi-a-GCST010782	0.09	0.04	0.04	0.46	13.43	17	0.71	18.28	18	0.44
	ebi-a-GCST010783	0.02	0.03	0.41	0.23	21.65	17	0.20	22.55	18	0.21
IL-1B	ebi-a-GCST010775	0.02	0.21	0.93	0.35	5.32	4	0.26	5.33	5	0.38
	ebi-a-GCST010776	-0.01	0.02	0.53	0.80	1.90	4	0.75	2.37	5	0.80
	ebi-a-GCST010777	0.00	0.02	0.80	0.83	2.46	4	0.65	2.53	5	0.77
	ebi-a-GCST010778	0.00	0.02	0.80	0.83	2.46	4	0.65	2.53	5	0.77
	ebi-a-GCST010779	-0.06	0.03	0.09	0.20	3.33	4	0.50	8.32	5	0.14
	ebi-a-GCST010780	-0.02	0.02	0.25	0.67	1.81	4	0.77	3.62	5	0.61
	ebi-a-GCST010781	-0.02	0.03	0.66	0.69	2.49	4	0.65	2.71	5	0.74
	ebi-a-GCST010782	-0.05	0.05	0.37	0.33	4.85	4	0.30	6.11	5	0.30
	ebi-a-GCST010783	-0.03	0.04	0.54	0.48	4.49	4	0.34	4.99	5	0.42
IL-1R1	ebi-a-GCST010775	0.05	0.12	0.68	0.94	8.70	16	0.93	8.88	17	0.94
	ebi-a-GCST010776	0.02	0.01	0.10	0.63	13.01	18	0.79	16.02	19	0.66
	ebi-a-GCST010777	0.01	0.01	0.29	0.69	14.02	18	0.73	15.24	19	0.71
	ebi-a-GCST010778	0.01	0.01	0.29	0.70	14.02	18	0.73	15.24	19	0.71
	ebi-a-GCST010779	0.00	0.02	0.88	0.70	15.19	18	0.65	15.22	19	0.71
	ebi-a-GCST010780	0.02	0.01	0.10	0.54	14.70	18	0.68	17.71	19	0.54
	ebi-a-GCST010781	-0.01	0.02	0.71	0.37	20.16	18	0.32	20.31	19	0.38
	ebi-a-GCST010782	-0.02	0.03	0.62	0.95	9.77	18	0.94	10.03	19	0.95
	ebi-a-GCST010783	0.01	0.02	0.65	0.92	10.96	18	0.90	11.18	19	0.92
IL-1Ra	ebi-a-GCST010775	-0.05	0.19	0.80	0.62	6.09	7	0.53	6.16	8	0.63
	ebi-a-GCST010776	-0.04	0.02	0.14	0.44	5.19	7	0.64	8.01	8	0.43
	ebi-a-GCST010777	-0.03	0.03	0.36	0.05	14.62	7	0.04	16.65	8	0.03
	ebi-a-GCST010778	-0.03	0.03	0.36	0.05	14.62	7	0.04	16.65	8	0.03
	ebi-a-GCST010779	-0.01	0.03	0.69	0.47	7.11	7	0.42	7.29	8	0.51
	ebi-a-GCST010780	-0.02	0.02	0.27	0.49	5.89	7	0.55	7.34	8	0.50
	ebi-a-GCST010781	0.03	0.05	0.57	0.28	11.47	8	0.18	11.97	9	0.22
	ebi-a-GCST010782	0.02	0.05	0.78	0.74	5.22	7	0.63	5.31	8	0.72
	ebi-a-GCST010783	-0.07	0.05	0.18	0.61	4.10	7	0.77	6.33	8	0.61
IL-1Racp	ebi-a-GCST010775	-0.03	0.07	0.65	0.46	28.24	26	0.35	28.47	27	0.39
	ebi-a-GCST010776	0.00	0.01	0.81	0.83	20.10	26	0.79	20.16	27	0.82
	ebi-a-GCST010777	-0.01	0.01	0.29	0.73	21.59	26	0.71	22.73	27	0.70
	ebi-a-GCST010778	-0.01	0.01	0.29	0.74	21.59	26	0.71	22.73	27	0.70
	ebi-a-GCST010779	0.00	0.01	0.77	0.55	25.80	26	0.47	25.88	27	0.53
	ebi-a-GCST010780	0.00	0.01	0.70	0.70	22.58	26	0.66	22.73	27	0.70
	ebi-a-GCST010781	0.03	0.01	0.06	0.11	32.35	26	0.18	37.24	27	0.09
	ebi-a-GCST010782	0.00	0.02	0.89	0.63	24.64	26	0.54	24.66	27	0.59
	ebi-a-GCST010783	0.01	0.01	0.45	0.47	27.13	26	0.40	27.74	27	0.42

COVID-19: Corona Virus Disease 2019; GWAS ID: Genome wide association study identity. IVW: Inverse variance weighted. SE: standard error. *p* val > 0.05 represent no significant pleiotropy and heterogeneity.

Supplementary table 4. The causal association of IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp with COVID-19.

Exposure	GWAS ID	Method	n SNP	Beta	SE	p val	OR	OR_lci95	OR_uci95
IL-1A	ebi-a-GCST010775	MR Egger	18	0.086	1.147	0.941	1.090	0.115	10.317
		Weighted median	18	-0.022	0.447	0.961	0.979	0.408	2.349
		IVW	18	-0.381	0.372	0.306	0.683	0.330	1.417
		Simple mode	18	0.208	0.702	0.771	1.231	0.311	4.877
		Weighted mode	18	0.208	0.637	0.748	1.231	0.353	4.290
	ebi-a-GCST010776	MR Egger	19	-0.139	0.103	0.194	0.870	0.711	1.064
		Weighted median	19	0.080	0.045	0.074	1.083	0.992	1.182
		IVW	19	0.002	0.041	0.961	1.002	0.926	1.085
		Simple mode	19	0.078	0.082	0.356	1.081	0.920	1.270
		Weighted mode	19	0.076	0.075	0.325	1.079	0.931	1.249
	ebi-a-GCST010777	MR Egger	19	-0.247	0.096	0.020	0.781	0.647	0.944
		Weighted median	19	0.044	0.050	0.384	1.045	0.947	1.153
		IVW	19	-0.007	0.039	0.849	0.993	0.919	1.072
		Simple mode	19	0.113	0.101	0.278	1.120	0.918	1.366
		Weighted mode	19	0.121	0.091	0.202	1.128	0.944	1.349
	ebi-a-GCST010778	MR Egger	19	-0.247	0.096	0.020	0.781	0.647	0.944
		Weighted median	19	0.044	0.052	0.400	1.045	0.944	1.157
		IVW	19	-0.007	0.039	0.849	0.993	0.919	1.072
		Simple mode	19	0.113	0.106	0.298	1.120	0.910	1.378
		Weighted mode	19	0.121	0.085	0.171	1.128	0.956	1.332
	ebi-a-GCST010779	MR Egger	19	-0.155	0.139	0.279	0.856	0.653	1.124
		Weighted median	19	0.029	0.067	0.666	1.029	0.902	1.174
		IVW	19	-0.008	0.055	0.887	0.992	0.891	1.105
		Simple mode	19	0.059	0.124	0.642	1.060	0.832	1.352
		Weighted mode	19	0.091	0.117	0.444	1.096	0.871	1.378
	ebi-a-GCST010780	MR Egger	19	-0.090	0.099	0.374	0.914	0.752	1.109
		Weighted median	19	0.059	0.041	0.150	1.061	0.979	1.150
		IVW	19	0.013	0.037	0.717	1.013	0.943	1.089
Simple mode		19	0.065	0.069	0.358	1.067	0.933	1.220	
Weighted mode		19	0.078	0.071	0.285	1.082	0.941	1.243	
ebi-a-GCST010781	MR Egger	19	-0.104	0.153	0.506	0.902	0.668	1.216	
	Weighted median	19	0.000	0.077	0.997	1.000	0.860	1.163	
	IVW	19	-0.019	0.056	0.728	0.981	0.880	1.094	
	Simple mode	19	0.046	0.150	0.765	1.047	0.780	1.405	
	Weighted mode	19	0.094	0.142	0.516	1.099	0.832	1.452	
ebi-a-GCST010782	MR Egger	19	-0.716	0.264	0.015	0.489	0.292	0.819	
	Weighted median	19	-0.189	0.129	0.144	0.828	0.643	1.066	
	IVW	19	-0.171	0.092	0.063	0.843	0.704	1.009	
	Simple mode	19	-0.372	0.255	0.161	0.689	0.418	1.136	
	Weighted mode	19	-0.275	0.234	0.254	0.759	0.480	1.200	
ebi-a-GCST010783	MR Egger	19	-0.169	0.176	0.350	0.844	0.598	1.192	
	Weighted median	19	-0.004	0.092	0.962	0.996	0.831	1.193	
	IVW	19	-0.035	0.073	0.630	0.965	0.837	1.114	
	Simple mode	19	0.095	0.166	0.572	1.100	0.795	1.523	
	Weighted mode	19	0.066	0.142	0.646	1.069	0.809	1.412	
ebi-a-GCST010775	MR Egger	6	-1.451	1.548	0.402	0.234	0.011	4.873	
	Weighted median	6	-0.615	0.886	0.488	0.541	0.095	3.069	
	IVW	6	-1.332	0.684	0.051	0.264	0.069	1.009	
	Simple mode	6	-2.398	1.464	0.162	0.091	0.005	1.602	
	Weighted mode	6	-0.192	1.125	0.871	0.825	0.091	7.487	
ebi-a-GCST010776	MR Egger	6	0.030	0.139	0.842	1.030	0.784	1.353	
	Weighted median	6	-0.072	0.089	0.421	0.931	0.781	1.109	
	IVW	6	-0.054	0.066	0.412	0.947	0.833	1.078	
	Simple mode	6	-0.051	0.116	0.677	0.950	0.758	1.192	

IL-1B	ebi-a-GCST010777	Weighted mode	6	-0.093	0.096	0.376	0.911	0.755	1.099
		MR Egger	6	0.105	0.132	0.472	1.110	0.858	1.437
		Weighted median	6	0.087	0.082	0.285	1.091	0.930	1.280
		IVW	6	0.074	0.065	0.255	1.077	0.948	1.223
		Simple mode	6	0.112	0.112	0.362	1.118	0.899	1.391
	ebi-a-GCST010778	Weighted mode	6	0.078	0.095	0.449	1.081	0.898	1.301
		MR Egger	6	0.105	0.132	0.472	1.110	0.858	1.437
		Weighted median	6	0.087	0.084	0.303	1.091	0.924	1.287
		IVW	6	0.074	0.065	0.255	1.077	0.948	1.223
		Simple mode	6	0.112	0.106	0.339	1.118	0.909	1.376
	ebi-a-GCST010779	Weighted mode	6	0.078	0.093	0.441	1.081	0.901	1.296
		MR Egger	6	0.487	0.199	0.070	1.628	1.102	2.403
		Weighted median	6	0.188	0.130	0.148	1.207	0.935	1.559
		IVW	6	0.102	0.127	0.425	1.107	0.862	1.421
		Simple mode	6	0.274	0.183	0.194	1.315	0.919	1.881
	ebi-a-GCST010780	Weighted mode	6	0.228	0.154	0.199	1.256	0.928	1.700
		MR Egger	6	0.155	0.113	0.242	1.168	0.936	1.457
		Weighted median	6	0.017	0.071	0.817	1.017	0.884	1.169
		IVW	6	0.023	0.056	0.682	1.023	0.917	1.142
		Simple mode	6	0.036	0.091	0.708	1.037	0.867	1.240
ebi-a-GCST010781	Weighted mode	6	0.026	0.082	0.760	1.027	0.874	1.206	
	MR Egger	6	-0.058	0.241	0.822	0.944	0.589	1.512	
	Weighted median	6	-0.215	0.141	0.126	0.806	0.612	1.063	
	IVW	6	-0.159	0.109	0.146	0.853	0.689	1.057	
	Simple mode	6	-0.225	0.207	0.328	0.799	0.532	1.199	
ebi-a-GCST010782	Weighted mode	6	-0.235	0.140	0.154	0.790	0.601	1.040	
	MR Egger	6	0.218	0.350	0.568	1.243	0.626	2.468	
	Weighted median	6	0.000	0.203	0.998	1.000	0.672	1.489	
	IVW	6	-0.093	0.172	0.589	0.911	0.650	1.277	
	Simple mode	6	-0.124	0.311	0.706	0.883	0.480	1.624	
ebi-a-GCST010783	Weighted mode	6	0.111	0.236	0.658	1.117	0.704	1.774	
	MR Egger	6	0.134	0.287	0.665	1.143	0.651	2.006	
	Weighted median	6	-0.159	0.169	0.345	0.853	0.612	1.187	
	IVW	6	-0.035	0.128	0.786	0.966	0.751	1.242	
	Simple mode	6	-0.122	0.267	0.667	0.885	0.524	1.494	
ebi-a-GCST010775	Weighted mode	6	-0.158	0.177	0.412	0.854	0.604	1.207	
	MR Egger	18	-0.194	0.674	0.777	0.823	0.220	3.086	
	Weighted median	18	0.112	0.416	0.787	1.119	0.495	2.529	
	IVW	18	0.061	0.302	0.841	1.062	0.588	1.919	
	Simple mode	18	0.256	0.767	0.743	1.292	0.287	5.806	
ebi-a-GCST010776	Weighted mode	18	0.221	0.728	0.765	1.247	0.299	5.196	
	MR Egger	20	-0.134	0.061	0.041	0.875	0.777	0.985	
	Weighted median	20	-0.027	0.040	0.501	0.974	0.900	1.053	
	IVW	20	-0.041	0.028	0.148	0.960	0.908	1.015	
	Simple mode	20	-0.015	0.079	0.854	0.985	0.843	1.151	
ebi-a-GCST010777	Weighted mode	20	-0.016	0.076	0.832	0.984	0.848	1.141	
	MR Egger	20	-0.112	0.068	0.116	0.894	0.783	1.021	
	Weighted median	20	-0.039	0.045	0.390	0.962	0.881	1.051	
	IVW	20	-0.045	0.031	0.143	0.956	0.899	1.015	
	Simple mode	20	-0.135	0.082	0.114	0.873	0.744	1.025	
ebi-a-GCST010778	Weighted mode	20	-0.024	0.081	0.770	0.976	0.834	1.143	
	MR Egger	20	-0.112	0.068	0.116	0.894	0.783	1.021	
	Weighted median	20	-0.039	0.041	0.347	0.962	0.888	1.043	
	IVW	20	-0.045	0.031	0.143	0.956	0.899	1.015	
	Simple mode	20	-0.135	0.085	0.128	0.873	0.739	1.032	
ebi-a-	Weighted mode	20	-0.024	0.081	0.771	0.976	0.834	1.144	
	MR Egger	20	0.041	0.097	0.678	1.042	0.862	1.260	
	Weighted median	20	0.028	0.059	0.641	1.028	0.915	1.155	

IL-1R1	ebi-a-GCST010779	IVW	20	0.027	0.044	0.531	1.028	0.944	1.119
		Simple mode	20	0.029	0.114	0.801	1.030	0.823	1.288
		Weighted mode	20	0.029	0.105	0.785	1.030	0.838	1.266
	ebi-a-GCST010780	MR Egger	20	-0.112	0.057	0.065	0.894	0.800	1.000
		Weighted median	20	0.001	0.037	0.980	1.001	0.931	1.076
		IVW	20	-0.024	0.026	0.352	0.976	0.928	1.027
		Simple mode	20	-0.004	0.071	0.955	0.996	0.867	1.144
	ebi-a-GCST010781	Weighted mode	20	-0.002	0.060	0.968	0.998	0.887	1.122
		MR Egger	20	0.039	0.113	0.735	1.040	0.833	1.297
		Weighted median	20	0.026	0.075	0.726	1.026	0.887	1.188
		IVW	20	0.002	0.053	0.968	1.002	0.903	1.112
	ebi-a-GCST010782	Simple mode	20	-0.006	0.146	0.970	0.995	0.747	1.325
		Weighted mode	20	0.028	0.127	0.826	1.029	0.801	1.321
		MR Egger	20	0.096	0.188	0.615	1.101	0.761	1.593
		Weighted median	20	0.080	0.107	0.456	1.083	0.878	1.336
	ebi-a-GCST010783	IVW	20	0.010	0.083	0.906	1.010	0.858	1.189
		Simple mode	20	0.108	0.183	0.563	1.114	0.778	1.595
		Weighted mode	20	0.136	0.187	0.477	1.145	0.794	1.653
MR Egger		20	0.092	0.125	0.469	1.097	0.859	1.401	
ebi-a-GCST010783	Weighted median	20	0.173	0.085	0.040	1.189	1.008	1.404	
	IVW	20	0.143	0.058	0.014	1.154	1.030	1.293	
	Simple mode	20	0.241	0.143	0.109	1.272	0.961	1.683	
	Weighted mode	20	0.235	0.131	0.089	1.265	0.978	1.635	
IL-1Ra	ebi-a-GCST010775	MR Egger	9	0.481	1.116	0.679	1.618	0.182	14.416
		Weighted median	9	-0.296	0.590	0.616	0.744	0.239	2.316
		IVW	9	0.214	0.433	0.621	1.238	0.530	2.891
		Simple mode	9	-0.363	0.906	0.699	0.695	0.103	4.692
	ebi-a-GCST010776	Weighted mode	9	-0.555	0.897	0.553	0.574	0.100	3.302
		MR Egger	9	0.184	0.136	0.220	1.202	0.920	1.570
		Weighted median	9	-0.073	0.067	0.282	0.930	0.819	1.057
		IVW	9	-0.030	0.050	0.546	0.971	0.881	1.069
	ebi-a-GCST010777	Simple mode	9	0.099	0.114	0.412	1.104	0.897	1.359
		Weighted mode	9	-0.075	0.101	0.479	0.928	0.755	1.140
		MR Egger	9	0.122	0.170	0.495	1.130	0.810	1.576
		Weighted median	9	0.037	0.073	0.611	1.038	0.908	1.185
	ebi-a-GCST010778	IVW	9	-0.029	0.072	0.685	0.971	0.843	1.118
		Simple mode	9	-0.005	0.110	0.961	0.995	0.806	1.227
		Weighted mode	9	0.031	0.073	0.686	1.031	0.880	1.208
		MR Egger	9	0.122	0.170	0.495	1.130	0.810	1.576
	ebi-a-GCST010779	Weighted median	9	0.037	0.073	0.611	1.038	0.903	1.192
		IVW	9	-0.029	0.072	0.685	0.971	0.843	1.118
Simple mode		9	-0.005	0.104	0.959	0.995	0.806	1.227	
Weighted mode		9	0.031	0.076	0.695	1.031	0.887	1.199	
ebi-a-GCST010780	MR Egger	9	0.120	0.184	0.535	1.127	0.787	1.616	
	Weighted median	9	0.160	0.099	0.106	1.173	0.967	1.423	
	IVW	9	0.050	0.073	0.492	1.051	0.912	1.212	
	Simple mode	9	0.177	0.134	0.223	1.193	0.915	1.557	
ebi-a-GCST010781	Weighted mode	9	0.175	0.108	0.143	1.191	0.950	1.495	
	MR Egger	9	0.097	0.105	0.387	1.101	0.897	1.352	
	Weighted median	9	0.022	0.059	0.706	1.023	0.914	1.144	
	IVW	9	-0.019	0.042	0.657	0.981	0.903	1.067	
ebi-a-GCST010781	Simple mode	9	0.082	0.100	0.435	1.086	0.908	1.298	
	Weighted mode	9	0.049	0.078	0.546	1.051	0.906	1.218	
	MR Egger	10	-0.113	0.312	0.726	0.893	0.485	1.645	
	Weighted median	10	-0.035	0.120	0.768	0.965	0.767	1.215	
ebi-a-GCST010781	IVW	10	0.060	0.101	0.554	1.062	0.870	1.296	
	Simple mode	10	0.031	0.174	0.864	1.031	0.744	1.429	
	Weighted mode	10	-0.023	0.126	0.858	0.977	0.758	1.260	

ebi-a-GCST010782	MR Egger	9	-0.090	0.292	0.767	0.914	0.516	1.619
	Weighted median	9	0.013	0.173	0.938	1.014	0.723	1.422
	IVW	9	-0.014	0.129	0.914	0.986	0.766	1.269
	Simple mode	9	0.171	0.265	0.537	1.187	0.702	2.005
	Weighted mode	9	0.010	0.196	0.962	1.010	0.690	1.477
ebi-a-GCST010783	MR Egger	9	0.523	0.275	0.099	1.687	0.984	2.892
	Weighted median	9	0.258	0.138	0.061	1.294	1.003	1.670
	IVW	9	0.141	0.101	0.163	1.152	0.944	1.404
	Simple mode	9	0.280	0.202	0.202	1.324	0.879	1.993
	Weighted mode	9	0.280	0.163	0.123	1.324	0.953	1.838
ebi-a-GCST010775	MR Egger	28	0.174	0.207	0.409	1.190	0.793	1.785
	Weighted median	28	0.141	0.180	0.432	1.152	0.810	1.637
	IVW	28	0.100	0.128	0.435	1.105	0.860	1.418
	Simple mode	28	0.061	0.332	0.855	1.063	0.554	2.039
	Weighted mode	28	0.143	0.171	0.410	1.154	0.825	1.615
ebi-a-GCST010776	MR Egger	28	0.010	0.023	0.647	1.011	0.967	1.056
	Weighted median	28	0.001	0.020	0.947	1.001	0.963	1.041
	IVW	28	0.015	0.014	0.280	1.015	0.988	1.043
	Simple mode	28	-0.014	0.034	0.678	0.986	0.923	1.053
	Weighted mode	28	0.004	0.020	0.864	1.004	0.964	1.044
ebi-a-GCST010777	MR Egger	28	0.028	0.024	0.263	1.028	0.981	1.078
	Weighted median	28	0.004	0.021	0.833	1.004	0.964	1.046
	IVW	28	0.007	0.014	0.633	1.007	0.979	1.036
	Simple mode	28	0.040	0.034	0.247	1.041	0.974	1.112
	Weighted mode	28	0.009	0.021	0.674	1.009	0.968	1.052
ebi-a-GCST010778	MR Egger	28	0.028	0.024	0.263	1.028	0.981	1.078
	Weighted median	28	0.004	0.021	0.837	1.004	0.963	1.048
	IVW	28	0.007	0.014	0.633	1.007	0.979	1.036
	Simple mode	28	0.040	0.035	0.258	1.041	0.972	1.114
	Weighted mode	28	0.009	0.019	0.634	1.009	0.972	1.047
IL-1Racp ebi-a-GCST010779	MR Egger	28	0.008	0.033	0.802	1.008	0.945	1.076
	Weighted median	28	-0.005	0.029	0.872	0.995	0.940	1.054
	IVW	28	0.016	0.020	0.430	1.016	0.977	1.057
	Simple mode	28	0.035	0.052	0.499	1.036	0.936	1.146
	Weighted mode	28	0.005	0.027	0.840	1.006	0.954	1.060
ebi-a-GCST010780	MR Egger	28	0.002	0.019	0.909	1.002	0.965	1.041
	Weighted median	28	-0.009	0.018	0.628	0.992	0.958	1.026
	IVW	28	0.008	0.012	0.482	1.008	0.985	1.032
	Simple mode	28	0.011	0.027	0.689	1.011	0.959	1.065
	Weighted mode	28	0.002	0.016	0.898	1.002	0.971	1.034
ebi-a-GCST010781	MR Egger	28	-0.001	0.043	0.986	0.999	0.918	1.088
	Weighted median	28	0.046	0.035	0.184	1.047	0.978	1.121
	IVW	28	0.067	0.028	0.018	1.069	1.011	1.130
	Simple mode	28	0.019	0.064	0.765	1.020	0.899	1.156
	Weighted mode	28	0.033	0.033	0.317	1.034	0.970	1.102
ebi-a-GCST010782	MR Egger	28	0.008	0.062	0.900	1.008	0.892	1.139
	Weighted median	28	-0.007	0.055	0.901	0.993	0.893	1.105
	IVW	28	0.015	0.038	0.695	1.015	0.943	1.092
	Simple mode	28	0.036	0.085	0.677	1.037	0.877	1.225
	Weighted mode	28	0.003	0.055	0.964	1.003	0.901	1.116
ebi-a-GCST010783	MR Egger	28	-0.005	0.047	0.918	0.995	0.908	1.090
	Weighted median	28	-0.001	0.041	0.988	0.999	0.922	1.083
	IVW	28	0.023	0.029	0.419	1.023	0.968	1.082
	Simple mode	28	0.018	0.064	0.782	1.018	0.898	1.155
	Weighted mode	28	0.009	0.039	0.818	1.009	0.935	1.089

COVID-19: Corona Virus Disease 2019; GWAS ID: Genome wide association study identity. IVW: Inverse

variance weighted. Beta: the regression coefficient based on the vitamin C raising effect allele. nsnp: the number of single-nucleotide polymorphism. SE: standard error. $p < 0.05$ represents the causal association of IL-1A, IL-1B, IL-1R1, IL-1Ra, and IL-1Racp levels with COVID-19. OR: Odds ratio. OR_lci95: Lower limit of 95% confidence interval for OR. OR_uci95: Upper limit of 95% confidence interval for OR.