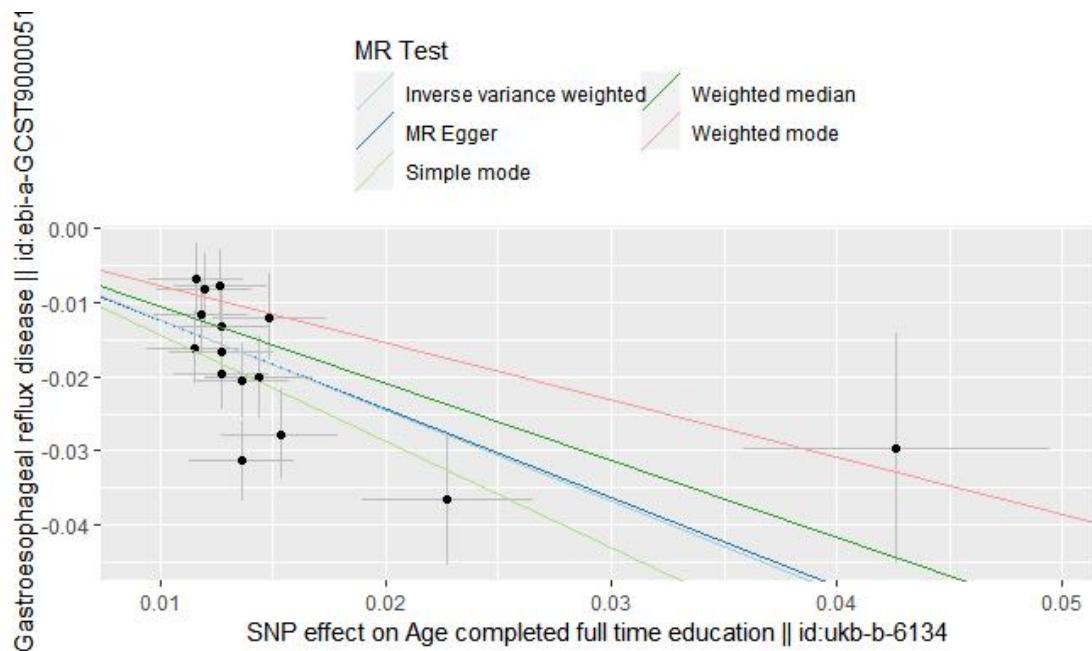
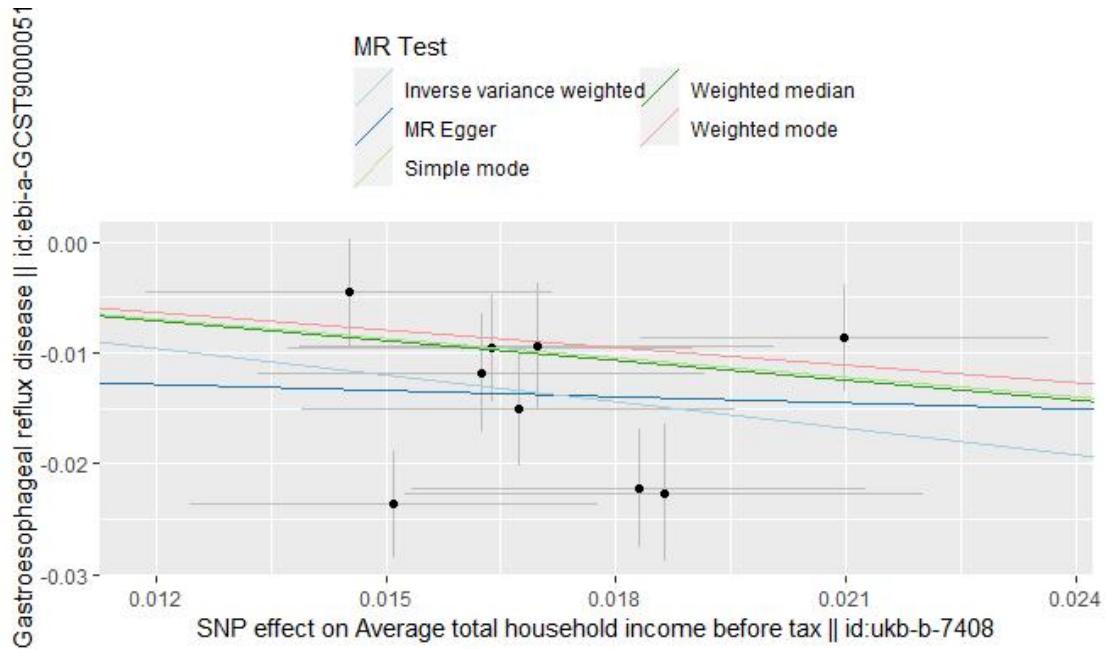


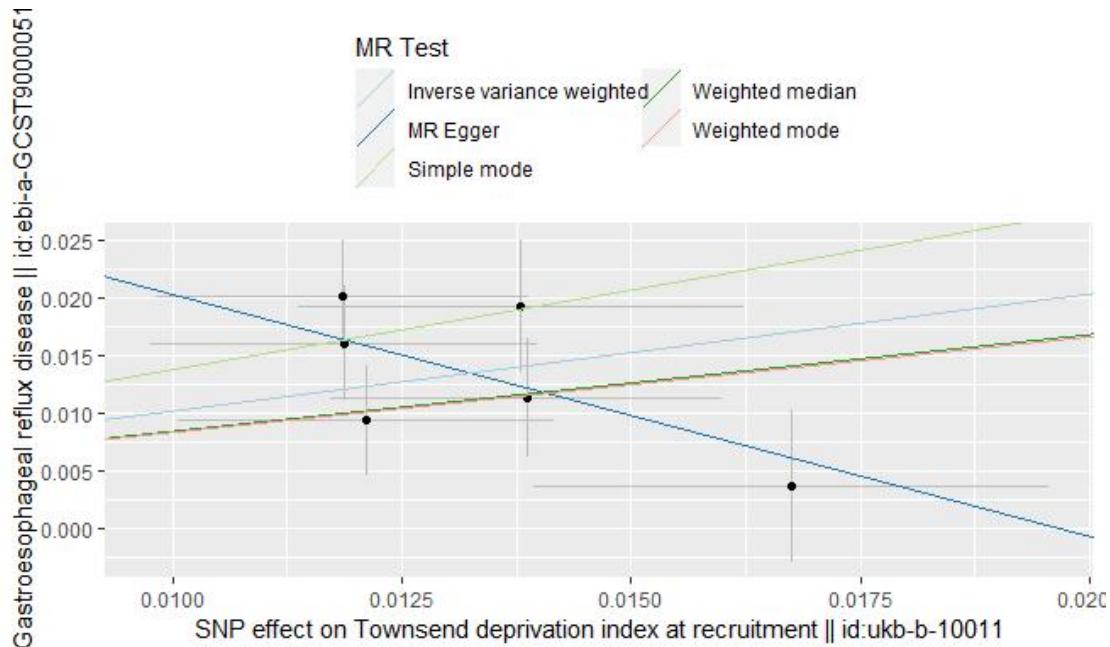
- **Supplementary Figures**

Supplementary Figure 1-12 Scatter plot figures

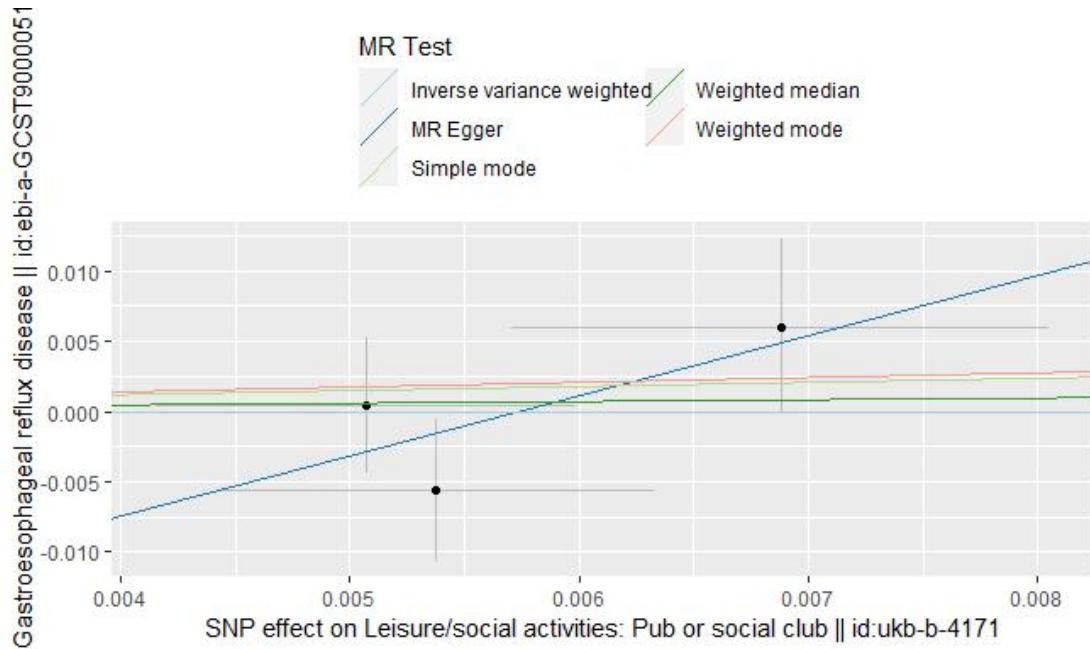




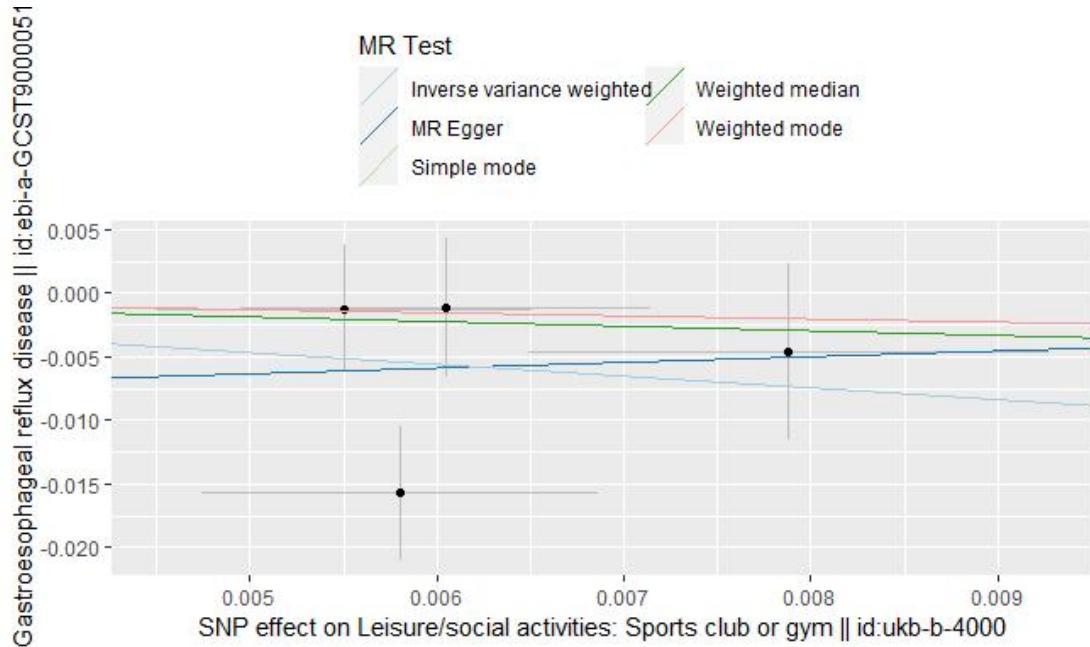
Supplementary Figure 3. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



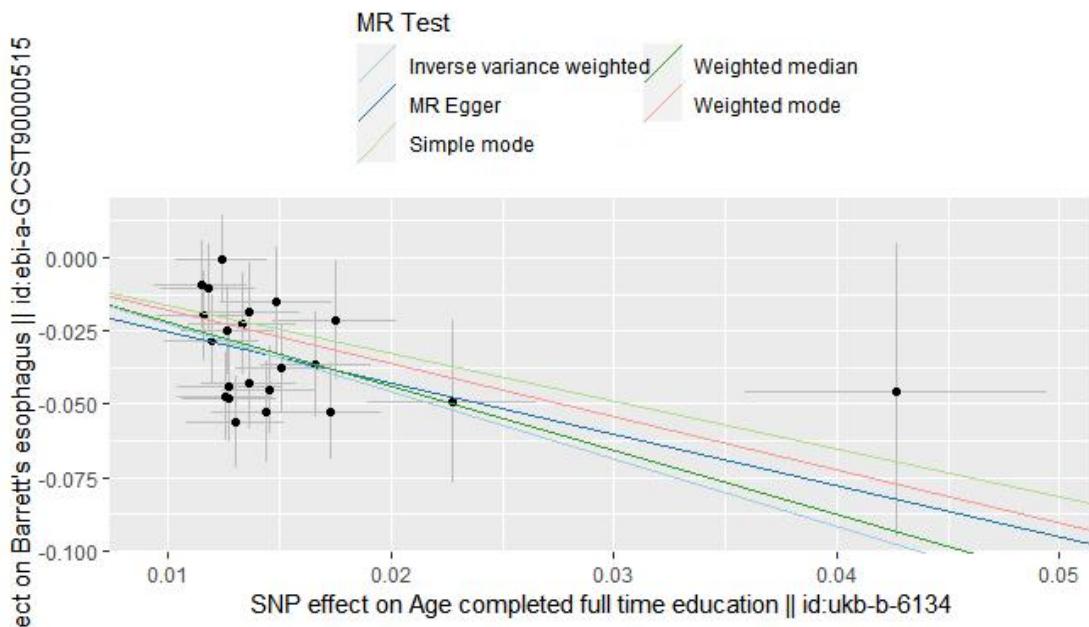
Supplementary Figure 4. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



Supplementary Figure 5. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



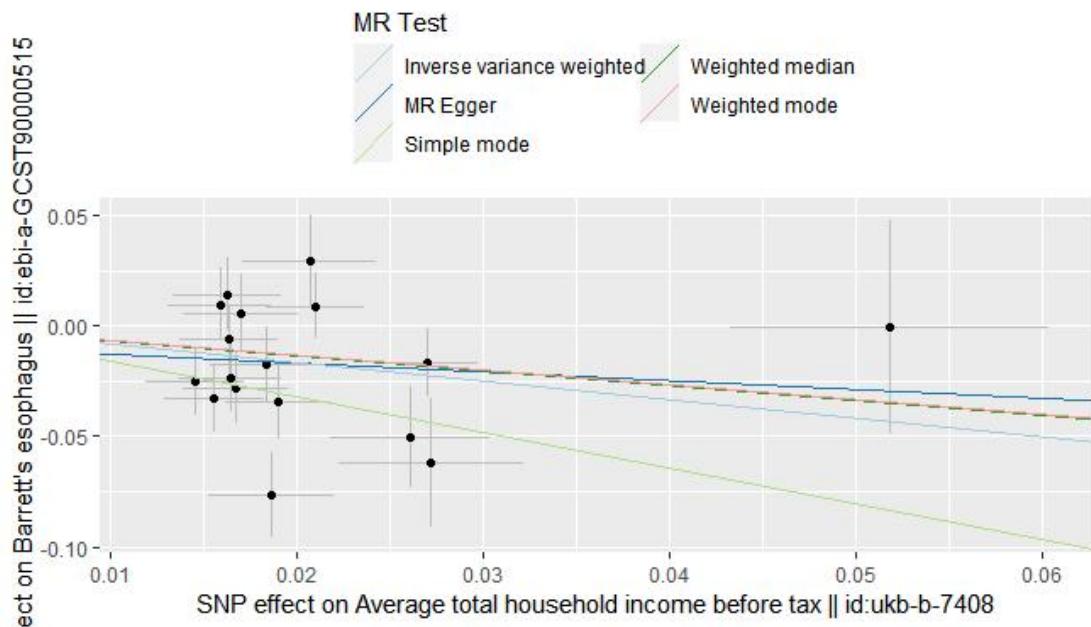
Supplementary Figure 6. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



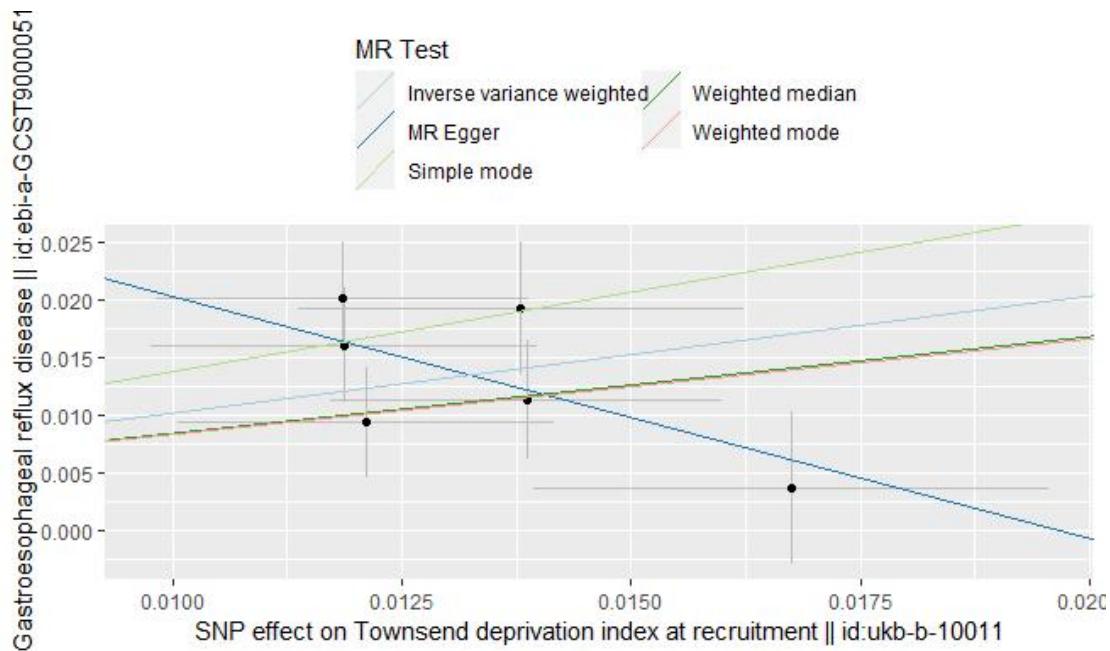
Supplementary Figure 7. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



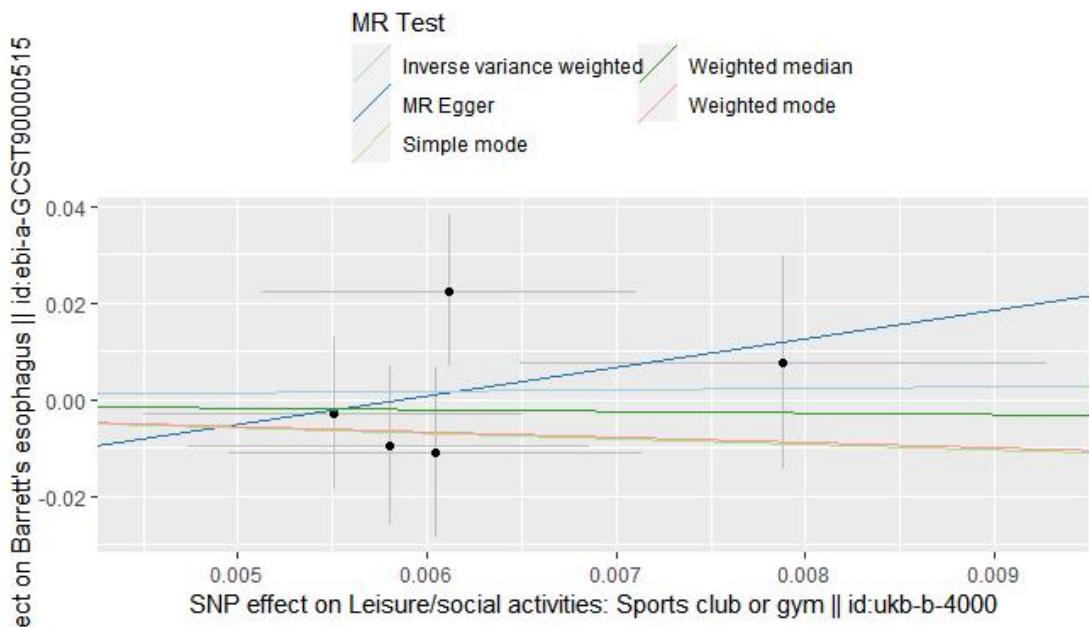
Supplementary Figure 8. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



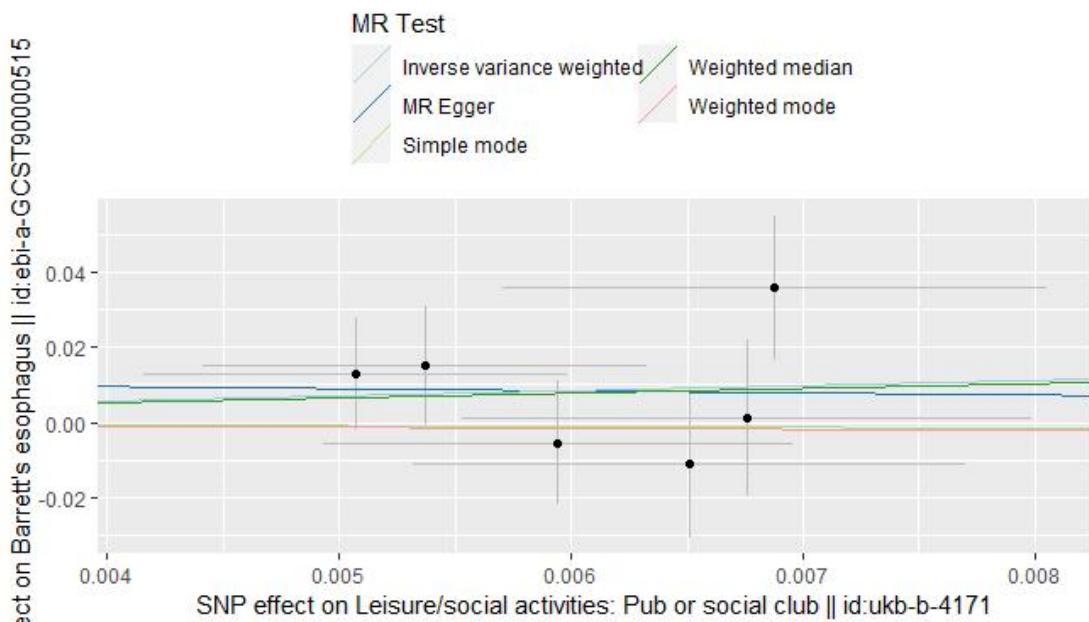
Supplementary Figure 9. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.



Supplementary Figure 10. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.

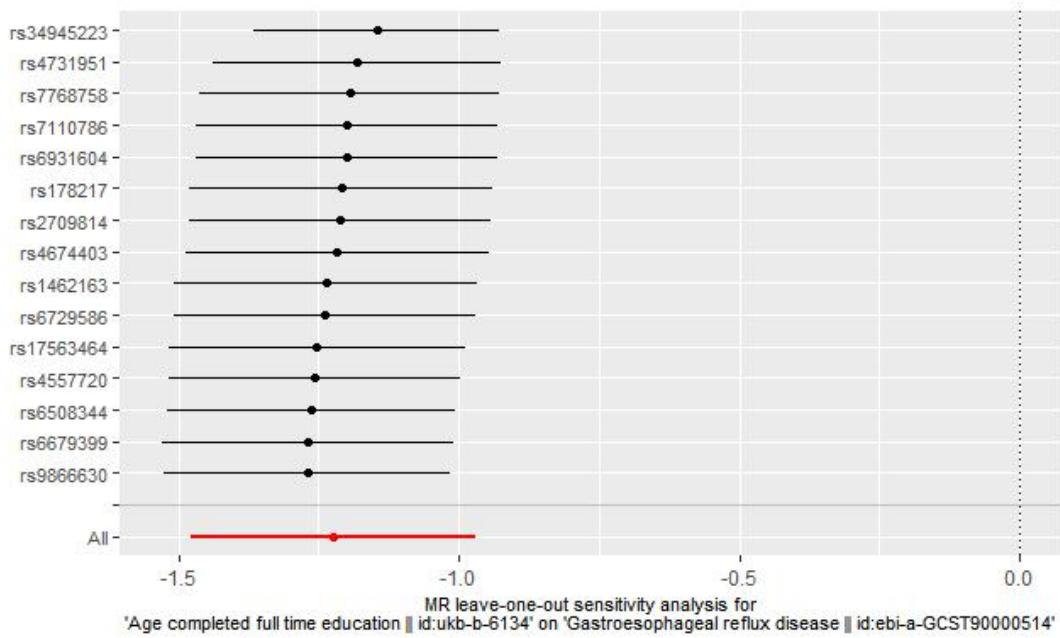


Supplementary Figure 11. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.

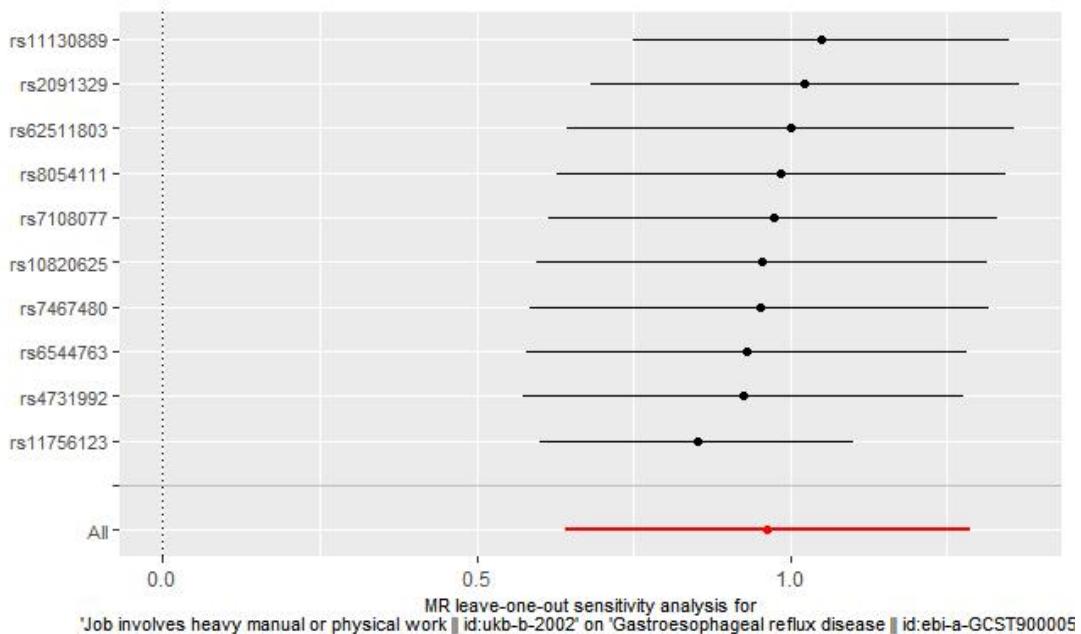


Supplementary Figure 12. SNP, single-nucleotide polymorphism; MR, Mendelian Randomization.

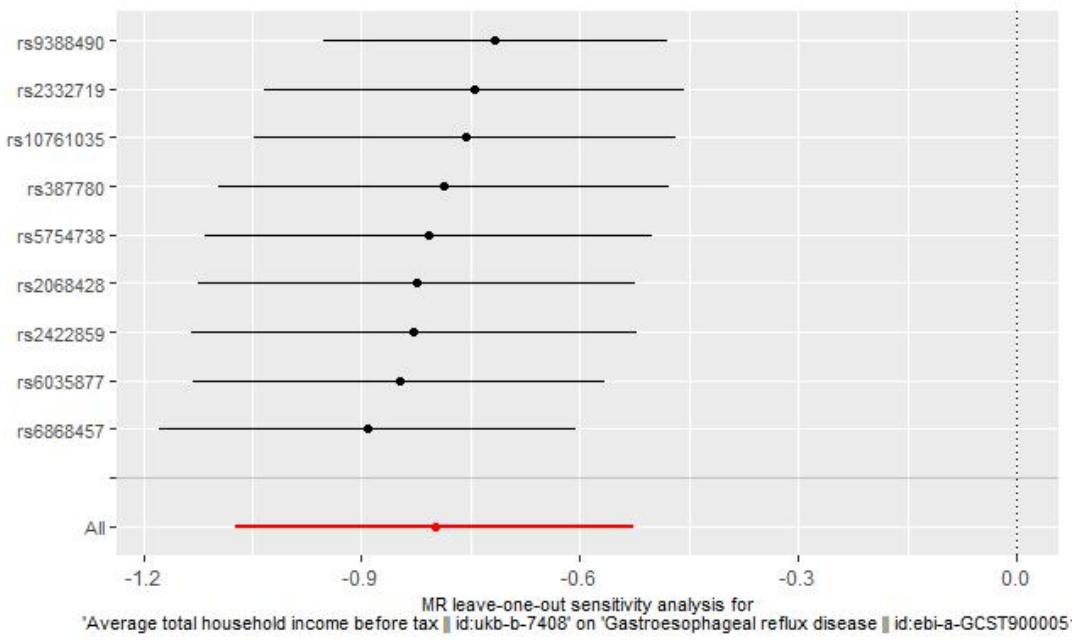
Supplementary Figure 13-24 leave-one-out sensitivity analysis



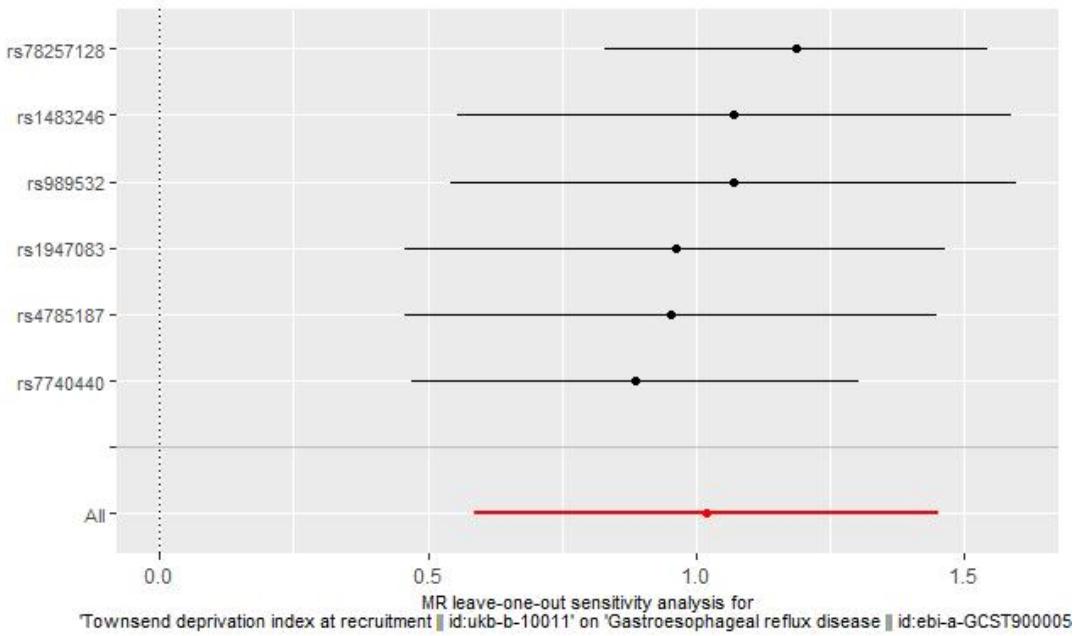
Supplementary Figure 13; MR, Mendelian Randomization.



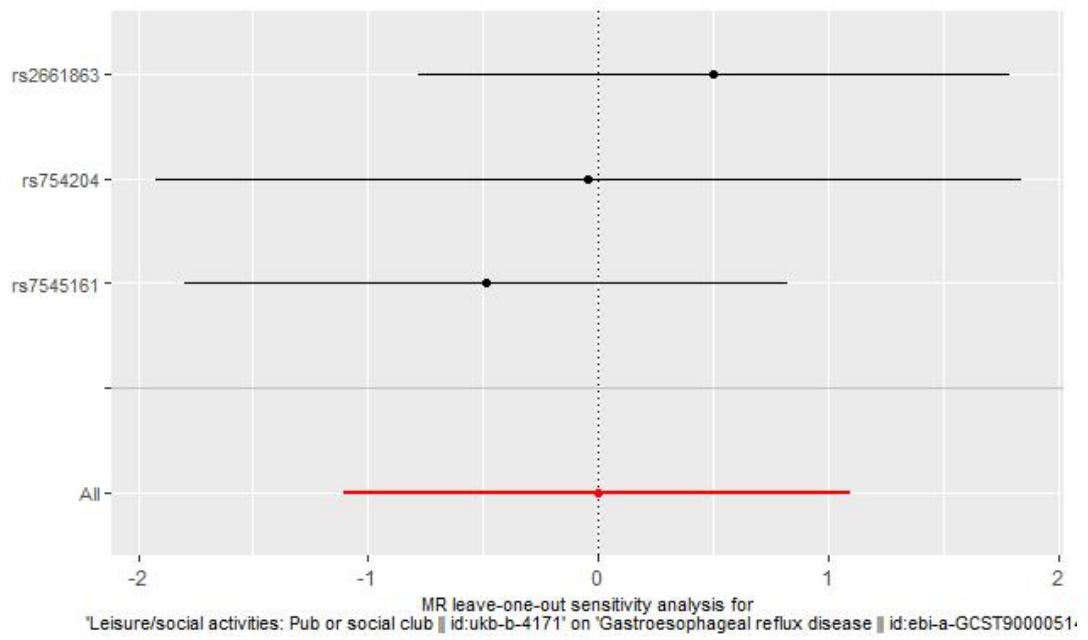
Supplementary Figure 14; MR, Mendelian Randomization.



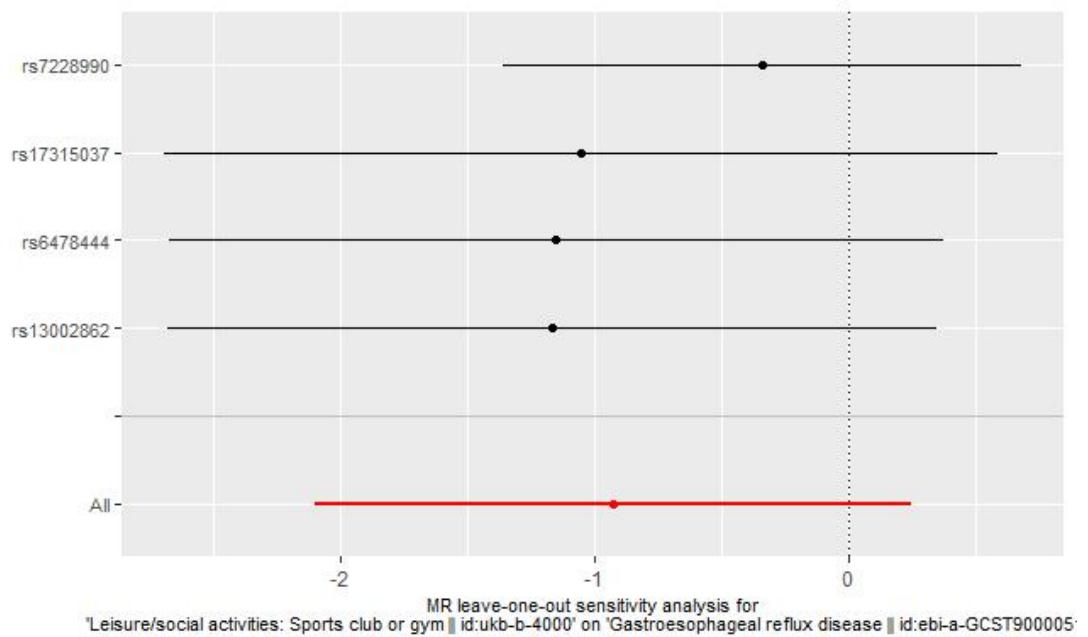
Supplementary Figure 15; MR, Mendelian Randomization.



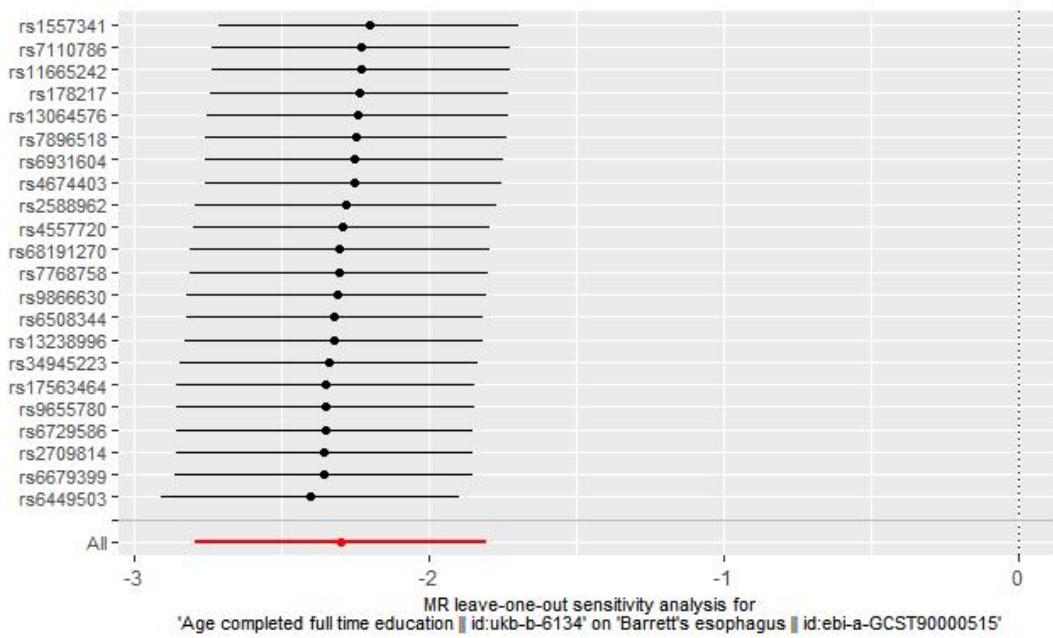
Supplementary Figure 16; MR, Mendelian Randomization.



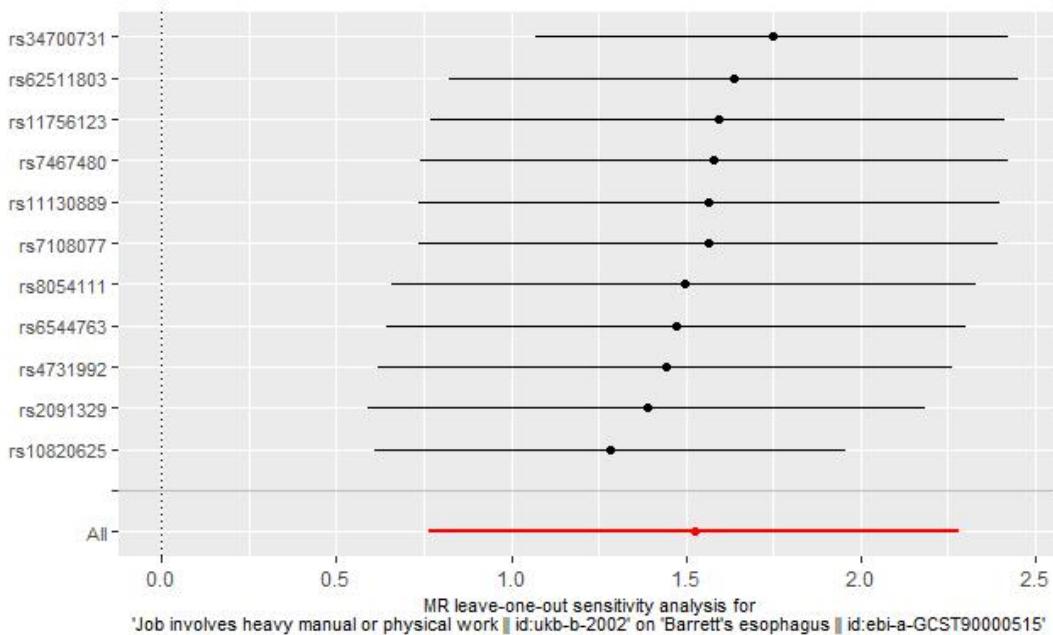
Supplementary Figure 17; MR, Mendelian Randomization.



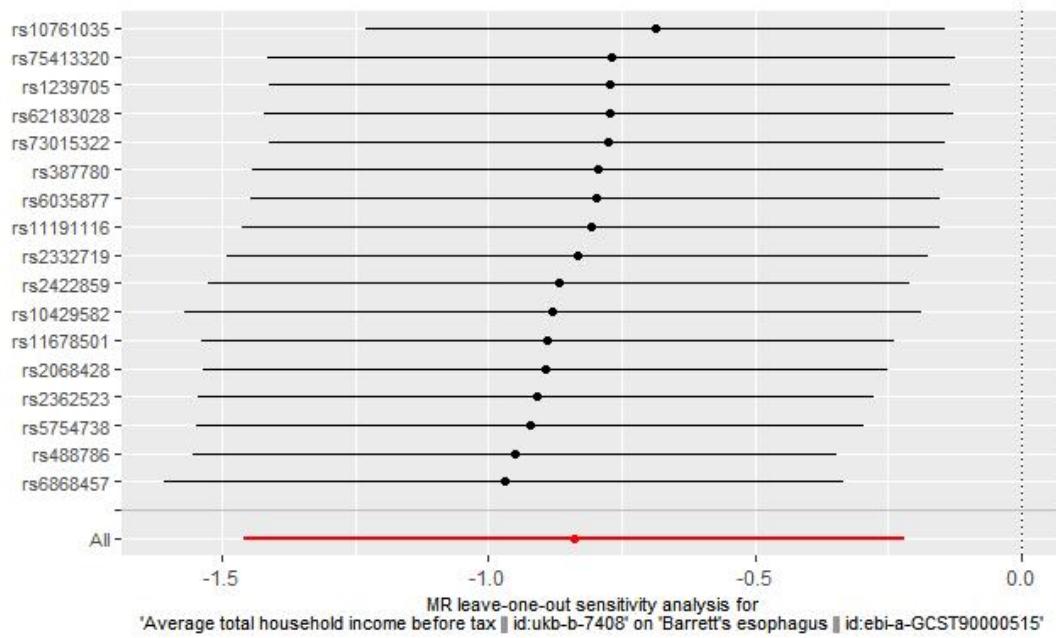
Supplementary Figure 18; MR, Mendelian Randomization.



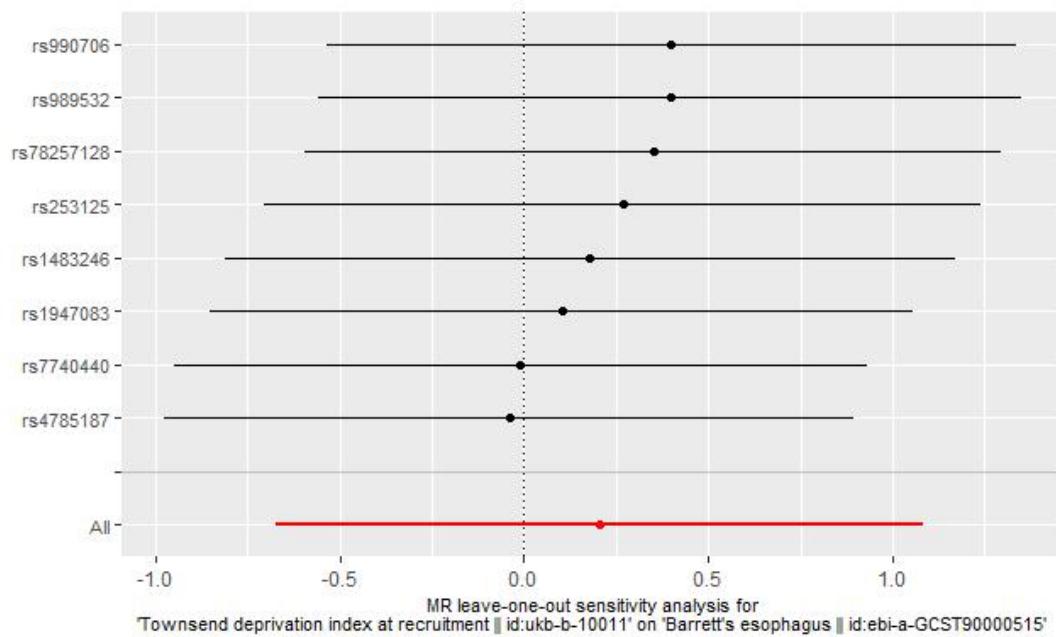
Supplementary Figure 19; MR, Mendelian Randomization.



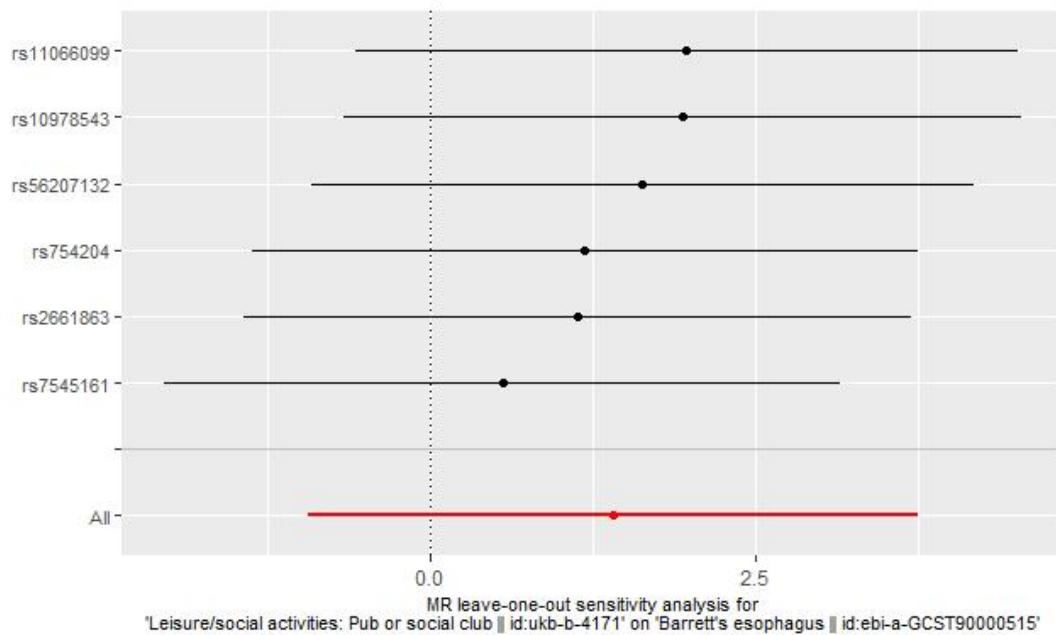
Supplementary Figure 20; MR, Mendelian Randomization.



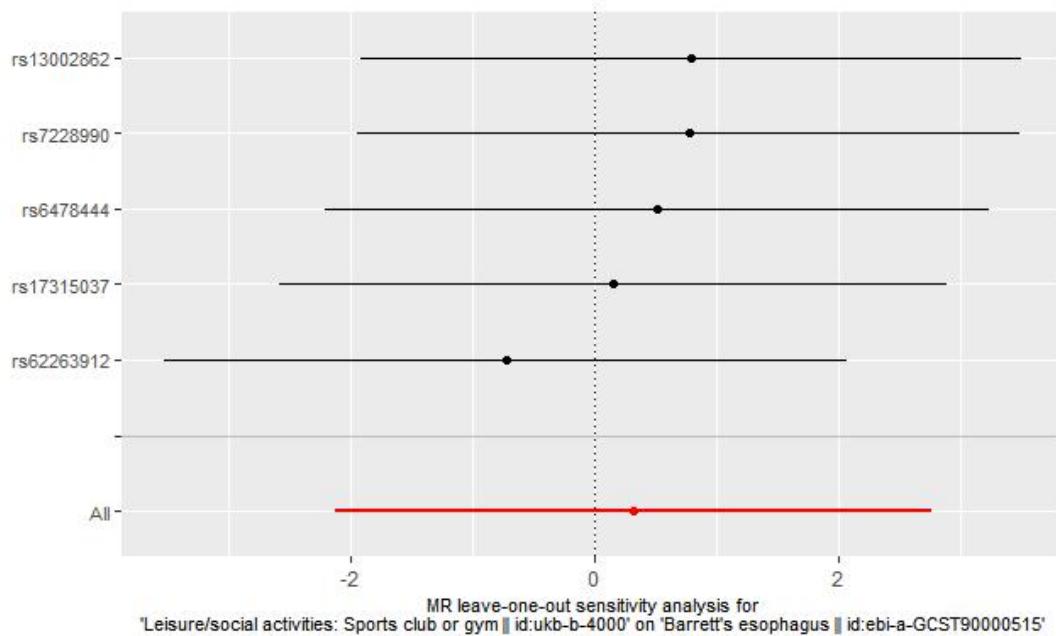
Supplementary Figure 21; MR, Mendelian Randomization.



Supplementary Figure 22; MR, Mendelian Randomization.

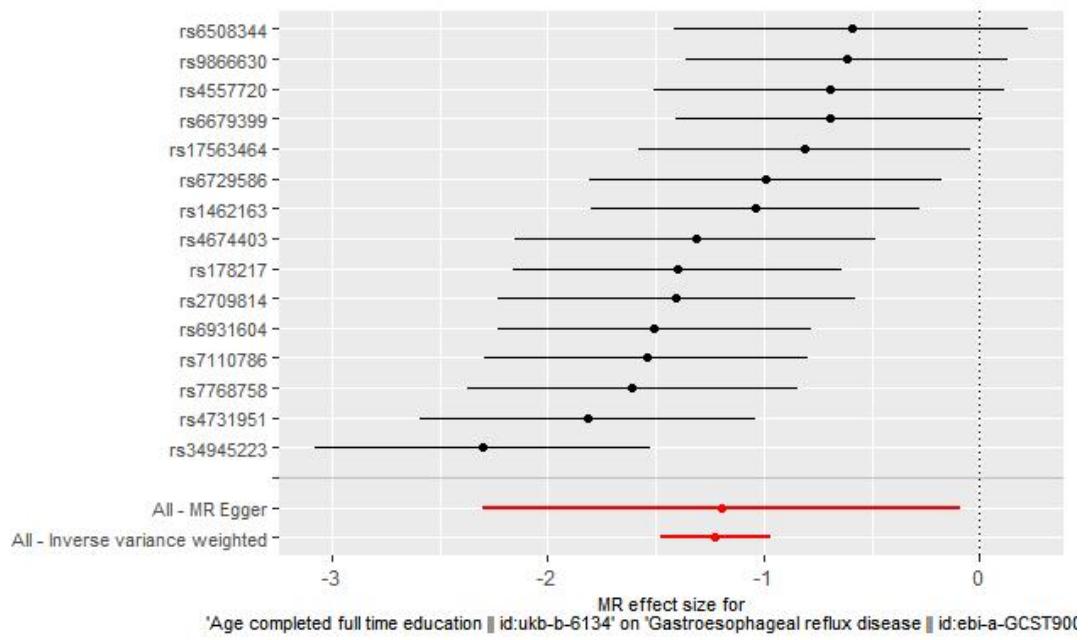


Supplementary Figure 23; MR, Mendelian Randomization.

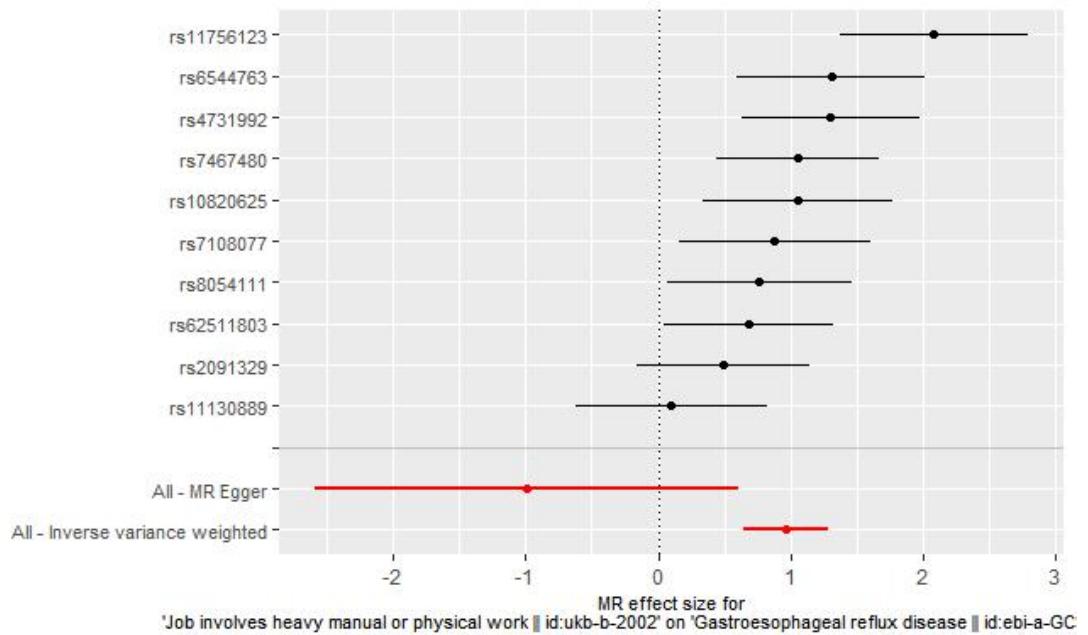


Supplementary Figure 24; MR, Mendelian Randomization.

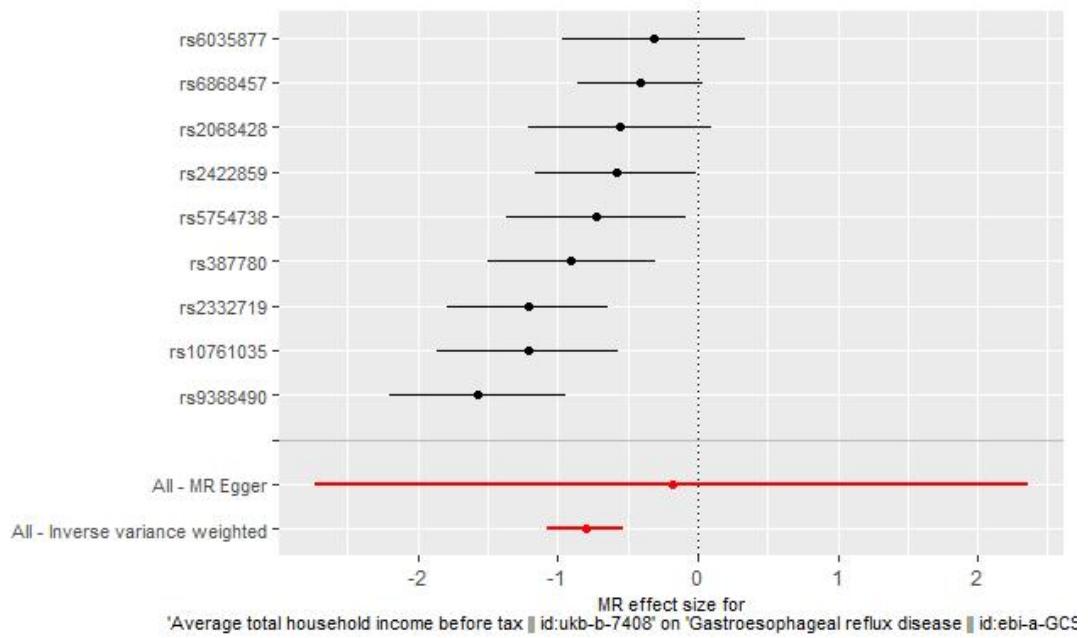
Supplementary Figure 25-36 Forest plot figure



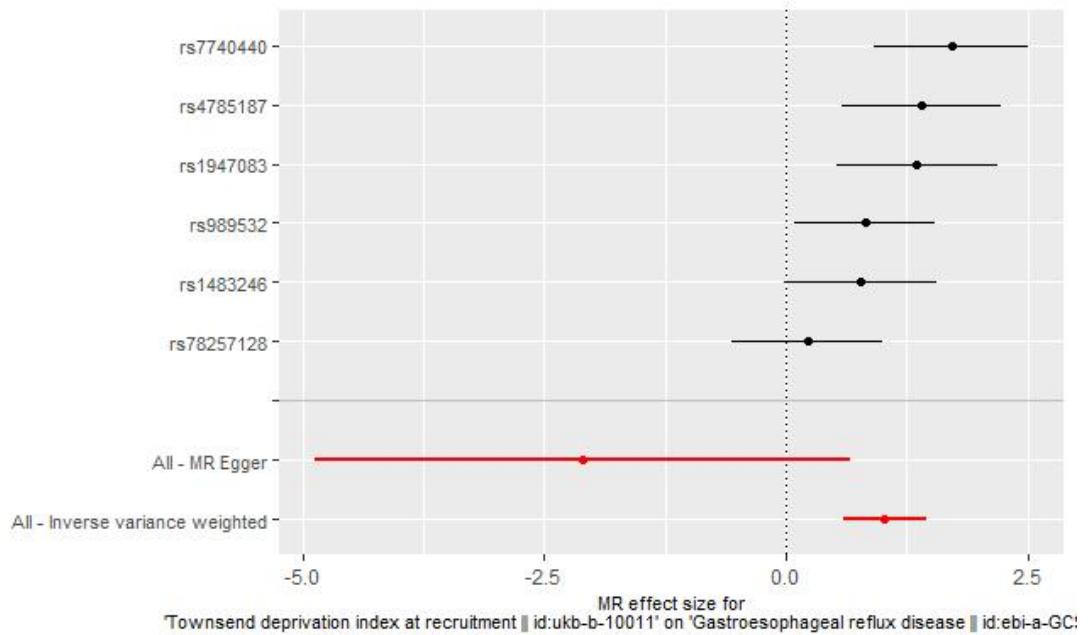
Supplementary Figure 25. MR, Mendelian Randomization.



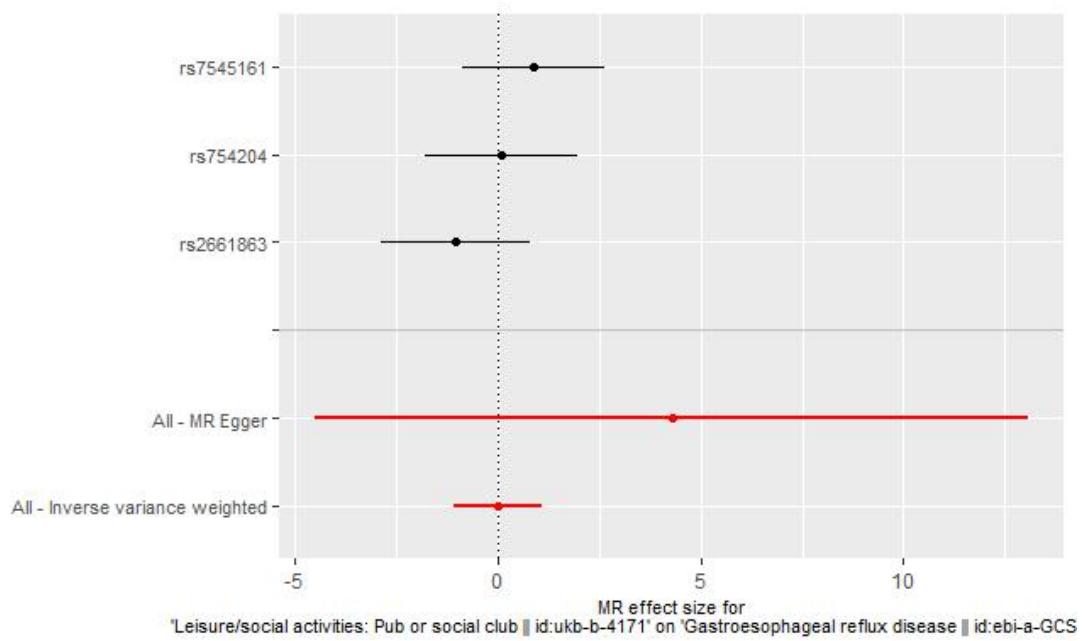
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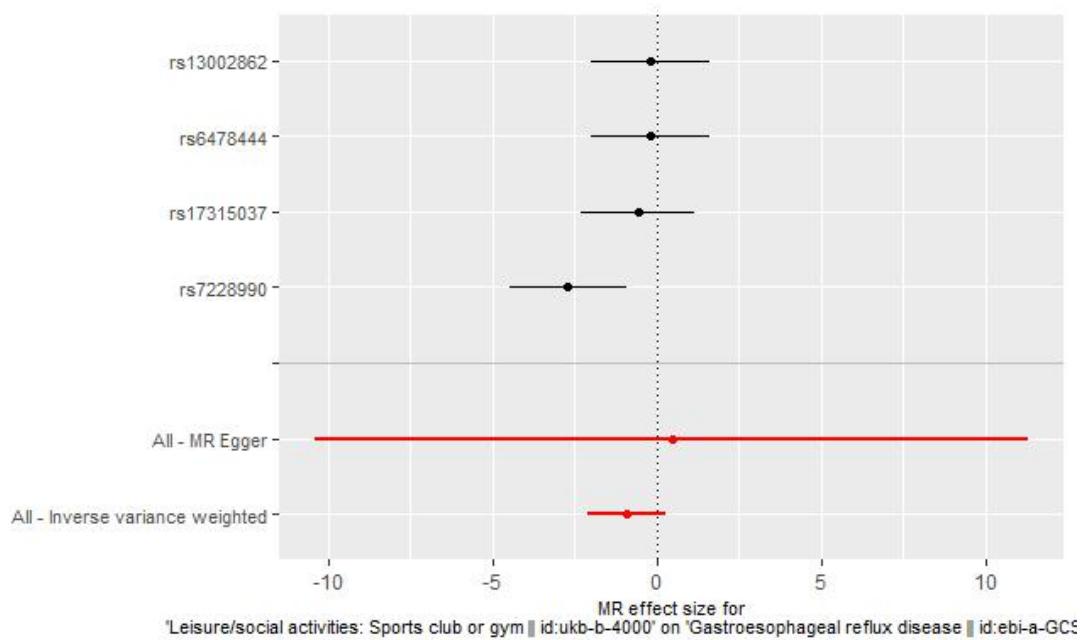
Supplementary Figure 27. MR, Mendelian Randomization.



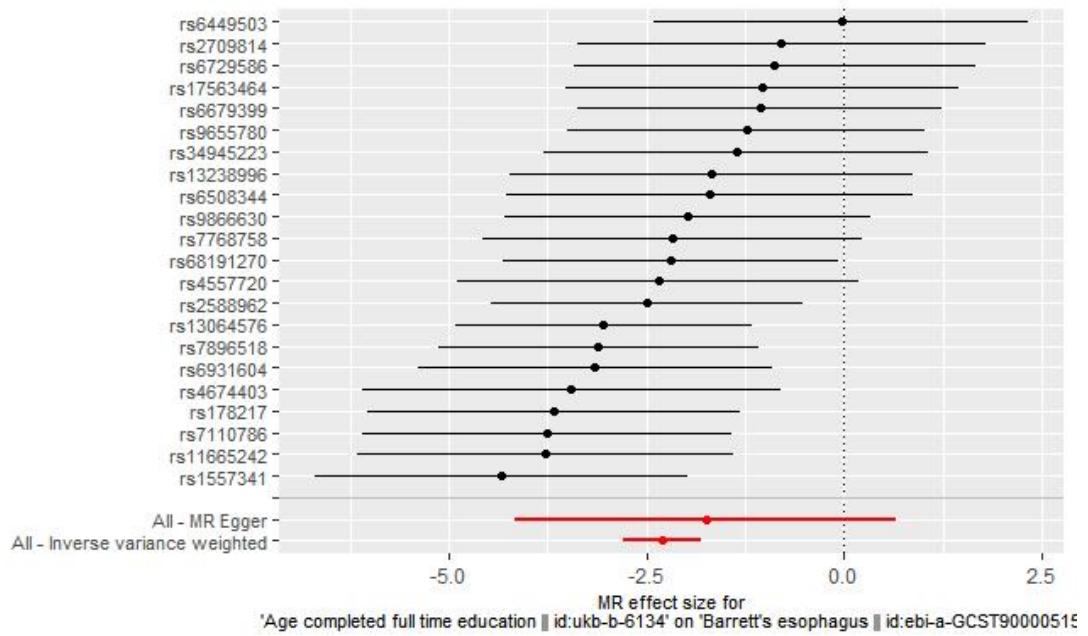
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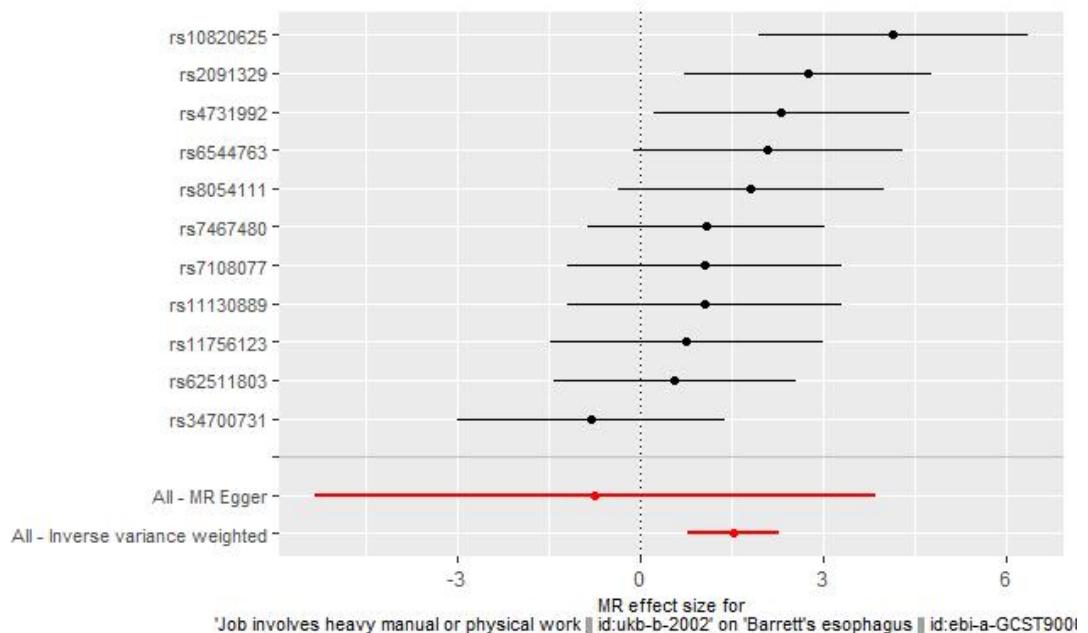
Supplementary Figure 29. MR, Mendelian Randomization.



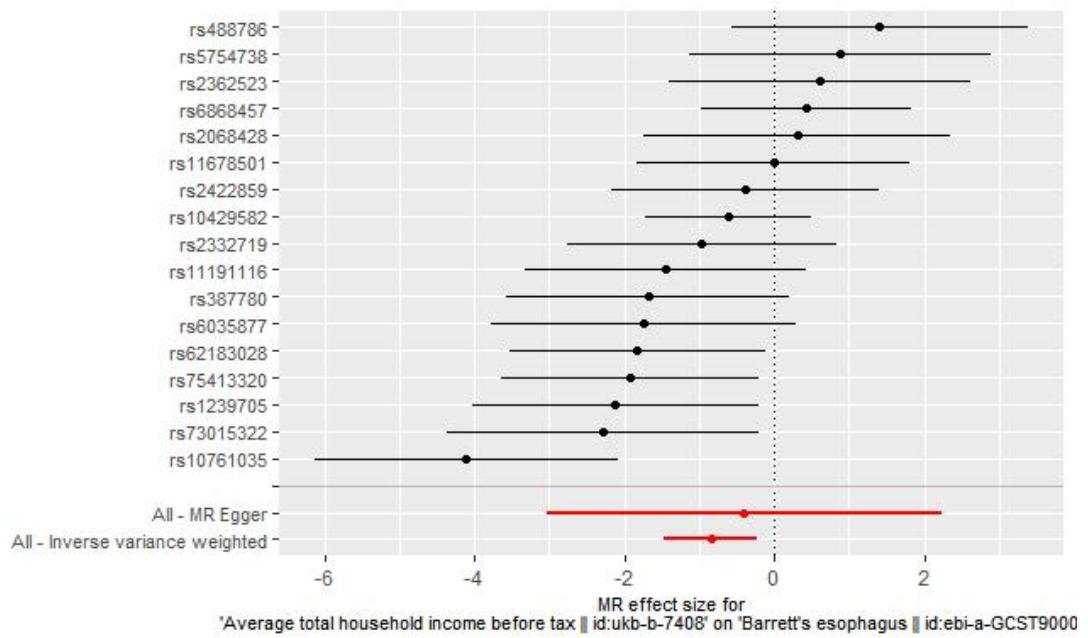
Supplementary Figure 30. MR, Mendelian Randomization.



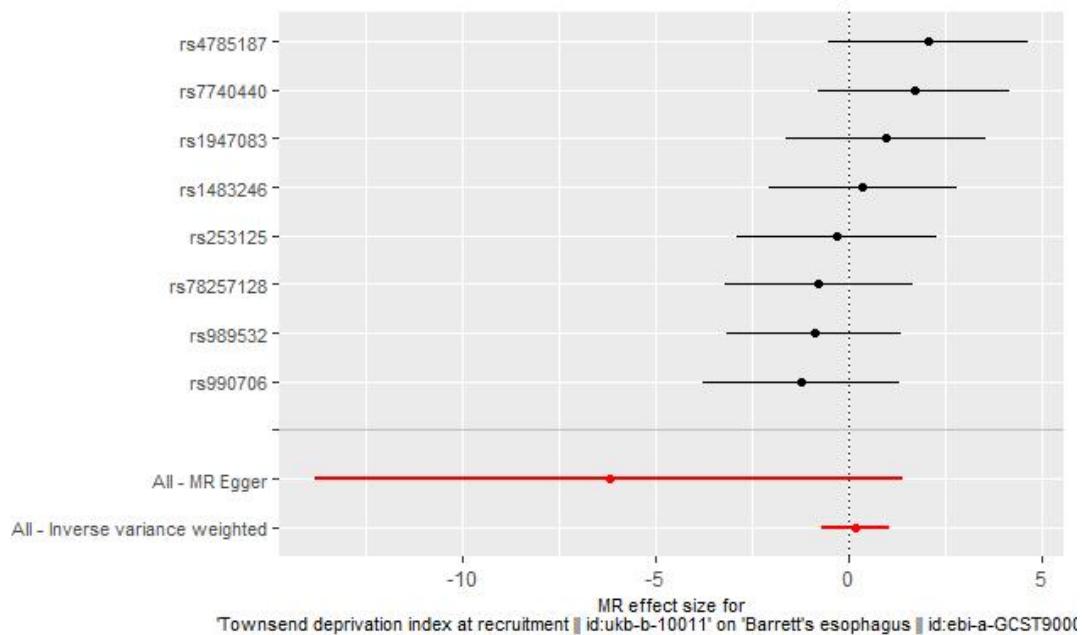
Supplementary Figure 31. MR, Mendelian Randomization.



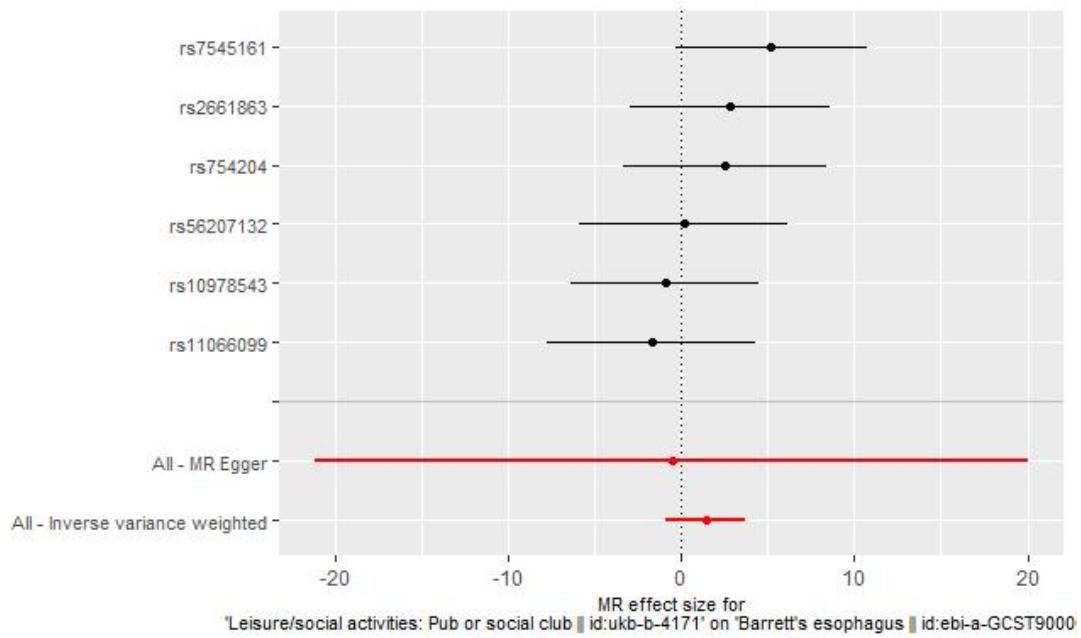
Supplementary Figure 32. MR, Mendelian Randomization.



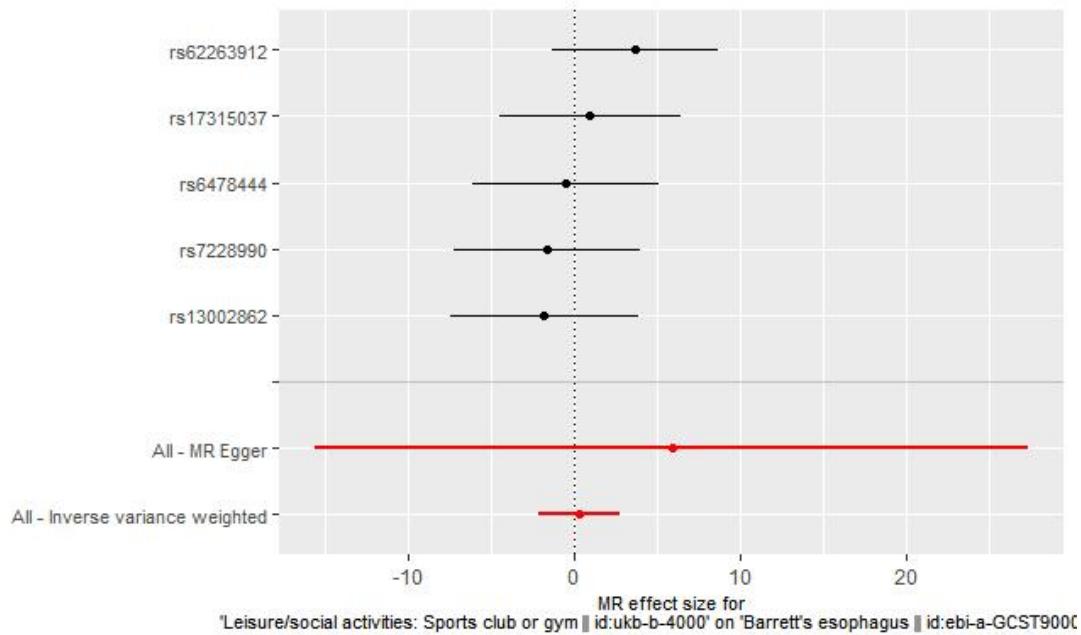
Supplementary Figure 33. MR, Mendelian Randomization.



Supplementary Figure 34. MR, Mendelian Randomization.

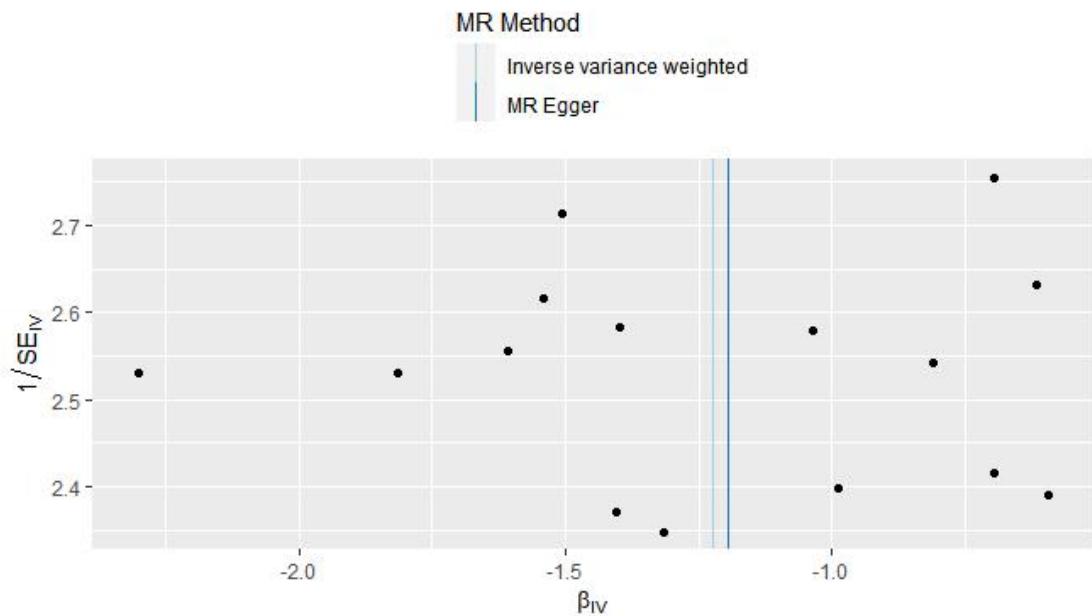


Supplementary Figure 35. MR, Mendelian Randomization.

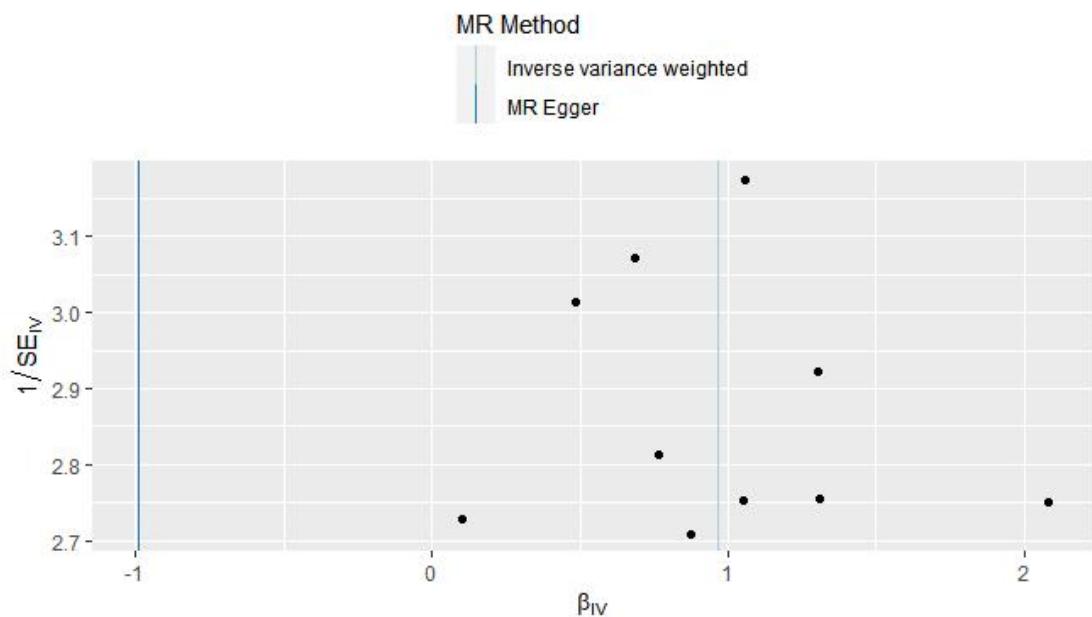


Supplementary Figure 36. MR, Mendelian Randomization.

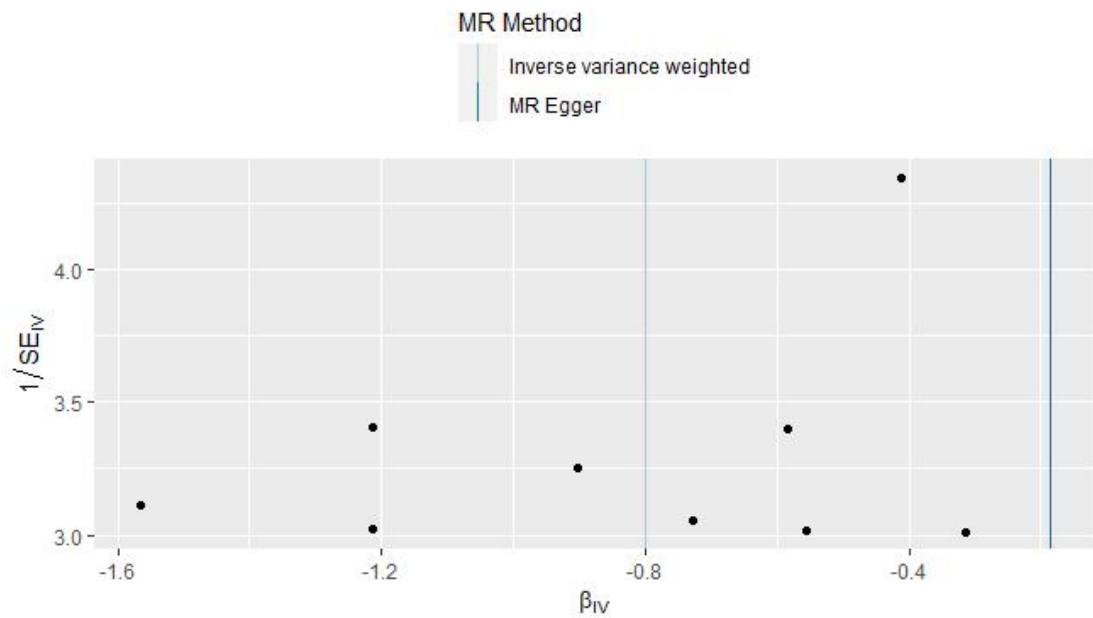
Supplementary Figure 37-48 Funnel plot figure



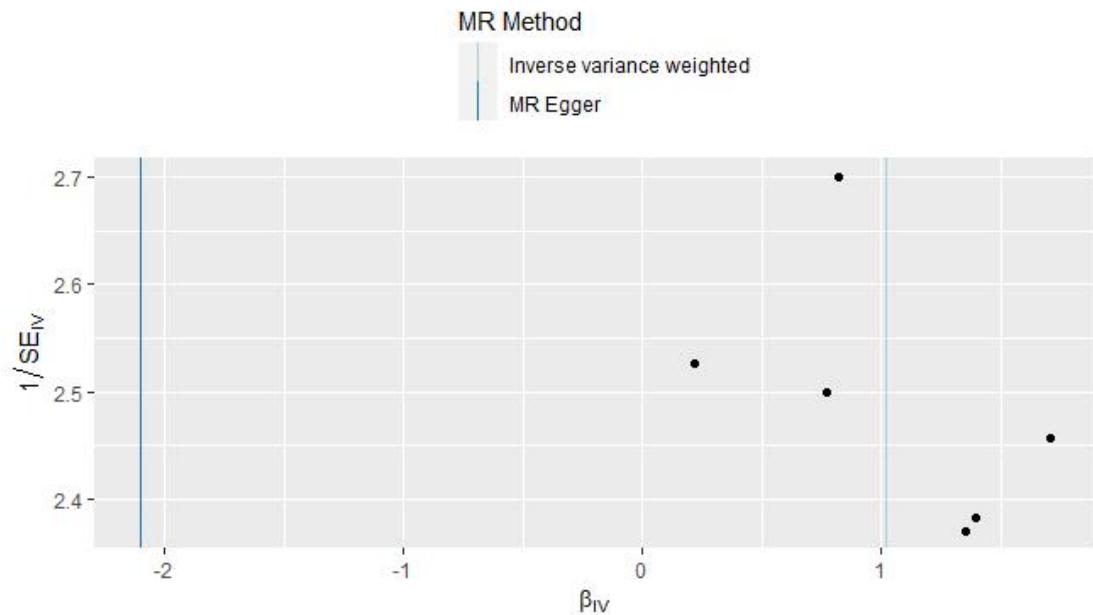
Supplementary Figure 37. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



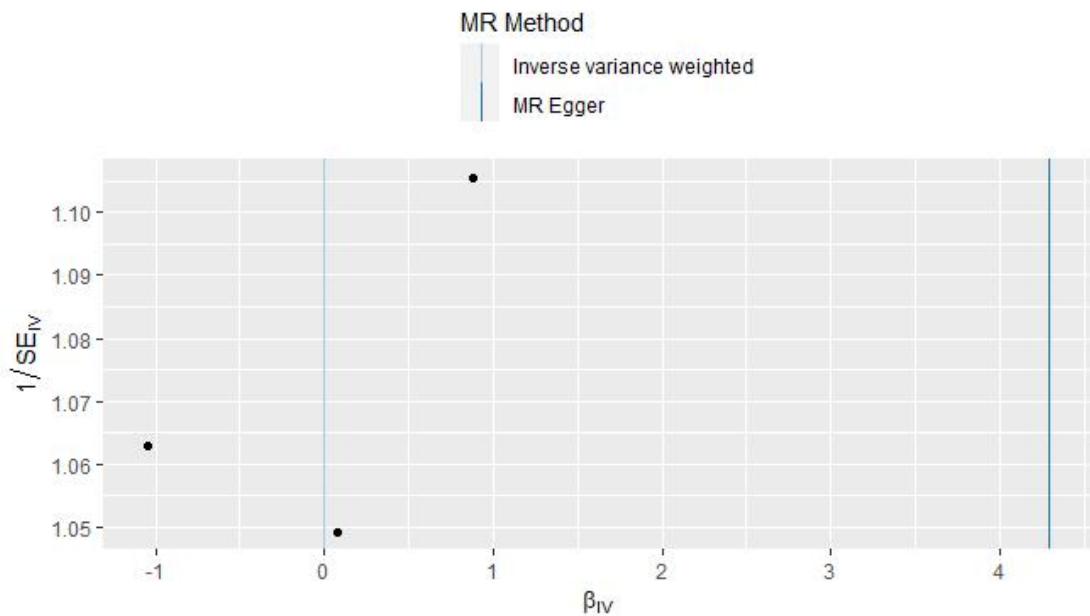
Supplementary Figure 38. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



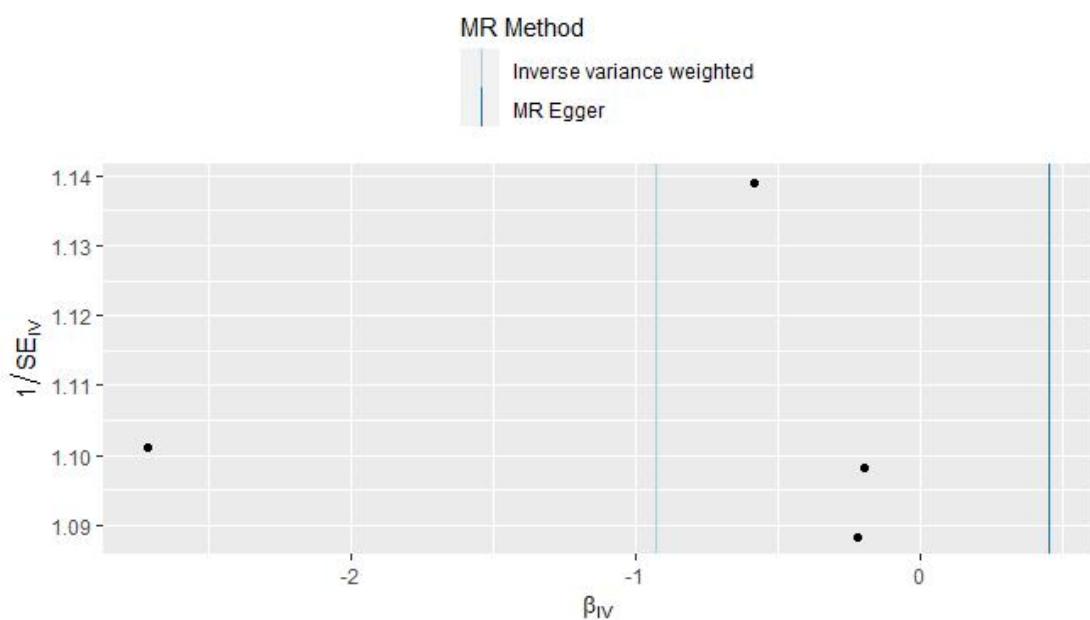
Supplementary Figure 39. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



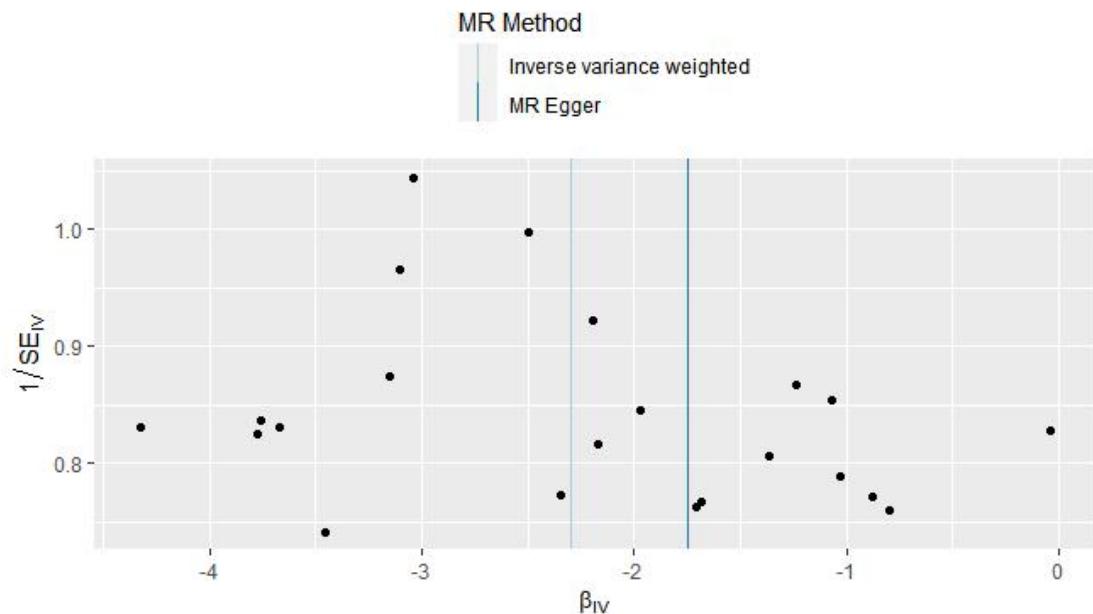
Supplementary Figure 40. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



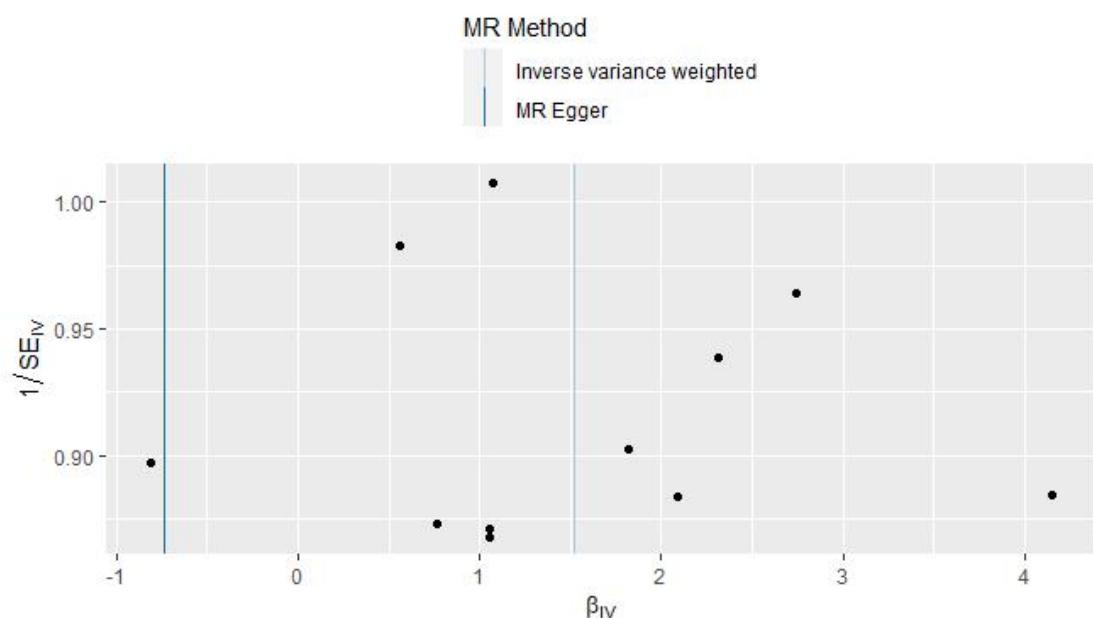
Supplementary Figure 41. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



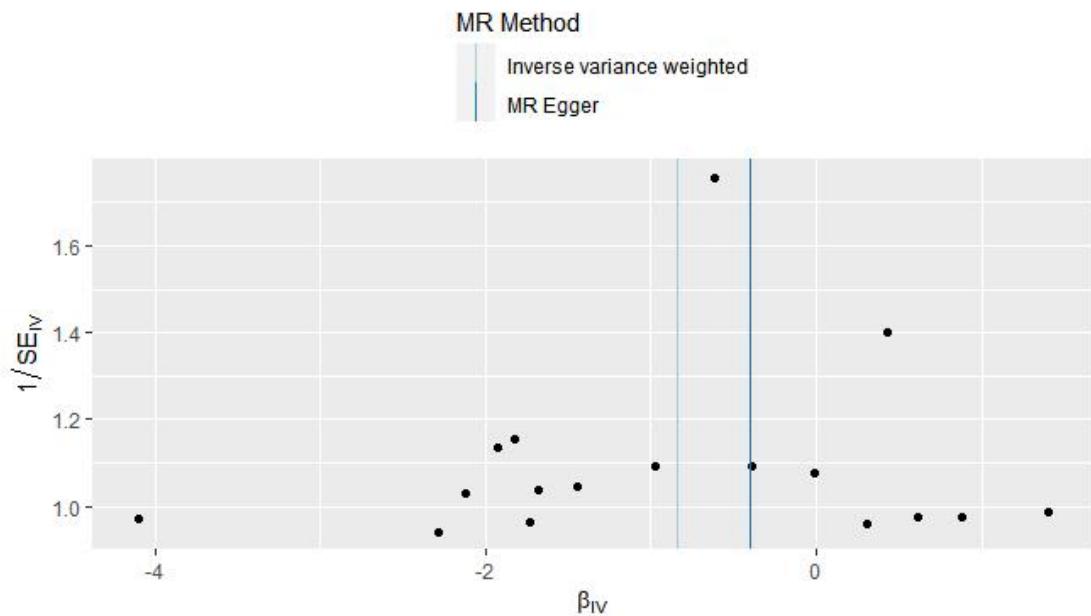
Supplementary Figure 42. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



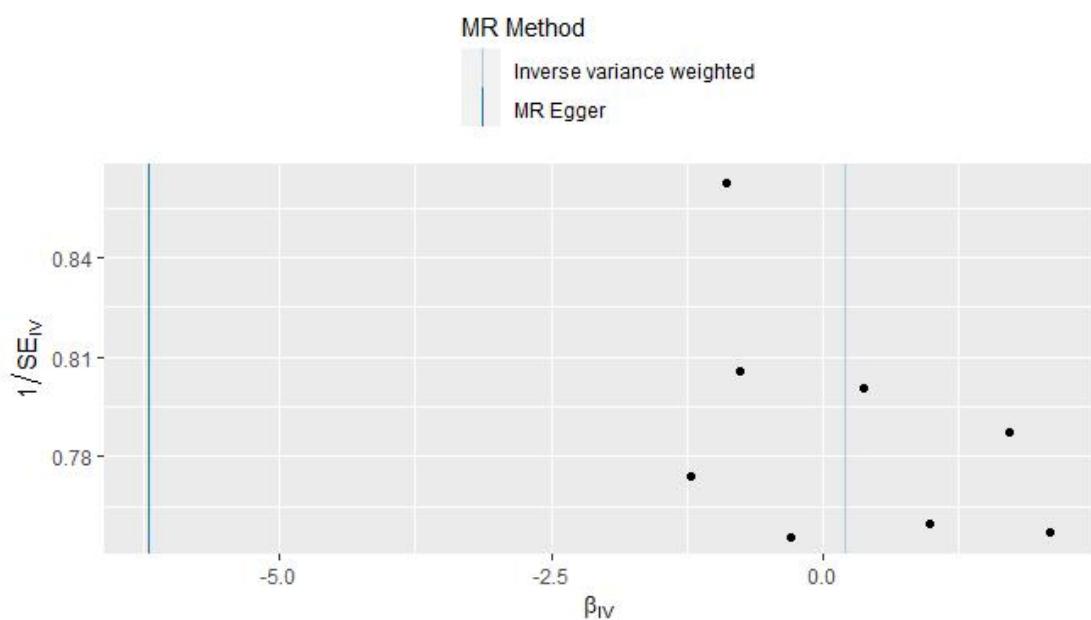
Supplementary Figure 43. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



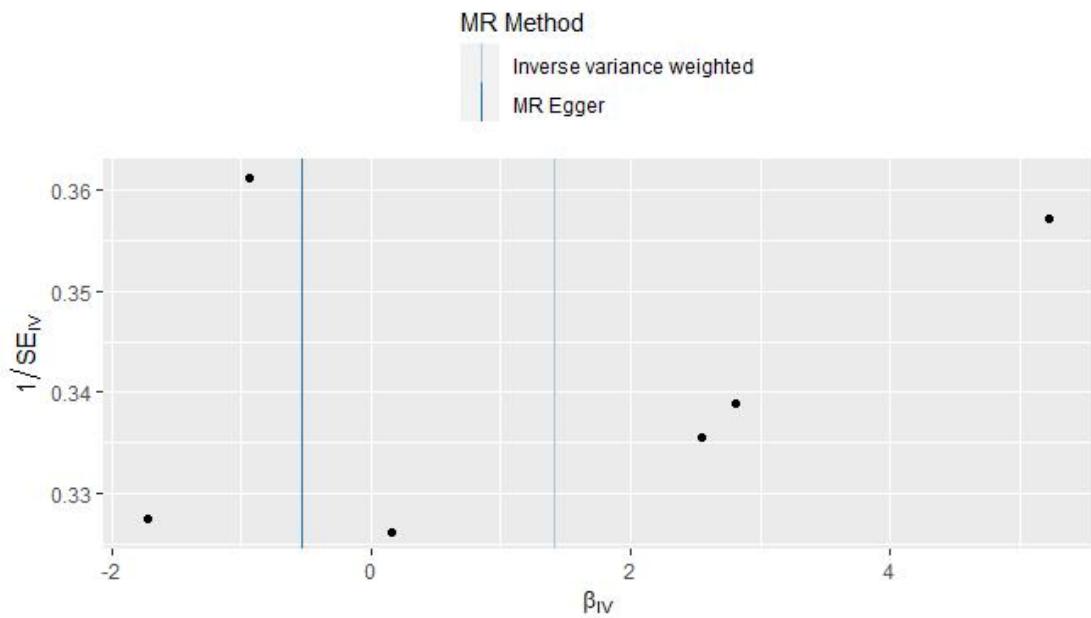
Supplementary Figure 44. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



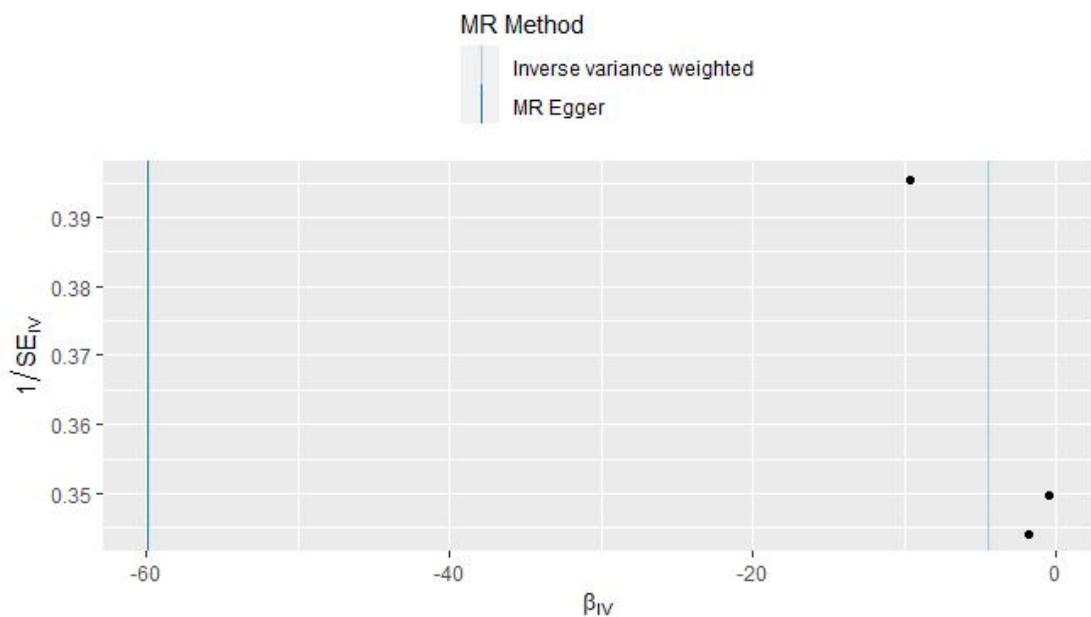
Supplementary Figure 45. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



Supplementary Figure 46. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



Supplementary Figure 47. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.



Supplementary Figure 48. SNP, single-nucleotide polymorphism; IV, Instrumental variable; SE: Standard Error; MR, Mendelian Randomization.

Supplementary Material

- **Supplementary Tables**

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Supplementary Table 1 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Age completed full time education and risk of Gastroesophageal reflux disease.

	SNP	Related confounding phenotypes	effect_allele	other_allele	effect_all_ele.outcome	other_all_ele.outcome	beta.exposure	beta.outcome	se.outcome	samplesize.outcome	samplesize.exposure	pval.effect_exposure	se.exposure	exp.osure	Age.complet.ed.full.time.education
1	rs10189857	BMI	G	A	G	A	-0.0124054	0.0203421	0.00485316	602604	Gastroesophageal reflux disease	307897	2.70E-09	0.00208474	Age.complet.ed.full.time.education
2	rs10760199	BMI	T	G	T	G	0.0115963	-0.0186891	0.00483678	602604	Gastroesophageal reflux disease	307897	2.50E-08	0.00208033	Age.complet.ed.full.time.education

3	rs109 5376	BMI 5	A	G	A	G	-0.0130 933	0.0257 032	0.004 8883 9	602604	Gastroesophageal reflux disease	307897	4.90E- 10	0.002 104
4	rs116 6524 2	ALCO HOL	G	A	G	A	-0.0125 523	0.0317 599	0.004 8818 7	602604	Gastroesophageal reflux disease	307897	2.10E- 09	0.002 09622
5	rs130 6457 6	BMI	T	C	T	C	0.0172 696	-0.042 7774	0.005 2936 8	602604	Gastroesophageal reflux disease	307897	2.40E- 14	0.002 26489

6	rs132 3899 6	BMI	G	A	G	A	-0.0133 142	0.0259 569	0.005 4862 8	602604	Gastroesophageal reflux disease	307897	3.20E- 08	0.002 40789
7	rs146 2163		T	C	T	C	-0.0127 222	0.0131 675	0.004 9340 9	602604	Gastroesophageal reflux disease	307897	1.90E- 09	0.002 11969
8	rs155 7341	ALCO HOL	C	A	C	A	0.0129 858	-0.034 9517	0.005 0292 5	602604	Gastroesophageal reflux disease	307897	3.70E- 09	0.002 20266

9	rs175 6346 4	A	C	A	C	-0.0148 391	0.0120 229	0.005 8379 8	602604	Gastroesophageal reflux disease	307897	3.50E- 09	0.002 51248	
1 0	rs178 217	C	T	C	T	-0.0143 669	0.0200 742	0.005 5639	602604	Gastroesophageal reflux disease	307897	2.10E- 09	0.002 39746	
1 1	rs258 8962	SMO KING	A	G	A	G	0.0150 825	-0.019 3965	0.004 8312 7	602604	Gastroesophageal reflux disease	307897	4.10E- 13	0.002 07972

1	rs270	C	T	C	T	-0.0114 761	0.0161 186	0.004 8384 4	602604	Gastroesophageal reflux disease	307897	4.20E- 08	0.002 0926
2	9814												
1	rs349	G	A	G	A	-0.0136 288	0.0313 79	0.005 3874 1	602604	Gastroesophageal reflux disease	307897	3.40E- 09	0.002 30526
3	4522 3												
1	rs455	G	A	G	A	0.0119 687	-0.008 31899	0.004 9532 4	602604	Gastroesophageal reflux disease	307897	2.00E- 08	0.002 13393
4	7720												

1	rs650	A	G	A	G	-0.0115 957	0.0068 7788	0.004 8491 7	602604	Gastroesophageal reflux disease	307897	2.80E-08	0.00208724
8	8344												
1	rs667	C	A	C	A	0.0426 608	-0.029 6751	0.015 4884	602604	Gastroesophageal reflux disease	307897	3.00E-10	0.00677016
9	9399												
2	rs672	A	G	A	G	-0.0117 953	0.0116 753	0.004 9163 2	602604	Gastroesophageal reflux disease	307897	2.40E-08	0.00211341
0	9586												

2	rs681 9127 0	SMO KING	C	A	C	A	0.0166 103	-0.021 12	0.005 7590 2	602604	Gastroesophageal reflux disease	307897	2.50E- 11	0.002 48918
2	rs693 1604		T	C	T	C	0.0136 265	-0.020 4973	0.005 0225 9	602604	Gastroesophageal reflux disease	307897	1.10E- 10	0.002 1131
2	rs711 0786		T	C	T	C	0.0127 141	-0.019 6005	0.004 8601 3	602604	Gastroesophageal reflux disease	307897	1.20E- 09	0.002 09121

2	rs965 7	5780	ALCO HOL	A	G	A	G	-0.0174 579	0.0217 935	0.006 4631 4	602604	Gastroesophageal reflux disease	307897	4.30E- 10	0.002 79646	Age com plet ed full time edu cati on Age com plet ed full time edu cati on
2	rs986 8	6630		A	G	A	G	-0.0126 597	0.0078 1251	0.004 8120 9	602604	Gastroesophageal reflux disease	307897	1.00E- 09	0.002 0719	Age com plet ed full time edu cati on

Supplementary Table 2 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Job involves heavy manual or physical work and risk of Gastroesophageal reflux disease.

Related confounding phenotypes															
SNP	effect_all_ele.exposure	other_all_ele.exposure	effect_all_ele.outcome	other_all_ele.outcome	beta.e xposure	beta.outcome	se.outcome	samplesize.outcome	pval.outcome	outcome	pvalue xposure	samplesize.exposure	se.exposure	exp osure	
rs1081158	Alcohol	T	C	T	C	0.0169887	-0.00013351	0.0048923	6026041	0.978228	Gastroesophageal reflux disease	2.70E-12	263615	0.00242996	
rs10820625		C	T	C	T	-0.0160068	-0.0168521	0.00581336	602604	0.00374516	Gastroesophageal reflux disease	2.90E-08	263615	0.00288515	

k

														Job invo lves heav
rs11														y man ual or phys ical wor k
3 1308		G	C	G	C	-0.024 9218	-0.002 52268	0.009 1374 4	602604	0.7824 85	Gastroesophag al reflux disease	4.50E- 08	263615	0.004 5572 5
rs11														Job invo lves heav
4 6638	Alcoh ol	A	C	A	C	0.0136 375	0.0318 018	0.004 9201 6	602604	1.02E- 10	Gastroesophag al reflux disease	2.50E- 08	263615	0.002 4480 6
														phys ical wor k

5	rs11 6789 79	BMI	C	T	C	T	-0.014 9934	-0.033 1273	0.005 4386 5	602604	1.12E- 09	Gastroesophag al reflux disease	3.00E- 08	263615 7063 7	0.002 y man ual or phys ical wor k Job invo lves heav
6	rs11 7267 86	Alcoh ol	G	T	G	T	0.0165 951	0.0057 5011	0.004 9864 6	602604	0.2488 51	Gastroesophag al reflux disease	2.40E- 11	263615 4834 4	0.002 y man ual or phys ical wor k Job invo lves heav
7	rs11 7561 23		T	A	T	A	-0.013 7486	-0.028 7505	0.004 9978 2	602604	8.79E- 09	Gastroesophag al reflux disease	3.20E- 08	263615 4865 8	0.002 lives heav y man ual

															or phys ical wor k
															Job invo lves heav
rs13															y man ual or phys ical wor k
8 2509 96	C	G	C	G	-0.013 3122	0.0098 0948	0.004 8185 2	602604	0.0417 715	Gastroesoph ageal reflux disease	2.70E- 08	263615	0.002 3930 7		Job invo lves heav
rs13															y man ual or phys ical wor k
9 7005 9	BMI	G	A	G	0.0180 816	0.0401 347	0.005 6884 8	602604	1.72E- 12	Gastroesoph ageal reflux disease	1.80E- 10	263615	0.002 8349 2		Job invo lves heav

															Job
1	rs20														invo
0	9132	G	A	G	A	-0.015 8895	-0.007 7522	0.005 2737	602604	0.1415 69	Gastroesophag	2.00E-09	263615	0.002 6500	lves
9											al reflux				heavy
											disease				y
															man
															ual
															or
															phys
															ical
															wor
															k
															Job
															invo
															lves
															heavy
1	rs23														y
1	1854	BMI	G	T	G	-0.014 2827	-0.025 5283	0.004 8790	602604	1.67E-07 5	Gastroesophag	3.70E-09	263615	0.002 4224	man
0											al reflux				ual
											disease				or
															phys
															ical
															wor
															k
															Job
															invo
1	rs28														lves
2	1934	BMI	C	T	C	0.0145 582	0.0252 013	0.005 0188	602604	5.13E-07 8	Gastroesophag	5.60E-09	263615	0.002 4973	heavy
8											al reflux				y
											disease				man
															ual

															or
															phys
															ical
															wor
															k
															Job
															invo
															lves
															heavy
1	rs34														y
3	7007	A	G	A	G	0.0197	-0.006	0.007	0.3834	Gastroesophag	1.50E-	0.003	4944	man	
	31					694	11605	0170	26	geal reflux	08	263615	5	ual	
								2		disease				or	
														phys	
														ical	
														wor	
														k	
														Job	
														invo	
														lves	
														heavy	
1	rs35														y
4	9991	BMI	G	C	G	-0.024	-0.042	0.005	3.00E-	Gastroesophag	7.40E-	0.002	5873	man	
	62					8598	6913	2233	16	geal reflux	22	263615	5	ual	
								2		disease				or	
														phys	
														ical	
														wor	
														k	

															Job
1	rs37	Alcoh	T	C	T	C	0.0135	0.0237	0.004	602604	1.50E-	Gastroesophag	3.70E-	0.002	invo
5	8535	ol					824	887	9450	9	06	al reflux	-08	263615	lves
	4											disease		0.4672	heavy
															y
															man
															ual
															or
															phys
															ical
															wor
															k
															Job
															invo
															lves
															heavy
1	rs45	BMI	A	G	A	G	-0.022	-0.033	0.004	602604	2.29E-	Gastroesophag	3.00E-	0.002	Job
6	8087						0715	8177	8202	6	12	al reflux	-20	263615	invo
	6											disease		0.3945	lves
															man
															ual
															or
															phys
															ical
															wor
															k
															Job
															invo
1	rs47														lves
7	3199	G	A	G	A		-0.017	-0.022	0.005	602604	0.0001	Gastroesophag	4.00E-	0.002	heav
	2						0407	2119	8293	4	38759	al reflux	-09	263615	y
												disease		0.895	man
															ual

															or
															phys
															ical
															wor
															k
															Job
															invo
															lves
															heav
1	rs62														y
8	5118	A	G	A	G	-0.021	-0.014	0.006							man
	03					1702	5097	8906	602604	0.0352	29	Gastroesophag	7.50E-	0.003	ual
								4				al reflux	-10	4396	or
												disease		8	phys
															ical
															wor
															k
															Job
															invo
															lves
															heav
1	rs65														y
9	4476	C	T	C	T	0.0139	0.0182	0.005				Gastroesophag	3.20E-	0.002	man
	3					692	768	072	602604	0.0003	14007	al reflux	-08	5258	ual
												disease		6	or
															phys
															ical
															wor
															k

														Job invo lves heav	
2	rs71 0807 7	G	A	G	A	-0.013 4395	-0.011 7868	0.004 9615 8	602604	0.0175 194	Gastroesophag al reflux disease	4.90E- 08	263615	0.002 4634 5	y man ual or phys ical wor k Job invo lves heav
2	rs74 6748 0	A	T	A	T	-0.015 5085	-0.016 3904	0.004 8854 4	602604	0.0007 93816	Gastroesophag al reflux disease	1.70E- 10	263615	0.002 4292	y man ual or phys ical wor k Job invo lves heav
2	rs80 5411 1	G	A	G	A	-0.015 349	-0.011 7556	0.005 4579	602604	0.0312 507	Gastroesophag al reflux disease	1.20E- 08	263615	0.002 6932	lves heav y man ual

or
phys
ical
wor
k

- **Supplementary Table 3** Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Average total household income before tax and risk of Gastroesophageal reflux disease.

	SNP	Related confounding phenotypes	effect_size.exposure	other_size.exposure	effect_size.outcome	other_size.outcome	beta_exposure	beta_outcome	samplesize.outcome	pval.outcome	outcome	outcome.deplicated	pval.exposure	samplesize.exposure	se.exposure	expouse
1	rs10761035	A G A G	0.0186 402	-0.022 6412	602604	0.000 24103 5	Gastroesophageal reflux disease	Gastroesophageal reflux disease	3.70E-08	397751	0.003 3851 2	Average total household income before tax	3.70E-08	397751	0.003 3851 2	seh old inco me befo re tax

														Ave		
														rage		
														total		
2	rs11 588 857	Alcoh ol	A	G	A	G	0.0213 402	-0.012 7023	602604	0.031 5435	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	4.90E- 11	397751	0.003 2461 5	seh old inco me befo re tax Ave rage total hou
3	rs11 665 242	Alcoh ol	G	A	G	A	-0.017 5478	0.031 7599	602604	7.73E -11	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	6.00E- 11	397751	0.002 6816 6	seh old inco me befo re tax Ave rage total hou
4	rs12 692 596	BMI	T	C	T	C	-0.015 4093	0.019 8715	602604	6.52E -05	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	1.70E- 08	397751	0.002 7334 7	hou seh old inco me befo

															re tax	
															Ave rage total hou	
															seh old inco me befo re tax	
5	rs13 002 946	BMI	A	T	A	T	0.0205 047	-0.033 6393	602604	6.50E- -10	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	7.30E- 12	397751	0.002 9928 5	Ave rage total hou seh old inco me befo re tax
6	rs14 213 34	BMI	C	A	C	A	0.0162 948	-0.020 2427	602604	3.12E- -05	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	1.00E- 09	397751	0.002 6702 8	Ave rage total hou seh old inco me befo re tax

														Ave	
														rage	
														total	
rs14														hou	
7	553	A	T	A	T	-0.017 2341	0.015 1799	602604	0.001 6761	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	9.20E- 11	397751	0.002 6596 1	seh old inco me befo re tax Ave rage total hou
rs20														hou	
8	684	T	C	T	C	0.0169 724	-0.009 41417	602604	0.094 1369	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	4.20E- 08	397751	0.003 0951 5	seh old inco me befo re tax Ave rage total hou
rs23														hou	
9	327	G	A	G	A	-0.018 3042	0.022 2214	602604	3.58E -05	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	5.70E- 10	397751	0.002 9532 3	hou seh old inco me befo

1	rs24 228 0	G	T	G	T	0.0163 711	-0.009 56311	602604	0.046 9732	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	6.20E- 10	397751	0.002 6471 8	seh old inco me befo re tax Ave rage total hou
1	rs38 778 1	C	T	C	T	0.0167 299	-0.015 0838	602604	0.003 35714	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	3.30E- 09	397751	0.002 8273 2	seh old inco me befo re tax Ave rage total hou

													Ave	
													rage	
													total	
1	rs57												hou	
2	547	G	A	G	A	-0.016	0.011	602604	0.026	Gastroesoph	Gastroesophag	2.70E-	0.002	
3	38					2464	8154		3937	ageal reflux	eal reflux	08	397751	seh
										disease	disease		9203	old
													8	inco
														me
														befo
														re
														tax
														Ave
														rage
														total
1	rs60												hou	
2	358	C	A	C	A	-0.014	0.004	602604	0.345	Gastroesoph	Gastroesophag	4.30E-	0.002	
3	77					5138	55135		124	ageal reflux	eal reflux	08	397751	seh
										disease	disease		6504	old
													2	inco
														me
														befo
														re
														tax
														Ave
														rage
														total
1	rs66												hou	
2	993	BMI	G	A	G	-0.019	0.020	602604	5.77E	Gastroesoph	Gastroesophag	3.50E-	0.002	
3	97					0757	122		-05	ageal reflux	eal reflux	12	397751	seh
										disease	disease		7415	old
													3	inco
														me
														befo

1	rs68457	C	T	C	T	0.0209 797	-0.008 64939	602604	0.073 0685	Gastroesophageal reflux disease	Gastroesophageal reflux disease	3.10E-15	397751	0.002 6601	Average total houseold income before tax
5	rs77001 07	BMI	C	A	C	A	-0.022 8296	0.035 5854	602604	3.81E-07	Gastroesophageal reflux disease	Gastroesophageal reflux disease	2.50E-09	397751 8299 2	Average total houseold income before tax
6															

														Ave		
														rage		
														total		
1	rs78													hou		
7	425	BMI	A	G	A	G	-0.025 3813	0.028 9491	602604	2.70E -06	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	7.90E- 14	397751	0.003 3968	seh old inco me befo re tax Ave rage total hou
6																
1	rs93													hou		
8	884	T	C	T	C		0.0151 05	-0.023 6637	602604	1.06E -06	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	1.50E- 08	397751	0.002 6686	seh old inco me befo re tax Ave rage total hou
90																
1	rs95													hou		
9	569	BMI	T	C	T	C	-0.015 3409	0.027 291	602604	1.60E -08	Gastroesoph ageal reflux disease	Gastroesophag eal reflux disease	8.50E- 09	397751	0.002 6641	seh old inco me befo
58																

- **Supplementary Table 4** Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Townsend deprivation index at recruitment and risk of Gastroesophageal reflux disease.

SNP	Related confounding phenotypes														Tow nsen d depr ivati on inde x at recr uitm ent
	effect_all_ele.exposure	other_all_ele.exposure	effect_all_ele.outcome	other_all_ele.outcome	beta.e_xposure	beta.outcome	se.outcome	samplesize.outcome	pval.outcome	outcome	pval.e_xposure	se.exposure	samplesize.exposure	exp	
rs113 1 3452 85	BMI C T C T	0.0163 127	0.0200 155	0.007 1161 4	602604	0.0049 1281	Gastroesophageal reflux disease	3.20E-08	0.002 9509 4	462464					

															Tow nse d depr ivati on inde x at recr uitm ent Tow nse d depr ivati on inde x at recr uitm ent Tow nse d depr ivati on inde x at recr uitm ent Tow nse d depr ivati on inde x at recr
2	rs118 5582 1	BMI	A	G	A	G	-0.012 5981	-0.025 9592	0.005 4394 7	602604	1.82E- 06	Gastroesophag al reflux disease	3.90E- 08	0.002 2925 9	462464
3	rs148 3246	C	T	C	T		-0.012 1102	-0.009 3388	0.004 8450 9	602604	0.0539 2	Gastroesophag al reflux disease	3.10E- 09	0.002 0440 9	462464
4	rs194 7083	A	G	A	G		0.0118 662	0.0160 533	0.005 0061 9	602604	0.0013 4273	Gastroesophag al reflux disease	1.90E- 08	0.002 1105	462464

															uitm ent
5	rs240 3326	BMI	G	A	G	A	0.0115 465	0.0191 822	0.004 8119 4	602604	6.71E- 05	Gastroesophag al reflux disease	1.20E- 08	0.002 0270 3	462464
6	rs253 125	C	T	C	T	T	0.0131 74	-0.005 70351	0.005 5683 2	602604	0.3057 03	Gastroesophag al reflux disease	1.90E- 08	0.002 3432	462464

7	rs386 5018	BMI	T	C	T	C	-0.013 4594	-0.017 8233	0.005 4528	602604	0.0010 8059	Gastroesophageal reflux disease	4.20E-09	0.002 2908 6	462464
8	rs478 5187		A	G	A	G	0.0137 996	0.0192 606	0.005 7920 2	602604	0.0008 83019	Gastroesophageal reflux disease	1.50E-08	0.002 4370 3	462464
9	rs693 1604	Alcohol	T	C	T	C	-0.012 1903	-0.020 4973	0.005 0225 9	602604	4.48E-05	Gastroesophageal reflux disease	4.10E-09	0.002 0730 9	462464

1	rs704 067	BMI	A	G	A	G	0.0120 267	0.0226 494	0.004 8543 8	602604	3.07E- 06	Gastroesophag al reflux disease	3.90E- 09	0.002 0424	462464
1	rs774 10440		A	G	A	G	0.0118 47	0.0202 381	0.004 8211 8	602604	2.70E- 05	Gastroesophag al reflux disease	5.70E- 09	0.002 0339 1	462464

1	rs782 5712 8	T	C	T	C	-0.016 7416	-0.003 64373	0.006 3	6265 3	602604	0.5824 09	Gastroesophageal reflux disease	2.40E-09 09	0.002 8067 6	462464
1	rs989 532	G	A	G	A	0.0138 559	0.0113 503	0.005 1313 6	602604	0.0269 699	Gastroesophageal reflux disease	7.40E-11 11	0.002 1275 6	462464	
1	rs990 706	T	C	T	C	0.0155 869	-0.012 9805	0.006 4345 7	602604	0.0436 626	Gastroesophageal reflux disease	7.90E-09 09	0.002 7014 1	462464	

Supplementary Table 5 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Leisure/social activities: Pub or social club and risk of Gastroesophageal reflux disease

Related confounding phenotypes															Leisure/social activities:
SNP	effect_all_ele.exposure	other_all_ele.exposure	effect_all_ele.outcome	other_all_ele.outcome	beta.e_xposure	beta.outcome	se.outcome	samplesize.outcome	pval.outcome	outcome	pval.exposure	se.exposure	samplesize.exposure	exposure	
rs11066099	G	C	G	C	0.0065 1032	-0.014 7389	0.006 3309 5	602604	0.0199 081	Gastroesophageal reflux disease	4.60E-08	0.001 1909 2	461369	Pub or social club	

2	rs13 9920	BMI	T	C	T	C	0.0061 4932	0.0071 2136	0.004 8137 9	602604	0.1390 42	Gastroesophag al reflux disease	1.20E- 11	0.000 9075 3	461369	Leisu re/soc ial activi ties: Pub or social club Leisu re/soc ial activi ties:
3	rs26 6186 3	C	T	C	T	-0.005 37368	0.0056 056	0.005 0562 5	602604	0.2675 83	Gastroesophag al reflux disease	1.90E- 08	0.000 9567 22	461369	Leisu re/soc ial activi ties: Pub or social club Leisu re/soc ial activi ties:	
4	rs75 4204	T	C	T	C	0.0050 7247	0.0004 13709	0.004 8349 9	602604	0.9318 12	Gastroesophag al reflux disease	3.00E- 08	0.000 9152 01	461369	Leisu re/soc ial activi ties: Pub or social club	

5	rs75 4516 1	T	C	T	C	-0.006 87787	-0.006 04065	0.006 2211 7	602604	0.3315 57	Gastroesophag eal reflux disease	4.20E- 09	0.001 1701 8	461369	Leisu re/soc ial activi ties: Pub or social club	
6	rs78 0094	Alcoh ol	C	T	C	T	0.0051 697	-0.003 84079	0.004 9459	602604	0.4374 18	Gastroesophag eal reflux disease	2.70E- 08	0.000 9301 37	461369	Leisu re/soc ial activi ties: Pub or social club
7	rs94 0159 3	BMI	C	A	C	A	-0.005 31642	-0.034 7468	0.004 8223 2	602604	5.79E- 13	Gastroesophag eal reflux disease	5.30E- 09	0.000 9107 46	461369	Leisu re/soc ial activi ties: Pub or social club

Supplementary Table 6 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Leisure/social activities: Sports club or gym and risk of Gastroesophageal reflux disease.

		Related confounding phenotypic traits															
SNP		effect_all_ele.exposure	other_all_ele.exposure	effect_allele.outcome	other_allele.outcome	beta.e xposure	beta.outcome	se.outcome	samplesize.outcome	pval.outcome	outcome	se.exposure	samplesize.exposure	pval.exposure	exposure		
rs11712056	BMI	C	T	C	T	-0.00597808	0.0424557	0.00483548	602604	1.63E-18	Gastroesophageal reflux disease	0.00096055	461369	4.90E-10	Leisure/social activities: Sports club or gym	Leisure/social activities: Sports club or gym	
rs13002862		A	C	A	C	0.0060452	-0.00117087	0.00550497	602604	0.831566	Gastroesophageal reflux disease	0.00109537	461369	3.40E-08	Leisure/social activities: Sports club or gym	Leisure/social activities: Sports club or gym	

rs17 3	3150 37	C	T	C	T	-0.007 88121	0.0045 8221	0.006 9191 6	602604	0.5078 11	Gastroesophag al reflux disease	0.001 3926 2	461369	1.50E-08	Leisu re/soc ial activi ties: Sport s club or gym Leisu re/soc ial	
rs62 4	2639 12	Smoki ng	A	G	A	G	-0.006 11858	-0.003 39804	0.005 0045 2	602604	0.4971 41	Gastroesophag al reflux disease	0.000 9922 44	461369	7.00E-10	activi ties: Sport s club or gym Leisu re/soc ial
rs64 5	7844 4	G	A	G	A	0.0055 0414	-0.001 21552	0.005 0577 5	602604	0.8100 77	Gastroesophag al reflux disease	0.001 0037 3	461369	4.20E-08	activi ties: Sport s club or gym	

- Supplementary Table 7 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Age completed full time education and risk of Barrett's esophagus.

1	rs10 189 857	BMI	G	A	G	A	-0.012 4054	0.031 531	0.015 1383	56429	0.037 264	Barrett's esophagus id:ebi-a-GCST900 00515	307897	2.70E-09	0.002 0847 4
2	rs10 760 199	BMI	T	G	T	G	0.0115 963	0.003 42634	0.015 179	56429	0.821 413	Barrett's esophagus id:ebi-a-GCST900 00515	307897	2.50E-08	0.002 0803 3
3	rs11 665 242	Alcoh ol	G	A	G	A	-0.012 5523	0.047 4094	0.015 2082	56429	0.001 82478	Barrett's esophagus id:ebi-a-GCST900 00515	307897	2.10E-09	0.002 0962 2

rs14 4 621 63	T	C	T	C	-0.012 7222	0.013 7455	0.015 4459	56429	0.373 513	 id:ebi-a-GCST900 00515	Barrett's esophagus	307897	1.90E-09	0.002 1196 9
rs17 5 563 464	A	C	A	C	-0.014 8391	0.015 247	0.018 8083	56429	0.417 565	 id:ebi-a-GCST900 00515	Barrett's esophagus	307897	3.50E-09	0.002 5124 8
rs17 6 821 7	C	T	C	T	-0.014 3669	0.052 7652	0.017 3023	56429	0.002 29145	 id:ebi-a-GCST900 00515	Barrett's esophagus	307897	2.10E-09	0.002 3974 6

7	rs25 889 62	Smok ing	A	G	A	G	0.0150 825	-0.037 6218	0.015 1151	56429	0.012 8097	Barrett's esophagus id:ebi-a-GCST900 00515	307897	4.10E- 13	0.002 0797 2	Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on
8	rs45 577 20		G	A	G	A	0.0119 687	-0.028 0723	0.015 5009	56429	0.070 1391	Barrett's esophagus id:ebi-a-GCST900 00515	307897	2.00E- 08	0.002 1339 3	Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on
9	rs46 744 03		G	T	G	T	-0.012 7249	0.043 9594	0.017 1779	56429	0.010 4954	Barrett's esophagus id:ebi-a-GCST900 00515	307897	4.90E- 08	0.002 3324	Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on Age com plet ed full time edu cati on

1	rs47 319 51	T	G	T	G	0.0153 215	-0.050 3112	0.018 7821	56429	0.007 39129	 id:ebi-a-GCST900 00515	Barrett's esophagus	3.40E-09	0.002 5923 6
1	rs65 083 44	A	G	A	G	-0.011 5957	0.019 7887	0.015 2033	56429	0.193 05	 id:ebi-a-GCST900 00515	Barrett's esophagus	2.80E-08	0.002 0872 4
1	rs66 793 99	C	A	C	A	0.0426 608	-0.045 4615	0.049 9448	56429	0.362 698	 id:ebi-a-GCST900 00515	Barrett's esophagus	3.00E-10	0.006 7701 6

1	rs67												Age		
3	295	A	G	A	G	-0.011 7953	0.010 3537	0.015 3002	56429	0.498 595	Barrett's esophagus id:ebi-a-GCST900 00515	307897	2.40E-08	0.002 1134 1	com plet ed full time edu cati on Age
1	rs71												com plet ed full time edu cati on Age		
4	107	T	C	T	C	0.0127 141	-0.047 7787	0.015 2041	56429	0.001 67517	Barrett's esophagus id:ebi-a-GCST900 00515	307897	1.20E-09	0.002 0912 1	com plet ed full time edu cati on Age
1	rs77												com plet ed full time edu cati on Age		
5	687	C	T	C	T	-0.022 7349	0.049 2636	0.027 8694	56429	0.077 117	Barrett's esophagus id:ebi-a-GCST900 00515	307897	2.10E-09	0.003 7937 1	com plet ed full time edu cati on

1	rs96 557 80	Alcoh ol	A	G	A	G	-0.017 4579	0.021 526	0.020 1497	56429	0.285 386	Barrett's esophagus id:ebi-a-GCST900 00515	307897	4.30E- 10	0.002 7964 6	Age com plet ed full time edu cati on Age com plet ed full time edu cati on
1	rs98 666 30		A	G	A	G	-0.012 6597	0.024 975	0.014 9754	56429	0.095 3674	Barrett's esophagus id:ebi-a-GCST900 00515	307897	1.00E- 09	0.002 0719	Age com plet ed full time edu cati on

- Supplementary Table 8 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Job involves heavy manual or physical work and risk of Barrett's esophagus.

SNP	Related confounding phenotypes														
	effect_all_ele.exposure	other_all_ele.exposure	effect_all_ele.outcome	other_all_ele.outcome	beta.e_xposure	beta.o_utcome	se.ou_tcome	samplesize_outcome	pval.outcome	outcome	pval_exposure	samplesize_exposure	se.exposure	exp_outcome	

															Job	
1	rs10 8115 8	Alcoh ol	T	C	T	C	0.0169 887	0.0038 6911	0.015 2839	56429	0.8001 52	Barrett's esophag us	2.70E- 12	263615	0.002 42996	y man ual or phys ical wor k
2	rs10 8206 25		C	T	C	T	-0.0160 068	-0.066 3494	0.018 092	56429	0.0002 45092	Barrett's esophag us	2.90E- 08	263615	0.002 88515	y man ual or phys ical wor k
3	rs11 1308 89		G	C	G	C	-0.0249 218	-0.026 1838	0.028 6138	56429	0.3601 52	Barrett's esophag us	4.50E- 08	263615	0.004 55725	lves heav y man ual

															or	
															phys	
															ical	
															wor	
															k	
															Job	
															invo	
															lves	
															heavy	
															y	
4	rs11 6638 24	Alcoh ol	A	C	A	C	0.0136 375	0.0335 832	0.015 306	56429	0.0282 273	Barrett's esophag us	2.50E- 08	263615	0.002 44806	man ual or phys ical wor k
															Job	
															invo	
															lves	
															heavy	
															y	
5	rs11 6789 79	BMI	C	T	C	T	-0.0149 934	-0.053 1341	0.017 2002	56429	0.0020 0729	Barrett's esophag us	3.00E- 08	263615	0.002 70637	man ual or phys ical wor k

															Job	
															invo	
															lves	
															heavy	
6	rs11 7267 86	Alcoh ol	G	T	G	T	0.0165 951	0.0153 067	0.015 5479	56429	0.3248 76	Barrett's esophag us	2.40E- 11	263615	0.002 48344	y man ual or phys ical wor k
7	rs11 7561 23		T	A	T	A	-0.0137 486	-0.010 4674	0.015 7484	56429	0.5062 66	Barrett's esophag us	3.20E- 08	263615	0.002 48658	y man ual or phys ical wor k
8	rs13 2509 96		C	G	C	G	-0.0133 122	0.0250 578	0.015 1571	56429	0.0982 902	Barrett's esophag us	2.70E- 08	263615	0.002 39307	lves heavy y man ual

or
phys-
ical
wor-
k

Job
invo
lves
heav

rs13
9 7005 BMI G A G A 0.0180 0.0486 0.017 56429 0.0062 Barrett's
9 9 816 319 7771 2587 esophag 1.80E-
us 10 263615 0.002 man
83492 ual

rs20
9132 G A G A -0.0158 -0.043 0.016 56429 0.0081 Barrett's
9 9 895 619 4801 2643 esophag
us 2.00E- 263615 0.002 man
09 65008 ual

															Job	
1	rs23 1854 0	BMI	G	T	G	T	-0.0142 827	-0.034 8159	0.015 2291	56429	0.0222 459	Barrett's esophag us	3.70E- 09	263615	0.002 42247	man ual or phys ical wor k Job invo lves heav
1	rs28 1934 8	BMI	C	T	C	T	0.0145 582	0.0322 053	0.015 7062	56429	0.0403 172	Barrett's esophag us	5.60E- 09	263615	0.002 49731	man ual or phys ical wor k Job invo lves heav
1	rs34 7007 31		A	G	A	G	0.0197 694	-0.015 9721	0.022 0407	56429	0.4686 58	Barrett's esophag us	1.50E- 08	263615	0.003 49445	lves heav y man ual

																	or
																	phys
																	ical
																	wor
																	k
																	Job
																	invo
																	lves
																	heav
1	rs35																y
4	9991	BMI	G	C	G	C	-0.0248	-0.049	0.016	56429	0.0024	Barrett's	7.40E-	263615	0.002	man	
	62						598	4108	34		9523	esophag	-22		5873	ual	
												us				or	
																phys	
																ical	
																wor	
																k	
																Job	
																invo	
																lves	
																heav	
1	rs37															y	
5	8535	BMI	T	C	T	C	0.0135	0.0135	0.015	56429	0.3823	Barrett's	3.70E-	263615	0.002	man	
	4						824	052	4584		12	esophag	-08		4672	ual	
												us				or	
																phys	
																ical	
																wor	
																k	

														Job		
														invo		
														lves		
1	rs45													heavy		
6	8087	BMI	A	G	A	G	-0.0220 715	-0.066 0802	0.015 1042	56429	1.21E- 05	Barrett's esophag us	3.00E- 20	263615	0.002 3945	man ual or phys ical wor k
1	rs47													Job		
7	3199		G	A	G	A	-0.0170 407	-0.039 451	0.018 1492	56429	0.0297 276	Barrett's esophag us	4.00E- 09	263615	0.002 895	invo lves heavy
1	rs62													y		
8	5118		A	G	A	G	-0.0211 702	-0.011 9118	0.021 5452	56429	0.5803 49	Barrett's esophag us	7.50E- 10	263615	0.003 43968	man ual or phys ical wor k
1	rs62													Job		
8	03													invo		
1	rs62													lves		
8	03													heavy		
1	rs62													y		
8	03													man ual		

1 9	rs65 4476 3	C	T	C	T	0.0139 692	0.0292 379	0.015 8076	56429	0.0643 695	Barrett's esophag us	3.20E- 08	263615	0.002 52586	Job invo lves heav y man ual or phys ical wor k
2 0	rs71 0807 7	G	A	G	A	-0.0134 395	-0.014 1655	0.015 4855	56429	0.3603 16	Barrett's esophag us	4.90E- 08	263615	0.002 46345	Job invo lves heav y man ual or phys ical wor k

														Job	
2	rs74													invo	
1	6748	A	T	A	T	-0.0155 085	-0.016 5796	0.015 3925	56429	0.2814 26	Barrett's esophag us	1.70E- 10	263615	0.002 4292	lves
2	rs80													heavy	
2	5411	G	A	G	A	-0.0153 49	-0.027 8481	0.017 0007	56429	0.1014 09	Barrett's esophag us	1.20E- 08	263615	0.002 6932	y
1														man	
														ual	
														or	
														physical	
														wor	
														k	
														Job	
														invo	
														lves	
														heavy	
														y	
														man	
														ual	
														or	
														physical	
														wor	
														k	

- Supplementary Table 9 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Average total household income before tax and risk of Barrett's esophagus.

	SNP	Related confounding phenotypes	effect_all_ele.exposure	other_all_ele.exposure	effect_all_ele.outcome	other_all_ele.outcome	beta.e_xposure	beta.outcome	se.o_utco	samplesize_outcome	pval.outcome	pvalue_xposure	samplesize_exposure	se.exposure	expposure
1	rs10429582	C T C T	0.0270 261	-0.016 5796	0.015 3925	56429	0.2814 26	Barrett's esophagus	7.60E-24	397751	0.002 68405	Average seh old inc om e bef ore tax Ave rag e	total 1 hou seh old inc om e bef ore tax Ave rag e		
2	rs10761035	A G A G	0.0186 402	-0.076 484	0.019 207	56429	6.83E-05	Barrett's esophagus	3.70E-08	397751	0.003 38512	total 1 hou seh old inc	total 1 hou seh old inc		

3	rs111 9111 6	T	C	T	C	-0.016 3993	0.0229 312	0.015 6668	56429	0.1432 8	Barrett's esophagus	3.50E- 09	397751	0.002 77643	
4	rs115 8885 7	Alcoh ol	A	G	A	G	0.0213 402	0.0072 9364	0.018 2288	56429	0.6890 69	Barrett's esophagus	4.90E- 11	397751	0.003 24615

															ore	
															tax	
															Ave	
															rag	
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															tota	
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															hou	
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															old	
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															om	
															e	
															bef	
															ore	
															tax	
															Ave	
															rag	
															e	
															tota	
															l	
															hou	
															seh	
															old	
															inc	
															om	
															e	
															bef	
5	rs116 6524 2	Alcoh ol	G	A	G	A	-0.017 5478	0.0474 094	0.015 2082	56429	0.0018 2478	Barrett's esophagus	6.00E- 11	397751	0.002 68166	Ave rag e tota l hou seh old inc om e bef ore tax
6	rs116 7850 1	C	T	C	T	-0.051 7926	0.0033 8915	0.048 2575	56429	0.9440 1	Barrett's esophagus	1.20E- 09	397751	0.008 51645	Ave rag e tota l hou seh old inc om e bef	

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															Ave	
															rag	
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															bef	
															ore	
															tax	
															Ave	
															rag	
															e	
															tota	
															l	
7	rs117	1433	Alcoh	A	G	A	G	0.0153	-0.038	0.015	56429	0.0133	Barrett's	9.40E-	397751	0.002
		7	ol					807	4626	5466		607	esophagus	09		67881
8	rs118	7775	Alcoh	G	T	G	T	-0.020	0.0533	0.015	56429	0.0007	Barrett's	1.20E-	397751	0.002
		8	ol					3602	468	8799		81124	esophagus	12		86697

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															tax			
															Ave			
															rag			
															e			
															tota			
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															e			
															bef			
															ore			
															tax			
															Ave			
															rag			
															e			
															tota			
															l			
9	rs119 1	1743	BMI	T	C	T	C	0.0227 135	-0.052 5662	0.016 5358	56429	0.0014 7816	Barrett's esophagus	3.10E- 15	397751	0.002 88021		
1	rs123 0	9705		G	A	G	A	0.0155 639	-0.036 3754	0.015 1036	56429	0.0160 232	Barrett's esophagus	4.70E- 09	397751	0.002 65785	hou seh old inc om e bef	

1	rs126 9259 6	BMI	T	C	T	C	-0.015 4093	0.0476 95	0.015 5212	56429	0.0021 1997	Barrett's esophagus	1.70E- 08	397751	0.002 73347	Ave rag e tota l hou seh old inc om e bef ore tax Ave rag e tota l hou seh old inc om e bef
1	rs128 8378 8	BMI	T	C	T	C	-0.018 8574	-0.003 70028	0.014 9768	56429	0.8048 56	Barrett's esophagus	1.40E- 12	397751	0.002 66308	Ave rag e tota l hou seh old inc om e bef

1	rs130 0294 3	BMI	A	T	A	T	0.0205 047	-0.053 1401	0.017 2051	56429	0.0020 109	Barrett's esophagus	7.30E- 12	397751	0.002 99285	ore tax	Ave rag e tota l hou seh old inc om e bef ore tax	
1	rs142 1334 4	BMI	C	A	C	A	0.0162 948	-0.021 9632	0.015 251	56429	0.1498 35	Barrett's esophagus	1.00E- 09	397751	0.002 67028	hou seh old inc om e bef	Ave rag e tota l hou seh old inc om e bef	

															ore
															tax
1	rs145	A	T	A	T	-0.017 2341	0.0171 412	0.015 168	56429	0.2584 4	Barrett's esophagus	9.20E- 11	397751	0.002 65961	seh old inc om e bef ore tax Ave rag e tota l hou
5	5350														
1	rs206	T	C	T	C	0.0169 724	0.0051 8713	0.017 6429	56429	0.7687 52	Barrett's esophagus	4.20E- 08	397751	0.003 09515	hou seh old inc om e bef
6	8428														

1	rs233															ore
7	2719	G	A	G	A	-0.018 3042	0.0177 633	0.016 7613	56429	0.2892 43	Barrett's esophagus	5.70E- 10	397751	0.002 95323	seh old inc om e bef ore tax Ave rag e tota l hou	tax
1	rs236															Ave
8	2523	G	A	G	A	0.0158 667	0.0095 996	0.016 255	56429	0.5548 13	Barrett's esophagus	1.40E- 08	397751	0.002 79836	hou seh old inc om e bef	rag

1	rs242																		
9	2859	G	T	G	T	0.0163 711	-0.006 30733	0.014 9776	56429	0.6736 68	Barrett's esophagus	6.20E- 10	397751	0.002 64718	seh old inc om e bef ore tax Ave rag e tota l hou				
2	rs282	BMI	C	A	C	A	-0.016 5601	0.0396 095	0.016 1309	56429	0.0140 689	Barrett's esophagus	2.90E- 09	397751	0.002 78855	hou seh old inc om e bef			

															ore tax
															Ave rag e tota l hou seh old inc om e bef ore tax Ave rag e tota l hou seh old inc om e bef
2	rs387		C	T	C	T	0.0167 299	-0.028 0818	0.016 1118	56429	0.0813 448	Barrett's esophagus	3.30E- 09	397751	0.002 82732
1	780														
2	rs411	BMI	A	G	A	G	-0.017 9965	0.0103 173	0.015 6503	56429	0.5097 44	Barrett's esophagus	1.10E- 10	397751	0.002 79169
2	5668														

														ore	
														tax	
														Ave	
														rag	
														e	
														tota	
														l	
														hou	
2	rs488	T	C	T	C	0.0206 878	0.0290 769	0.020 9333	56429	0.1648 25	Barrett's esophagus	6.70E- 09	397751	0.003 56762	seh old
3	786													inc	
														om	
														e	
														bef	
														ore	
														tax	
														Ave	
														rag	
														e	
														tota	
														l	
2	rs575	G	A	G	A	-0.016 2464	-0.014 3007	0.016 6343	56429	0.3899 47	Barrett's esophagus	2.70E- 08	397751	0.002 92038	hou seh
4	4738													old	
														inc	
														om	
														e	
														bef	

															ore
															tax
															Ave
															rag
															e
															tota
															l
															hou
2	rs589	Alcoh	G	A	G	A	0.0156	-0.025	0.016	56429	0.1111	Barrett's	4.40E-	397751	0.002
5	914	ol					098	894	2557		78	esophagus	08		85076
															seh
															old
															inc
															om
															e
															bef
															ore
															tax
															Ave
															rag
															e
															tota
															l
2	rs603		C	A	C	A	-0.014	0.0251	0.015	56429	0.0945	Barrett's	4.30E-	397751	0.002
6	5877						5138	541	047		823	esophagus	08		65042
															hou
															seh
															old
															inc
															om
															e
															bef

															ore tax	
															Ave rag e tota l hou inc om e bef ore tax Ave rag e tota l hou seh old inc om e bef ore tax Ave rag e tota l hou seh old inc om e bef	
2	rs621															
7	8302	T	G	T	G	-0.018 9832	0.0346 632	0.016 4578	56429	0.0351 876	Barrett's esophagus	3.20E- 11	397751	0.002 86114	seh old inc om e bef ore tax Ave rag e tota l hou seh old inc om e bef	
2	rs642	Smoki	T	G	T	G	0.0192 858	-0.030 9935	0.016 9206	56429	0.0669 962	Barrett's esophagus	5.00E- 11	397751	0.002 93518	hou seh old inc om e bef
8	9636	ng														

2 9	rs669 9397	BMI	G	A	G	A	-0.019 0757	0.0214 816	0.015 4816	56429	0.1652 72	Barrett's esophagus	3.50E- 12	397751	0.002 74153	seh old inc om e bef ore tax Ave rag e tota l hou	ore tax	
3 0	rs686 8457	C	T	C	T		0.0209 797	0.0089 3787	0.014 9911	56429	0.5510 35	Barrett's esophagus	3.10E- 15	397751	0.002 6601	hou seh old inc om e bef	Ave rag e tota l hou	

														ore tax
														Ave rag e tota l hou inc om e bef ore tax Ave rag e tota l hou old inc om e bef
3	rs730													
1	1532	T	G	T	G	-0.027	0.0620	0.028	56429	0.0317	Barrett's	4.10E-	397751	0.004
2	2					1853	315	879		154	esophagus	08		95368
3	rs754					-0.026	0.0501	0.022	56429	0.0290	Barrett's	9.50E-	397751	0.004
2	1332	C	T	C	T	1129	929	9923		335	esophagus	10		26829
0	0													

															ore	
															tax	
3	rs770 3 0107	BMI	C	A	C	A	-0.022 8296	0.0929 077	0.021 843	56429	2.11E- 05	Barrett's esophagus	2.50E- 09	397751	0.003 82992	seh old inc om e bef ore tax
3	rs784 4 256	BMI	A	G	A	G	-0.025 3813	0.0089 3944	0.019 4429	56429	0.6456 75	Barrett's esophagus	7.90E- 14	397751	0.003 39684	hou seh old inc om e bef
															Ave rag e tota l hou	

3	rs789 56518	BMI	G	A	G	A	0.0148 13	-0.045 1136	0.015 04	56429	0.0027 0352	Barrett's esophagus	4.10E- 08	397751	0.002 70047	seh old inc om e bef ore tax Ave rag e tota l hou		
3	rs938 68490	Nervo us	T	C	T	C	0.0151 05	-0.021 958	0.015 1386	56429	0.1469 3	Barrett's esophagus	1.50E- 08	397751	0.002 66861	hou seh old inc om e bef		

3 7	rs955 6958	BMI	T	C	T	C	-0.015 3409	0.0294 665	0.015 2257	56429	0.0529 529	Barrett's esophagus	8.50E- 09	397751	0.002 6641	seh old inc om e bef ore tax	Ave rag e tota l hou		
3 8	rs968 050	Alcoh ol	T	C	T	C	0.0223 829	-0.046 5845	0.014 979	56429	0.0018 7098	Barrett's esophagus	3.10E- 17	397751	0.002 65073	hou seh old inc om e bef	Ave rag e tota l hou		

3 9	rs989 1103	Alcoh ol	T	C	T	C	-0.023 4517	-0.023 6481	0.018 0562	56429	0.1903	Barrett's esophagus	9.40E- 14	397751	0.003 14818
															Ave rag e tota l hou seh old inc om e bef ore tax

- Supplementary Table 10 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Townsend deprivation index at recruitment and risk of Barrett's esophagus.

SNP	Related phenotypic confounding variables														
	confounder	effect_all_ele.exposure	other_all_ele.exposure	effect_all_ele.outcome	other_all_ele.outcome	beta_exposure	beta_outcome	se_outcome	sample_size_outcome	pval_outcome	outcome	pval_exposure	se_exposure	sample_size_exposure	exposure
	undifferentiated phenotype	exposure	exposure	outcome	outcome	exposure	outcome	exposure	size_outcome	outcome	exposure	exposure	exposure	size_exposure	exposure

															Tow nsen d depri vatio n inde x at recru itme nt	
1	rs113 3452 85	BMI	C	T	C	T	0.0163 127	0.0008 26945	0.022 6093	56429	0.9708 24	Barrett's esophag us	3.20E- 08	0.002 95094	462464	
2	rs118 5582 1	BMI	A	G	A	G	-0.012 5981	-0.034 3493	0.017 2571	56429	0.0465 415	Barrett's esophag us	3.90E- 08	0.002 29259	462464	

3	rs148 3246	C	T	C	T	-0.012 1102	-0.004 56071	0.015 1265	56429	0.7630 29	Barrett's esophag us	3.10E- 09	0.002 04409	462464	Tow nsen d depri vatio n inde x at recru itme nt	
4	rs194 7083	A	G	A	G	0.0118 662	0.0116 892	0.015 62	56429	0.4542 48	Barrett's esophag us	1.90E- 08	0.002 1105	462464	Tow nsen d depri vatio n inde x at recru itme nt	
5	rs240 3326	BMI	G	A	G	A	0.0115 465	0.0163 561	0.015 033	56429	0.2765 88	Barrett's esophag us	1.20E- 08	0.002 02703	462464	Tow nsen d depri vatio n inde x at recru

													itme		
													nt		
6	rs253 125	C	T	C	T	0.0131 74	-0.003 81172	0.017 4319	56429	0.8269 12	Barrett's esophag us	1.90E- 08	0.002 3432	462464	
7	rs386 5018	BMI	T	C	T	C	-0.013 4594	-0.038 7711	0.017 1256	56429	0.0235 787	Barrett's esophag us	4.20E- 09	0.002 29086	462464

8	rs478 5187	A	G	A	G	0.0137 996	0.0287 684	0.018 2185	56429	0.1143 18	Barrett's esophag us	1.50E- 08	0.002 43703	462464	Tow nsen d depri vatio n inde x at recru itme nt	
9	rs693 1604	Alcoh ol	T	C	T	C	-0.012 1903	-0.042 3081	0.015 579	56429	0.0066 1317	Barrett's esophag us	4.10E- 09	0.002 07309	462464	Tow nsen d depri vatio n inde x at recru itme nt
1 0	rs704 067	BMI	A	G	A	G	0.0120 267	0.0589 608	0.015 3505	56429	0.0001 22543	Barrett's esophag us	3.90E- 09	0.002 0424	462464	Tow nsen d depri vatio n inde x at recru

													itme		
1	rs774	A	G	A	G	0.0118 47	0.0202 887	0.015 0443	56429	0.1774 65	Barrett's esophag us	5.70E- 09	0.002 03391	462464	Two nsen d depri vatio n inde x at recru itme nt
1	0440													Tow nsen d depri vatio n inde x at recru itme nt	
1	rs782	T	C	T	C	-0.016 7416	0.0127 768	0.020 782	56429	0.5386 85	Barrett's esophag us	2.40E- 09	0.002 80676	462464	Two nsen d depri vatio n inde x at recru itme nt
2	5712													Tow nsen d depri vatio n inde x at recru itme nt	
2	8													Tow nsen d depri vatio n inde x at recru itme nt	

																Tow nse d depri vatio n inde x at recru itme nt
1	rs989		G	A	G	A	0.0138	-0.012	0.016	56429	0.4473	Barrett's esophag us	7.40E- 11	0.002	462464	
3	532						559	2055	0622		2					
1	rs990		T	C	T	C	0.0155	-0.019	0.020	56429	0.3441	Barrett's esophag us	7.90E- 09	0.002	462464	
4	706						869	0536	1392							

- Supplementary Table 11 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Leisure/social activities: Pub or social club and risk of Barrett's esophagus.

SN P	Relat ed confon undi	effect_all ele.expos ure	other_all ele.expos ure	effect_all ele.outco me	other_all ele.outco me	beta.e xposu re	beta.o utco me	se.o utco me	samplesi ze.outco me	pval. outco me	outco me	pval.e xposu re	se.ex posu re	samplesi ze.expos ure	expo sure

															Leisu re/soc ial	
4	rs13 9920	BMI	T	C	T	C	0.0061 4932	0.0186 796	0.015 1511	56429	0.2176 18	Barrett's esophag us	1.20E- 11	0.000 90753	461369	activit ies: Pub or social club Leisu re/soc ial
5	rs26 6186 3		C	T	C	T	-0.005 37368	-0.015 0832	0.015 858	56429	0.3415 35	Barrett's esophag us	1.90E- 08	0.000 95672 2	461369	activit ies: Pub or social club Leisu re/soc ial
6	rs35 8645 06	Alcoh ol	G	T	G	T	0.0082 214	0.0275 618	0.022 4755	56429	0.2200 84	Barrett's esophag us	2.00E- 09	0.001 37156	461369	ial activit ies: Pub or

social
club

1	rs94 0159 3	Smoki ng	C	A	C	A	-0.005 31642	-0.048 8776	0.015 0461	56429	0.0011 6011	Barrett's esophag us	5.30E- 09	0.000 91074 6	Leisu re/soc ial	activit ies: Pub or social club			
3																			
•																			
•																			
•																			
●	Supplementary Table 12 Association of the SNPs used as candidate genetic instruments from the GWAS for Mendelian randomization analyses of Leisure/social activities: Sports club or gymDataset and risk of Barrett's esophagus.																		

SN	Relat ed confon ding phen	effect_all ele.expos ure	other_all ele.expos ure	effect_all ele.outco me	other_all ele.outco me	beta.e xposu re	beta.o utco me	se.o utco me	samplesi ze.outco me	pval. outco me	outco me	se.ex posu re	samplesi ze.expos ure	pval.e xposu re	expo sure	
P																

- Supplementary Table 13 UVMR analyse for genetically causal associations of Age completed full time education with BMI.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Body mass index	Age completed full time education	MR Egger	32	0.698	1.16 (0.55, 2.44)
2	Body mass index	Age completed full time education	Weighted median	32	0.002	0.76 (0.64, 0.91)
3	Body mass index	Age completed full time education	Inverse variance weighted	32	4.32E-05	0.74 (0.64, 0.85)
4	Body mass index	Age completed full time education	Simple mode	32	0.247	0.80 (0.56, 1.16)
5	Body mass index	Age completed full time education	Weighted mode	32	0.198	0.80 (0.58, 1.11)

- Supplementary Table 14 UVMR analyse for genetically causal associations of Age completed full time education with MDD.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Major Depressive Disorder	Age completed full time education	MR Egger	35	0.668	0.74 (0.19, 2.91)
2	Major Depressive Disorder	Age completed full time education	Weighted median	35	0.009	0.64 (0.46, 0.89)
3	Major Depressive Disorder	Age completed full time education	Inverse variance weighted	35	0.002	0.63 (0.47, 0.84)
4	Major Depressive Disorder	Age completed full time education	Simple mode	35	0.189	0.57 (0.25, 1.30)
5	Major Depressive Disorder	Age completed full time education	Weighted mode	35	0.068	0.50 (0.24, 1.03)

- Supplementary Table 15 UVMR analyse for genetically causal associations of Age completed full time education with Cigarettes smoked per day.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Cigarettes smoked per day	Age completed full time education	MR Egger	36	0.391	0.74 (0.37, 1.46)
2	Cigarettes smoked per day	Age completed full time education	Weighted median	36	<0.001	0.67 (0.54, 0.83)

3	Cigarettes smoked per day	Age completed full time education	Inverse variance weighted	36	1.93E-08	0.62 (0.52, 0.73)
4	Cigarettes smoked per day	Age completed full time education	Simple mode	36	0.218	0.73 (0.45, 1.19)
5	Cigarettes smoked per day	Age completed full time education	Weighted mode	36	0.219	0.76 (0.50, 1.17)

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- Supplementary Table 16 UVMR analyse for genetically causal associations of Age completed full time education with Alcoholic drinks per week.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Alcoholic drinks per week	Age completed full time education	MR Egger	36	0.960	1.01 (0.75, 1.35)
2	Alcoholic drinks per week	Age completed full time education	Weighted median	36	0.869	0.99 (0.92, 1.08)
3	Alcoholic drinks per week	Age completed full time education	Inverse variance weighted	36	0.492	1.03 (0.95, 1.10)
4	Alcoholic drinks per week	Age completed full time education	Simple mode	36	0.658	0.96 (0.79, 1.16)
5	Alcoholic drinks per week	Age completed full time education	Weighted mode	36	0.575	0.95 (0.79, 1.14)

- Supplementary Table 17 UVMR analyse for genetically causal associations of Average total household income before tax with Sleep duration.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Sleep duration	Age completed full time education	MR Egger	39	0.650	1.09 (0.76, 1.54)
2	Sleep duration	Age completed full time education	Weighted median	39	0.577	1.02 (0.96, 1.08)
3	Sleep duration	Age completed full time education	Inverse variance weighted	39	0.548	1.03 (0.94, 1.12)
4	Sleep duration	Age completed full time education	Simple mode	39	0.804	1.02 (0.90, 1.15)
5	Sleep duration	Age completed full time education	Weighted mode	39	0.738	1.02 (0.91, 1.14)

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- Supplementary Table 18 UVMR analyse for genetically causal associations of Average total household income before tax with BMI.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Body mass index	Average total household income before tax	MR Egger	34	0.275	1.28 (0.83, 1.97)
2	Body mass index	Average total household income before tax	Weighted median	34	<0.001	0.81 (0.72, 0.90)
3	Body mass index	Average total household income before tax	Inverse variance weighted	34	3.66E-09	0.77 (0.71, 0.84)
4	Body mass index	Average total household income before tax	Simple mode	34	0.133	0.84 (0.68, 1.05)
5	Body mass index	Average total household income before tax	Weighted mode	34	0.112	0.83 (0.67, 1.04)

- Supplementary Table 19 UVMR analyse for genetically causal associations of Average total household income before tax with MDD.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Major Depressive Disorder	Average total household income before tax	MR Egger	42	0.706	1.18 (0.51, 2.69)
2	Major Depressive Disorder	Average total household income before tax	Weighted median	42	0.001	0.69 (0.55, 0.86)
3	Major Depressive Disorder	Average total household income before tax	Inverse variance weighted	42	4.85E-05	0.68 (0.56, 0.82)
4	Major Depressive Disorder	Average total household income before tax	Simple mode	42	0.162	0.66 (0.37, 1.17)
5	Major Depressive Disorder	Average total household income before tax	Weighted mode	42	0.640	0.88 (0.52, 1.49)

- Supplementary Table 20 UVMR analyse for genetically causal associations of Average total household income before tax with Cigarettes smoked per day.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Cigarettes smoked per day	Average total household income before tax	MR Egger	43	0.337	1.28 (0.78, 2.09)
2	Cigarettes smoked per day	Average total household income before tax	Weighted median	43	<0.001	0.75 (0.64, 0.88)
3	Cigarettes smoked per day	Average total household income before tax	Inverse variance weighted	43	1.91E-07	0.71 (0.63, 0.81)
4	Cigarettes smoked per day	Average total household income before tax	Simple mode	43	0.177	0.76 (0.51, 1.13)
5	Cigarettes smoked per day	Average total household income before tax	Weighted mode	43	0.345	0.83 (0.58, 1.21)

- Supplementary Table 21 UVMR analyse for genetically causal associations of Average total household income before tax with Alcoholic drinks per week.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Alcoholic drinks per week	Average total household income before tax	MR Egger	40	0.845	0.98 (0.80, 1.20)
2	Alcoholic drinks per week	Average total household income before tax	Weighted median	40	0.757	1.01 (0.95, 1.07)
3	Alcoholic drinks per week	Average total household income before tax	Inverse variance weighted	40	0.085	1.04 (0.99, 1.09)
4	Alcoholic drinks per week	Average total household income before tax	Simple mode	40	0.912	1.01 (0.89, 1.14)
5	Alcoholic drinks per week	Average total household income before tax	Weighted mode	40	0.798	0.99 (0.89, 1.10)

- Supplementary Table 22 UVMR analyse for genetically causal associations of Average total household income before tax with Sleep duration.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Sleep duration	Average total household income before tax	MR Egger	40	0.997	1.00 (0.85, 1.17)
2	Sleep duration	Average total household income before tax	Weighted median	40	0.024	1.05 (1.01, 1.10)
3	Sleep duration	Average total household income before tax	Inverse variance weighted	40	0.015	1.05 (1.01, 1.09)
4	Sleep duration	Average total household income before tax	Simple mode	40	0.214	1.06 (0.97, 1.15)
5	Sleep duration	Average total household income before tax	Weighted mode	40	0.183	1.05 (0.98, 1.14)

- Supplementary Table 23 UVMR analyse for genetically causal associations of BMI with Gastroesophageal reflux disease.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Gastroesophageal reflux disease	Body mass index	MR Egger	63	0.003	1.44 (1.14, 1.80)
2	Gastroesophageal reflux disease	Body mass index	Weighted median	63	1.05E-32	1.68 (1.55, 1.83)
3	Gastroesophageal reflux disease	Body mass index	Inverse variance weighted	63	9.17E-41	1.65 (1.53, 1.77)
4	Gastroesophageal reflux disease	Body mass index	Simple mode	63	1.77E-07	1.73 (1.44, 2.08)
5	Gastroesophageal reflux disease	Body mass index	Weighted mode	63	3.17E-11	1.71 (1.50, 1.95)

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 - Supplementary Table 24 UVMR analyse for genetically causal associations of MDD with Gastroesophageal reflux disease.

	Outcome	Exposure	Method	nSNP	Pval	OR (95% CI)
1	Gastroesophageal reflux disease	Major Depressive Disorder	MR Egger	4	0.306	1.41 (0.86, 2.31)
2	Gastroesophageal reflux disease	Major Depressive Disorder	Weighted median	4	6.04E-09	1.46 (1.29, 1.67)
3	Gastroesophageal reflux disease	Major Depressive Disorder	Inverse variance weighted	4	1.68E-14	1.47 (1.33, 1.62)
4	Gastroesophageal reflux disease	Major Depressive Disorder	Simple mode	4	0.025	1.46 (1.22, 1.75)
5	Gastroesophageal reflux disease	Major Depressive Disorder	Weighted mode	4	0.019	1.46 (1.24, 1.71)

- Supplementary Table 25 UVMR analyse for genetically causal associations of Cigarettes smoked per day with Gastroesophageal reflux disease.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Gastroesophageal reflux disease	Cigarettes smoked per day	MR Egger	6	0.428	0.93 (0.78, 1.10)
2	Gastroesophageal reflux disease	Cigarettes smoked per day	Weighted median	6	0.136	1.04 (0.99, 1.10)
3	Gastroesophageal reflux disease	Cigarettes smoked per day	Inverse variance weighted	6	0.266	1.09 (0.93, 1.28)
4	Gastroesophageal reflux disease	Cigarettes smoked per day	Simple mode	6	0.121	1.65 (0.97, 2.79)
5	Gastroesophageal reflux disease	Cigarettes smoked per day	Weighted mode	6	0.472	1.02 (0.97, 1.08)

- Supplementary Table 26 UVMR analyse for genetically causal associations of Alcoholic drinks per week with Gastroesophageal reflux disease.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Gastroesophageal reflux disease	Alcoholic drinks per week	MR Egger	9	0.670	0.77 (0.24, 2.44)
2	Gastroesophageal reflux disease	Alcoholic drinks per week	Weighted median	9	0.380	0.87 (0.64, 1.19)
3	Gastroesophageal reflux disease	Alcoholic drinks per week	Inverse variance weighted	9	0.349	0.87 (0.65, 1.16)
4	Gastroesophageal reflux disease	Alcoholic drinks per week	Simple mode	9	0.953	1.02 (0.60, 1.71)
5	Gastroesophageal reflux disease	Alcoholic drinks per week	Weighted mode	9	0.795	0.95 (0.65, 1.39)

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- Supplementary Table 27 UVMR analyse for genetically causal associations of Sleep duration with Gastroesophageal reflux disease.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Gastroesophageal reflux disease	Sleep duration	MR Egger	44	0.191	0.59 (0.27, 1.29)
2	Gastroesophageal reflux disease	Sleep duration	Weighted median	44	0.086	0.83 (0.67, 1.03)
3	Gastroesophageal reflux disease	Sleep duration	Inverse variance weighted	44	0.001	0.72 (0.59, 0.88)
4	Gastroesophageal reflux disease	Sleep duration	Simple mode	44	0.980	1.01 (0.63, 1.61)
5	Gastroesophageal reflux disease	Sleep duration	Weighted mode	44	0.482	0.88 (0.61, 1.27)

- Supplementary Table 28 UVMR analyse for genetically causal associations of BMI with Barrett's esophagus.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Barrett's esophagus	Body mass index	MR Egger	63	0.050	1.81 (1.01, 3.23)
2	Barrett's esophagus	Body mass index	Weighted median	63	3.61E-05	1.71 (1.33, 2.21)
3	Barrett's esophagus	Body mass index	Inverse variance weighted	63	2.51E-10	1.87 (1.54, 2.27)
4	Barrett's esophagus	Body mass index	Simple mode	63	<0.001	3.22 (1.73, 5.99)
5	Barrett's esophagus	Body mass index	Weighted mode	63	0.089	1.55 (0.94, 2.55)

- Supplementary Table 29 UVMR analyse for genetically causal associations of MDD with Barrett's esophagus.

Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
Barrett's esophagus	Major Depressive Disorder	Inverse variance weighted	2	0.005	1.95 (1.22, 3.09)

- Supplementary Table 30 UVMR analyse for genetically causal associations of Cigarettes smoked per day with Barrett's esophagus.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Barrett's esophagus	Cigarettes smoked per day	MR Egger	14	0.816	1.04 (0.77, 1.39)
2	Barrett's esophagus	Cigarettes smoked per day	Weighted median	14	0.072	1.16 (0.99, 1.36)
3	Barrett's esophagus	Cigarettes smoked per day	Inverse variance weighted	14	0.021	1.24 (1.03, 1.49)
4	Barrett's esophagus	Cigarettes smoked per day	Simple mode	14	0.721	1.08 (0.70, 1.68)
5	Barrett's esophagus	Cigarettes smoked per day	Weighted mode	14	0.149	1.14 (0.96, 1.34)

- Supplementary Table 31 UVMR analyse for genetically causal associations of Alcoholic drinks per week with Barrett's esophagus.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Barrett's esophagus	Alcoholic drinks per week	MR Egger	18	0.417	2.51 (0.29, 21.91)
2	Barrett's esophagus	Alcoholic drinks per week	Weighted median	18	0.219	1.56 (0.77, 3.19)
3	Barrett's esophagus	Alcoholic drinks per week	Inverse variance weighted	18	0.354	1.34 (0.72, 2.51)
4	Barrett's esophagus	Alcoholic drinks per week	Simple mode	18	0.308	2.10 (0.53, 8.43)
5	Barrett's esophagus	Alcoholic drinks per week	Weighted mode	18	0.241	1.78 (0.70, 4.52)

- Supplementary Table 32 UVMR analyse for genetically causal associations of Sleep duration with Barrett's esophagus.

	Outcome	Exposure	Method	nSNP	Pval	OR ((95% CI)
1	Barrett's esophagus	Sleep duration	MR Egger	23	0.070	0.00 (1.40E-05, 1.17)
2	Barrett's esophagus	Sleep duration	Weighted median	23	0.382	0.63 (0.22, 1.78)

3	Barrett's esophagus	Sleep duration	Inverse variance weighted	23	0.380	0.55 (0.14, 2.10)
4	Barrett's esophagus	Sleep duration	Simple mode	23	0.842	0.81 (0.10, 6.27)
5	Barrett's esophagus	Sleep duration	Weighted mode	23	0.870	0.87 (0.17, 4.51)

Supplementary Table 33 The results of the mediation effect test.

Exposure	Mediator	Outcome	Total Effect: β_c (95% CI) *	Direct Effect: β_a (95% CI) †	Direct Effect: β_b (95% CI) †	Mediation Effect (95% CI) §	Proportion mediated
Educational attainment	MDD	GERD	-1.468 (1.694, -1.242)	-0.465(-0.321, -0.143)	0.343(-0.059, 0.212)	-0.159 (-0.278, -0.057)	10.83%
Educational attainment	MDD	BE	-2.203 (-2.906, -1.501)	-0.465(-0.004, 0.043)	0.502(-0.142, 0.869)	-0.233 (-0.482, -0.053)	10.58%
Educational attainment	Smoking	BE	-2.203 (-2.906, -1.501)	0.161(-0.026, 0.029)	0.259(-0.26, 0.778)	-0.077 (-0.161, -0.005)	3.50%
Household income	Sleep	GERD	-0.048 (-1.074, -0.525)	-0.363(-0.066, -0.01)	-0.096(-0.631, 0.439)	0.017 (0.001, 0.041)	9.75%

*Total Effect β_c : the effect of educational attainment on the risk of EC, BE, and GERD. Only the results for which the total effect is statistically significant are shown here.

†Direct effect β_a : the effect of educational attainment on the risk of modifiable risk factors.

‡Direct effect β_b : the effect of modifiable risk factors on the risk of EC, BE, and GERD after adjusting two educational attainment traits.

§Mediation effect: $\beta_a \times \beta_b$; 95% CI of mediation effect: $\beta_a \times \beta_b \pm 1.96 \times \sqrt{\beta_b^2 \times se_a^2 + \beta_a^2 \times se_b^2}$.

- Supplementary Table 34 Heterogeneity and pleiotropy evaluations for genetically causal associations of socioeconomic traits with Gastroesophageal reflux disease risk.

Exposure	Outcome	nSNP	Heterogeneity		Pleiotropy		
			Q	P	Intercept	SE	P
Age completed full time education	Gastroesophageal reflux disease	27	7.569	0.134	0.01	0.0541	0.342
Job involves heavy manual or physical work	Gastroesophageal reflux disease	10	20.339	0.016	0.032	0.013	0.041
Average total household income before tax	Gastroesophageal reflux disease	9	15.49	0.05	-0.011	0.022	0.649

Townsend deprivation index at recruitment	Gastroesophageal reflux disease	6	9.074	0.106	0.041	0.019	0.09
Leisure/social activities: Pub or social club	Gastroesophageal reflux disease	3	2.179	0.336	-0.024	0.025	0.512
Leisure/social activities: Sports club or gym	Gastroesophageal reflux disease	4	5.27	0.153	-0.009	0.034	0.825

- Supplementary Table 35 Heterogeneity and pleiotropy evaluations for genetically causal associations of socioeconomic traits with Barrett's esophagus.

Exposure	Outcome	nSNP	Heterogeneity		Pleiotropy		
			Q	P	Intercept	SE	P
Age completed full time education	Barrett's esophagus	12	8.714	0.648	-0.011	0.019	0.579
Job involves heavy manual or physical work	Barrett's esophagus	11	13.894	0.178	0.037	0.038	0.353
Average total household income before tax	Barrett's esophagus	17	34.101	0.005	-0.009	0.026	0.736
Townsend deprivation index at recruitment	Barrett's esophagus	8	6.652	0.466	0.086	0.052	0.148
Leisure/social activities: Pub or social club	Barrett's esophagus	6	4.152	0.528	0.012	0.063	0.861
Leisure/social activities: Sports club or gym	Barrett's esophagus	3	6.978	0.031	0.324	0.498	0.633

- Supplementary Table 36 Pleiotropy evaluations for genetically causal associations of Age completed full time education with modifiable risk factors.

Exposure	Outcome	nSNP	Pleiotropy		
			Intercept	SE	P
Age completed full time education	Body mass index	32	-0.006	0.005	0.237
Age completed full time education	Major Depressive Disorder	35	-0.002	0.01	0.814
Age completed full time education	Cigarettes smoked per day	36	-0.003	0.005	0.603
Age completed full time education	Alcoholic drinks per week	36	<0.001	0.002	0.904
Age completed full time education	Sleep duration]	39	-0.001	0.002	0.364

- Supplementary Table 37 Pleiotropy evaluations for genetically causal associations of Average total household income before tax with modifiable risk factors.

Exposure	Outcome	nSNP	Pleiotropy
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			Intercept	SE	P
Average total household income before tax	Body mass index	34	-0.01	0.004	0.027
Average total household income before tax	Major Depressive Disorder	42	-0.011	0.008	0.191
Average total household income before tax	Cigarettes smoked per day	43	-0.012	0.005	0.021
Average total household income before tax	Alcoholic drinks per week	40	0.001	0.002	0.546
Average total household income before tax	Sleep duration]	40	0.001	0.002	0.687

- Supplementary Table 38 Pleiotropy evaluations for genetically causal associations of modifiable risk factors with Gastroesophageal reflux disease.

Exposure	Outcome	nSNP	Pleiotropy		
			Intercept	SE	P
Body mass index	Gastroesophageal reflux disease	63	0.004	0.003	0.218
Major Depressive Disorder	Gastroesophageal reflux disease	4	0.002	0.013	0.88
Cigarettes smoked per day	Gastroesophageal reflux disease	6	0.019	0.008	0.067
Alcoholic drinks per week	Gastroesophageal reflux disease	9	0.002	0.009	0.833
Sleep duration]	Gastroesophageal reflux disease	44	-0.002	0.005	0.708

- Supplementary Table 39 Pleiotropy evaluations for genetically causal associations of modifiable risk factors with Barrett's esophagus.

Exposure	Outcome	nSNP	Pleiotropy		
			Intercept	SE	P
Body mass index	Barrett's esophagus	63	0.001	0.008	0.903
Major Depressive Disorder	Barrett's esophagus	2	0.015	0.01	0.154
Cigarettes smoked per day	Barrett's esophagus	14	-0.01	0.017	0.563
Alcoholic drinks per week	Barrett's esophagus	18	-0.01	0.017	0.563
Sleep duration]	Barrett's esophagus	23	-0.009	0.012	0.48