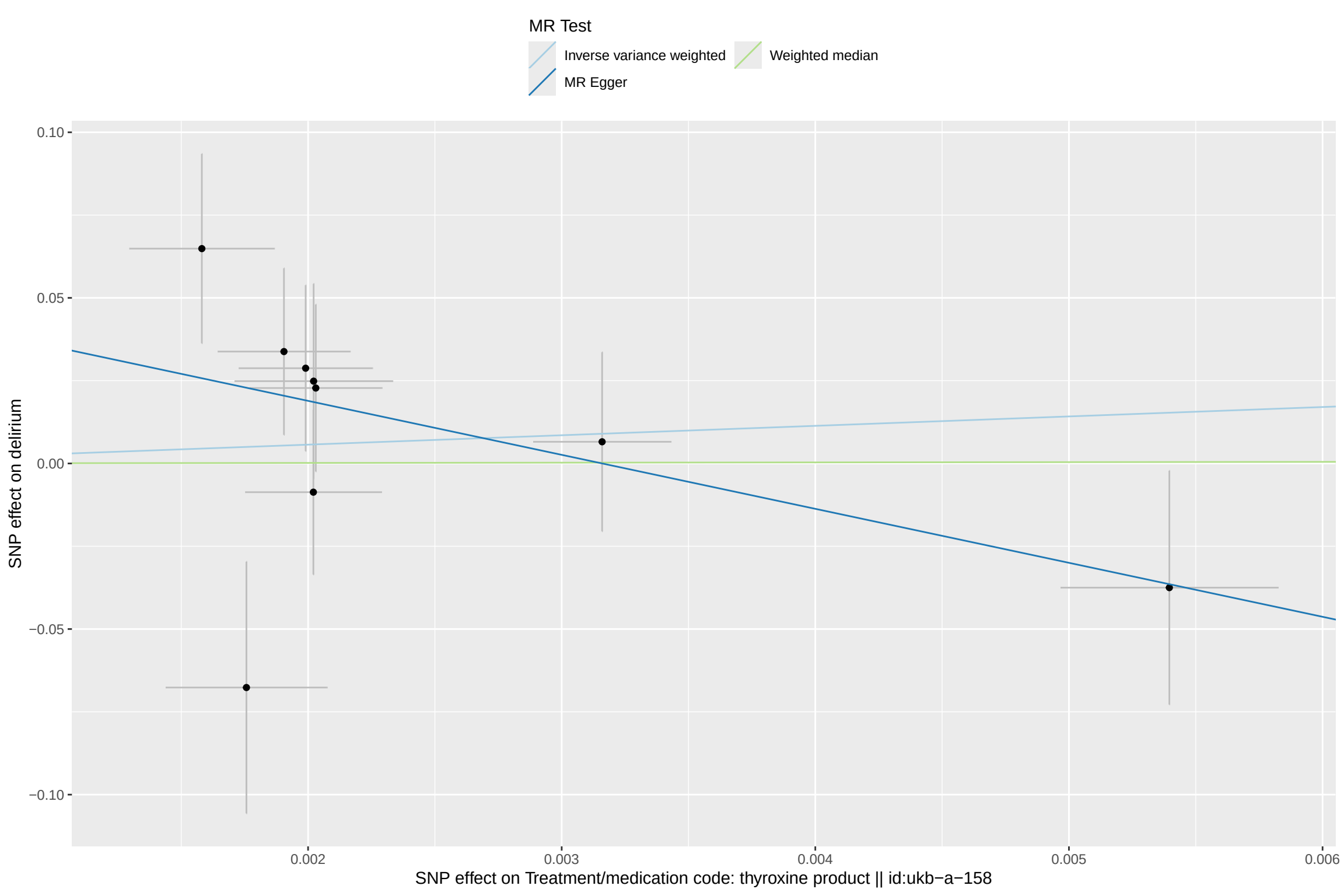
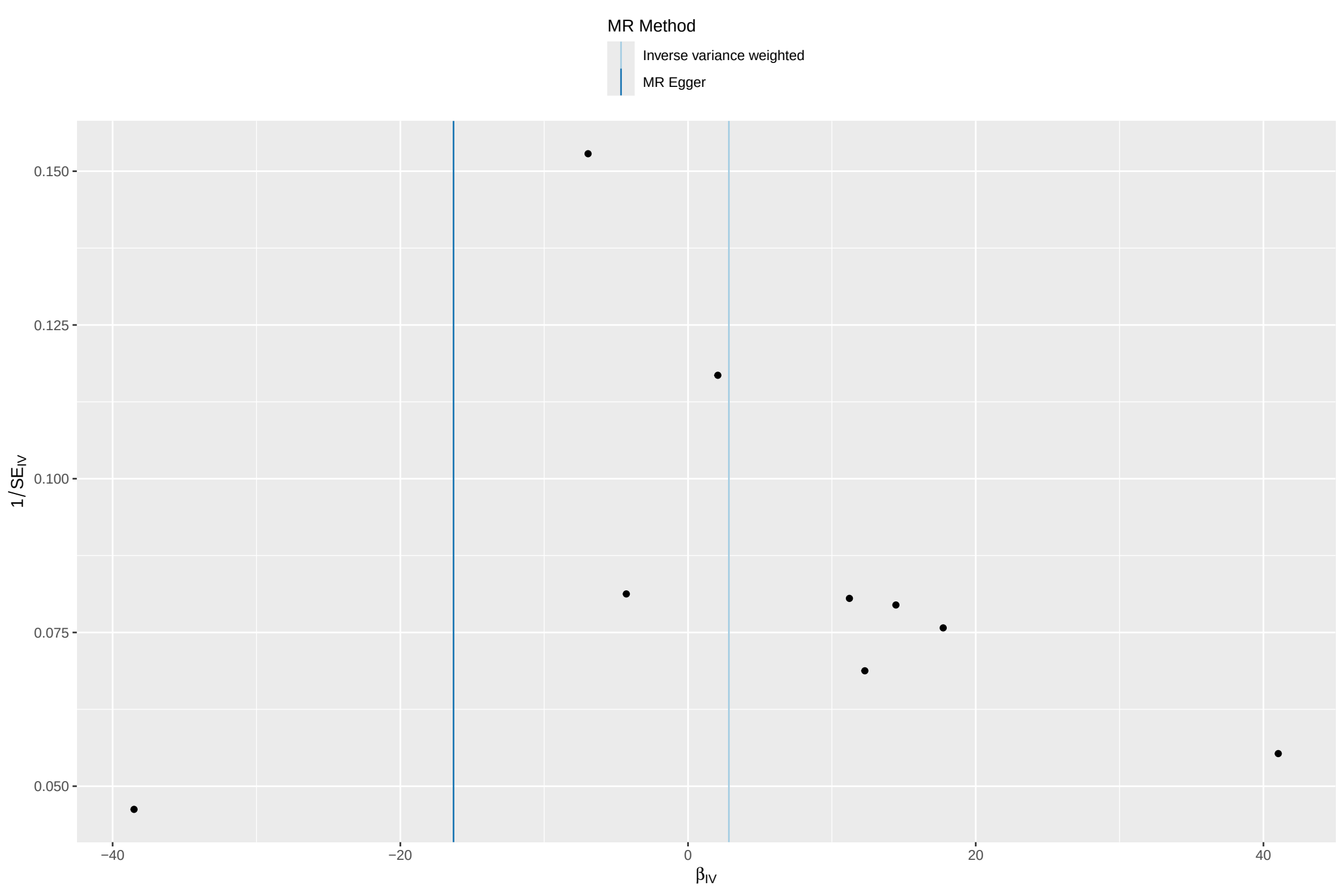
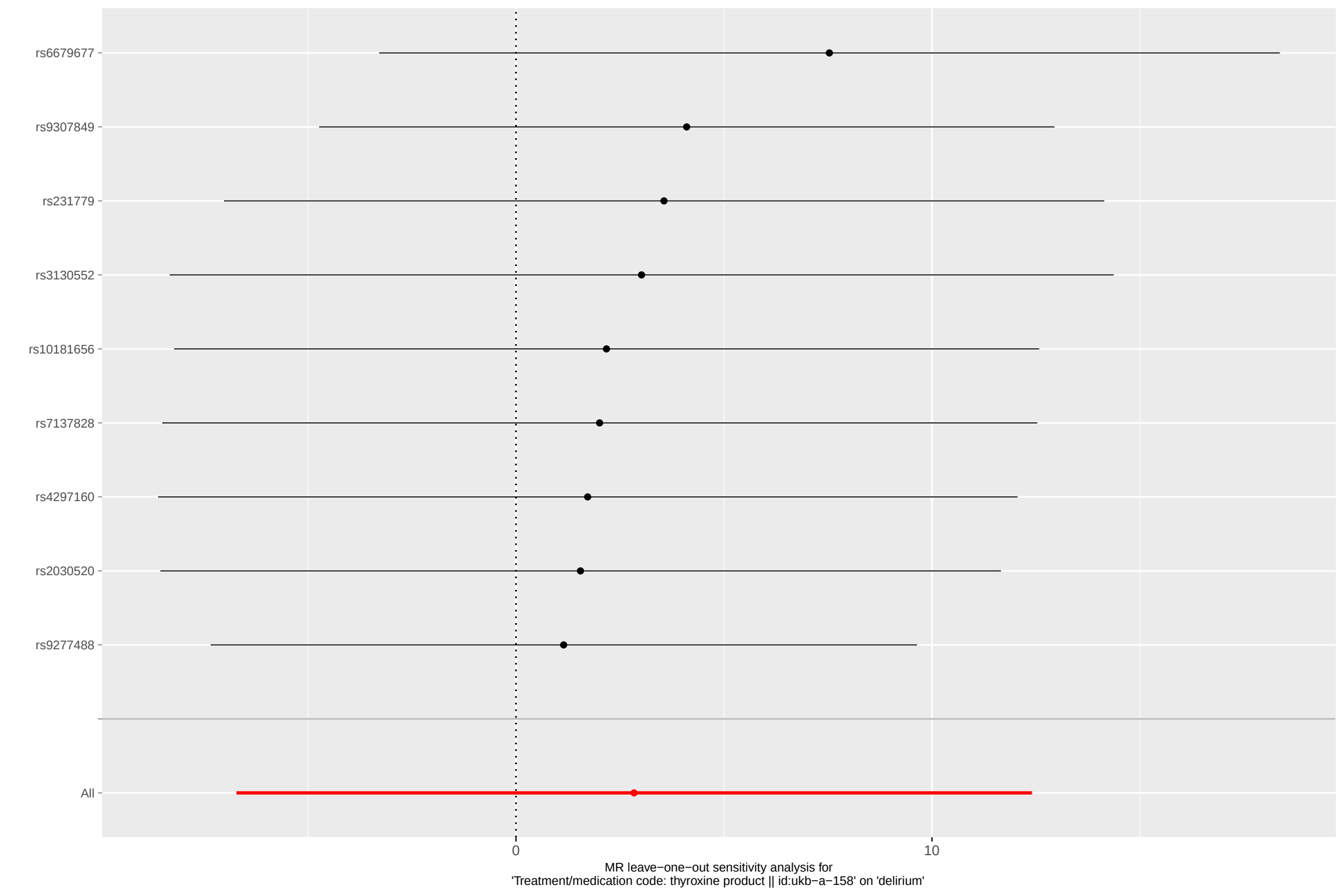
	method	nsnp	b	se	pval
1	MR Egger	9	−16.3010488103409	11.3861828122943	0.195339999580618
2	Weighted median	9	0.080033499915876	5.29387579360599	0.98793793734855
3	Inverse variance weighted	9	2.83955635618228	4.87450138977475	0.560207935933683
4	Simple mode	9	11.5515554458364	8.0560256127842	0.189506367051704
5	Weighted mode	9	−2.58495839351618	5.43787561527877	0.647237375600855

	type	method	Q	Q_df	Q_pval
1	heterogeneity	MR Egger	9.31528843326968	7	0.230805206893754
2	heterogeneity	Inverse variance weighted	13.7003619053626	8	0.0899174996029159

	type	exposure	egger_intercept	se	pval
1	pleiotropy_test	Treatment/medication code: thyroxine product || id:ukb−a−158	0.05150491	0.02837327	0.1123457


MR Test
Inverse variance weighted
Weighted median
MR Egger
0.10
0.05
0.00
−0.05
−0.10
SNP effect on delirium
0.002
0.003
0.004
0.005
0.006
SNP effect on Treatment/medication code: thyroxine product || id:ukb−a−158


MR Method
Inverse variance weighted
MR Egger
0.150
0.125
0.100
0.075
0.050
1/SE_IV
−40
−20
0
20
40
β_IV


rs6679677
rs9307849
rs231779
rs3130552
rs10181656
rs7137828
rs4297160
rs2030520
rs9277488
All
0
10
MR leave−one−out sensitivity analysis for
'Treatment/medication code: thyroxine product || id:ukb−a−158' on 'delirium'