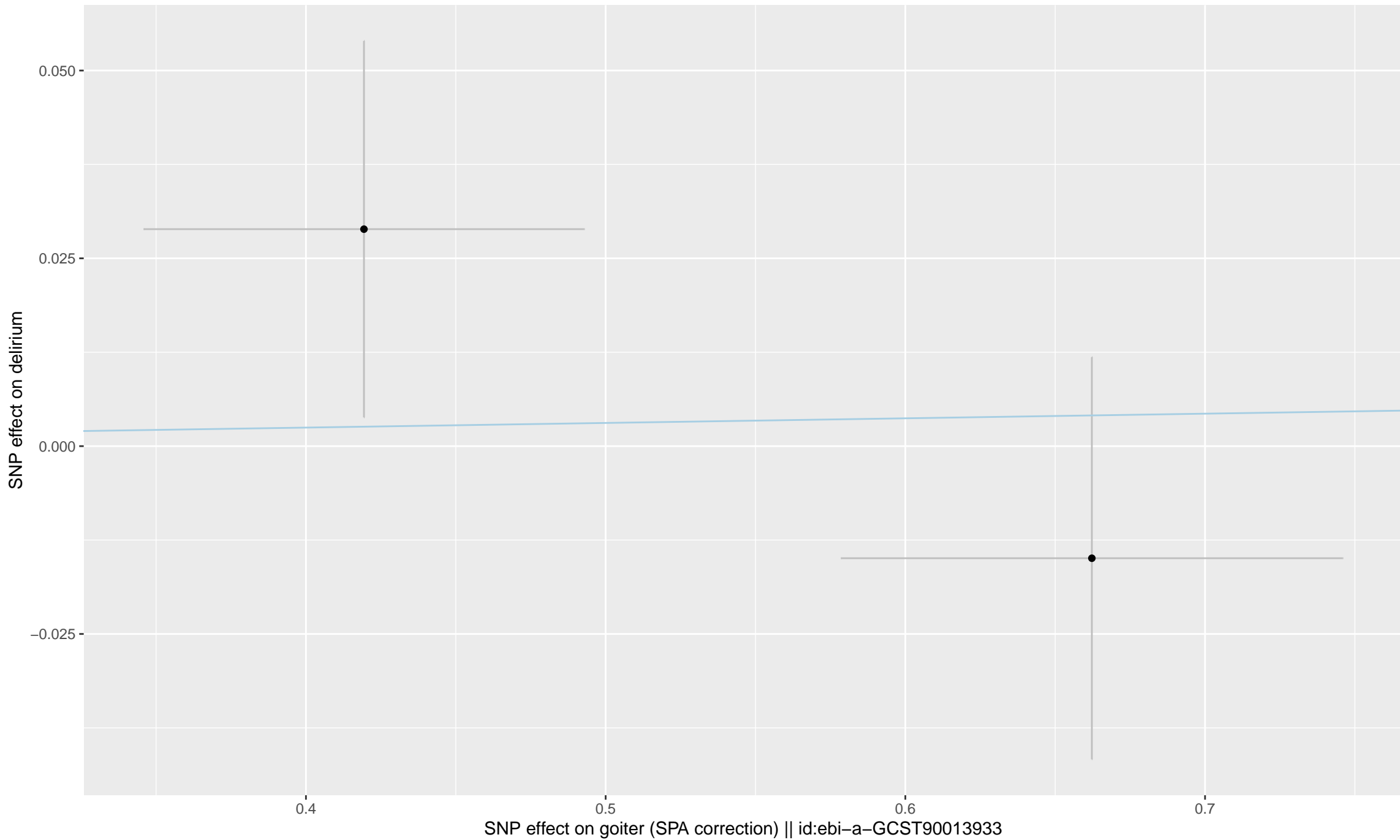


		method	nsnp	b	se	pval	
1		Inverse variance weighted	2	0.00617537161860756	0.0424251928889275	0.884269481219748	
2							
3							
4							
5							
6		type	method	Q	Q_df	Q_pval	
7	1	heterogeneity	Inverse variance weighted	1.60629996373517	1	0.205012694057542	
8	2						
9	3						
10		type	exposure		egger_intercept	se	pval
11	1	pleiotropy_test	goiter (SPA correction) id:ebi-a-GCST90013933		NA	NA	NA
12	6						
13	7						
14	8						
15	9						
16							
17							
18							
19							
20							
21							

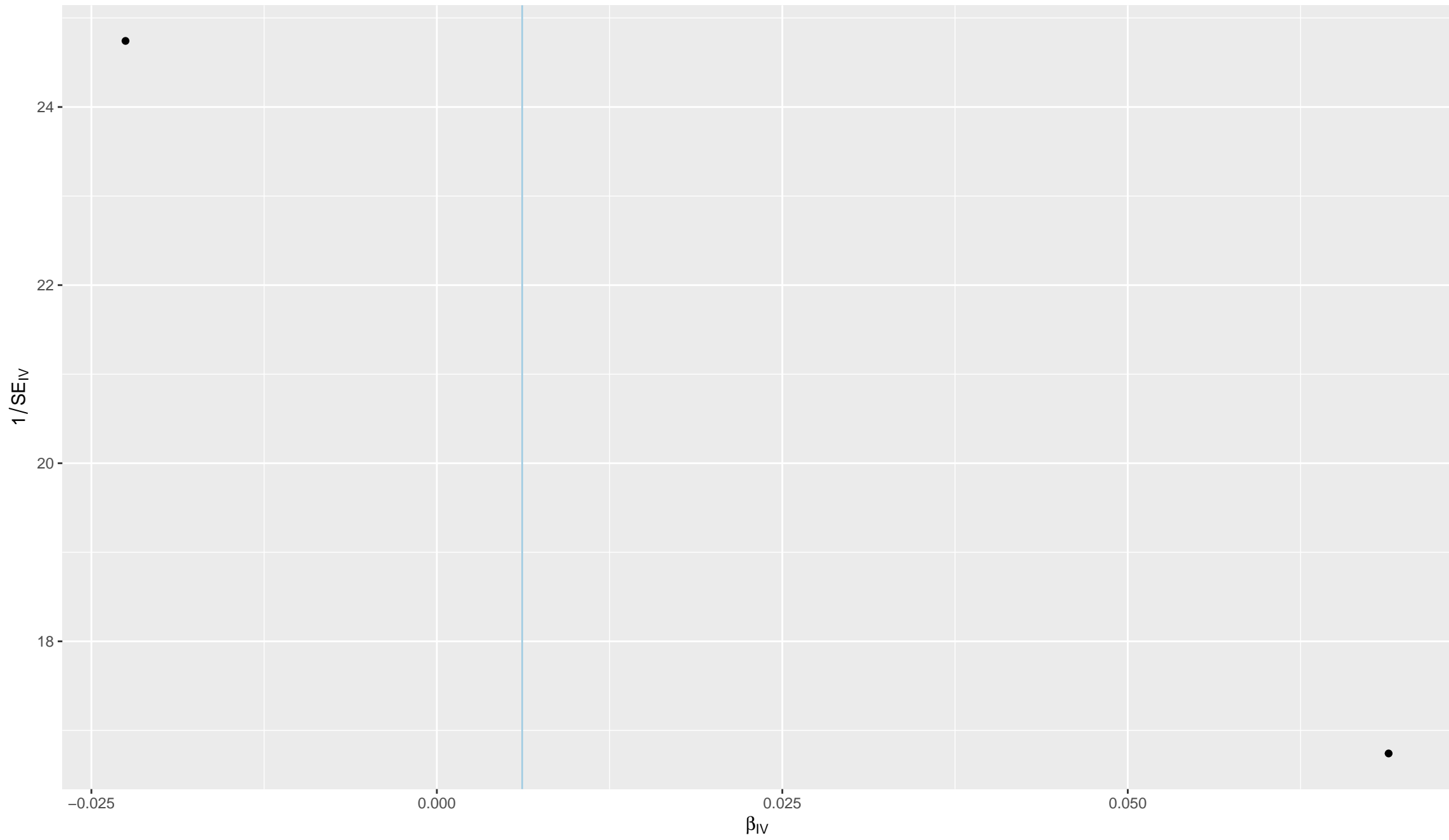
MR Test

Inverse variance weighted



MR Method

- Inverse variance weighted
- MR Egger



Insufficient number of SNPs