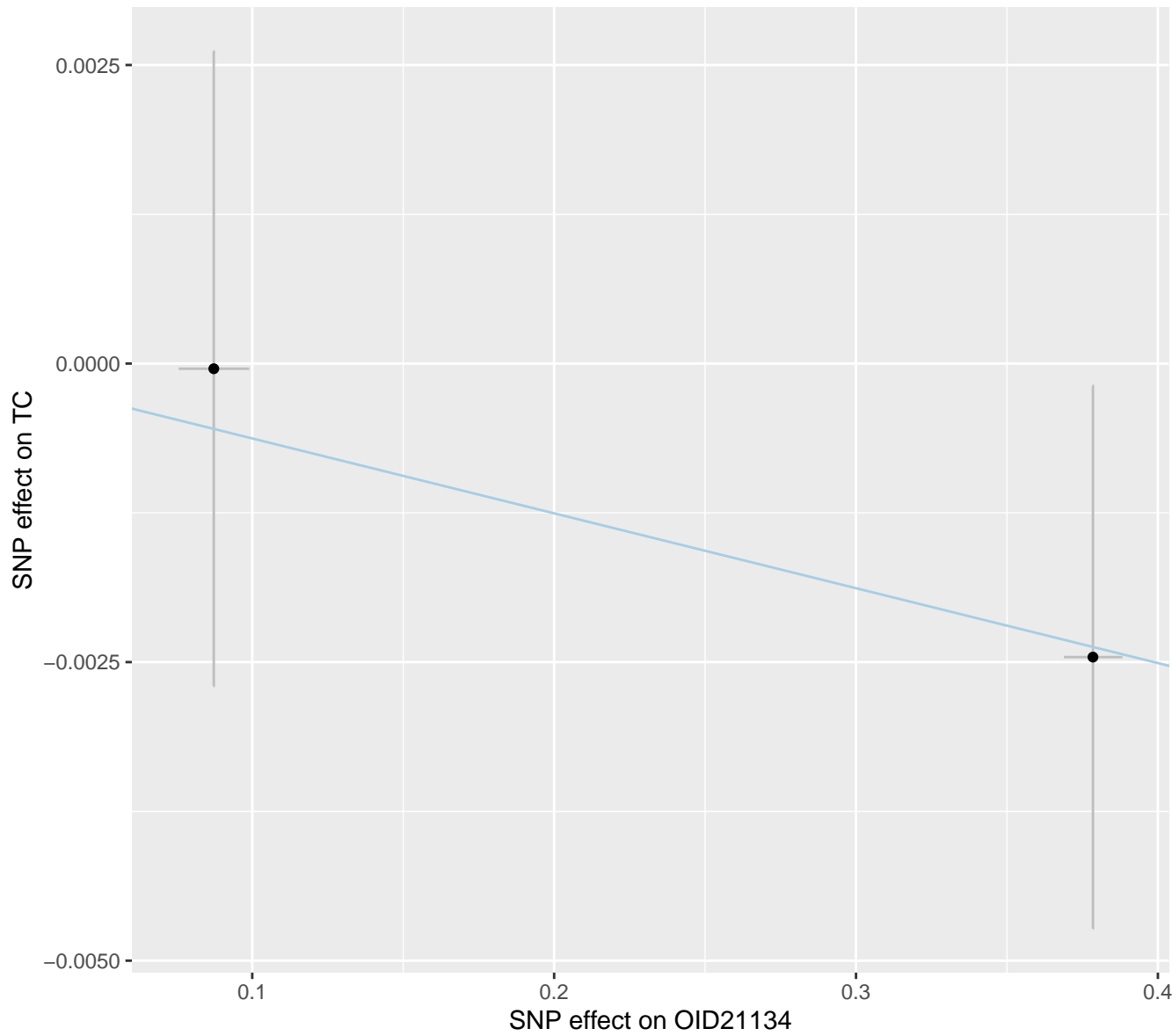


MR Estimate



Inverse variance weighted



Insufficient number of SNPs