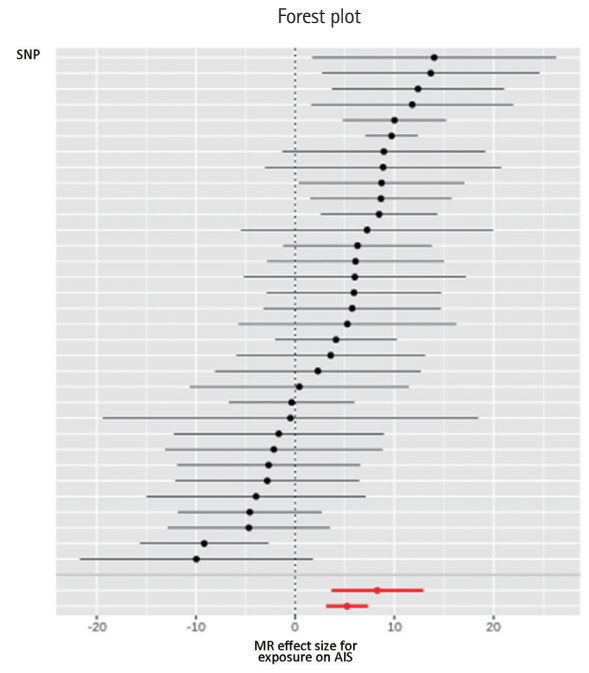
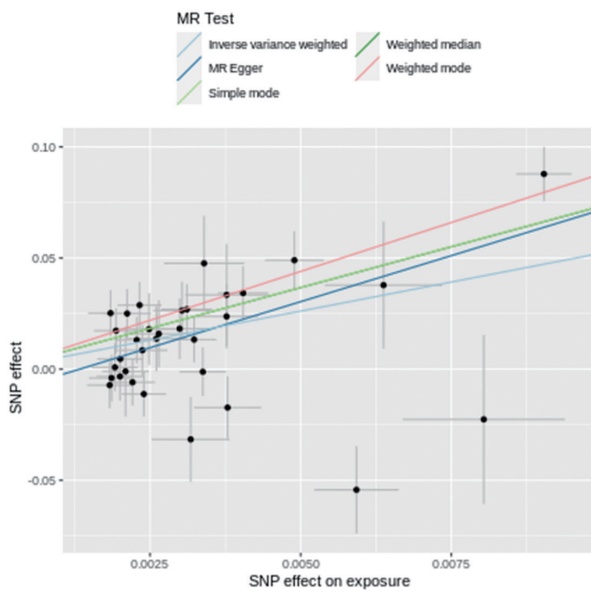


A



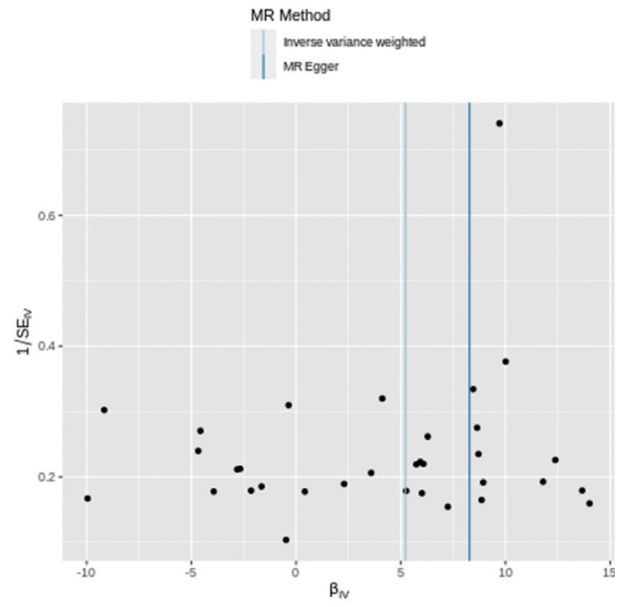
B

Comparison of results using different MR methods



C

Funnel plot of IVW and MR-Egger regression



D

Supplementary Figure 9. The leave-one-out analysis, single nucleotide polymorphism (SNP) analysis, sensitivity analysis, and funnel plot for cardiac arrhythmias, chronic obstructive pulmonary disease co-morbidities and any ischemic stroke (AIS). (A) Leave-one-out analysis, (B) forest plot, (C) comparison of results using different Mendelian randomization (MR) methods, (D) funnel plot of inverse-variance weighted (IVW) and MR-Egger regression. SE, standard error.