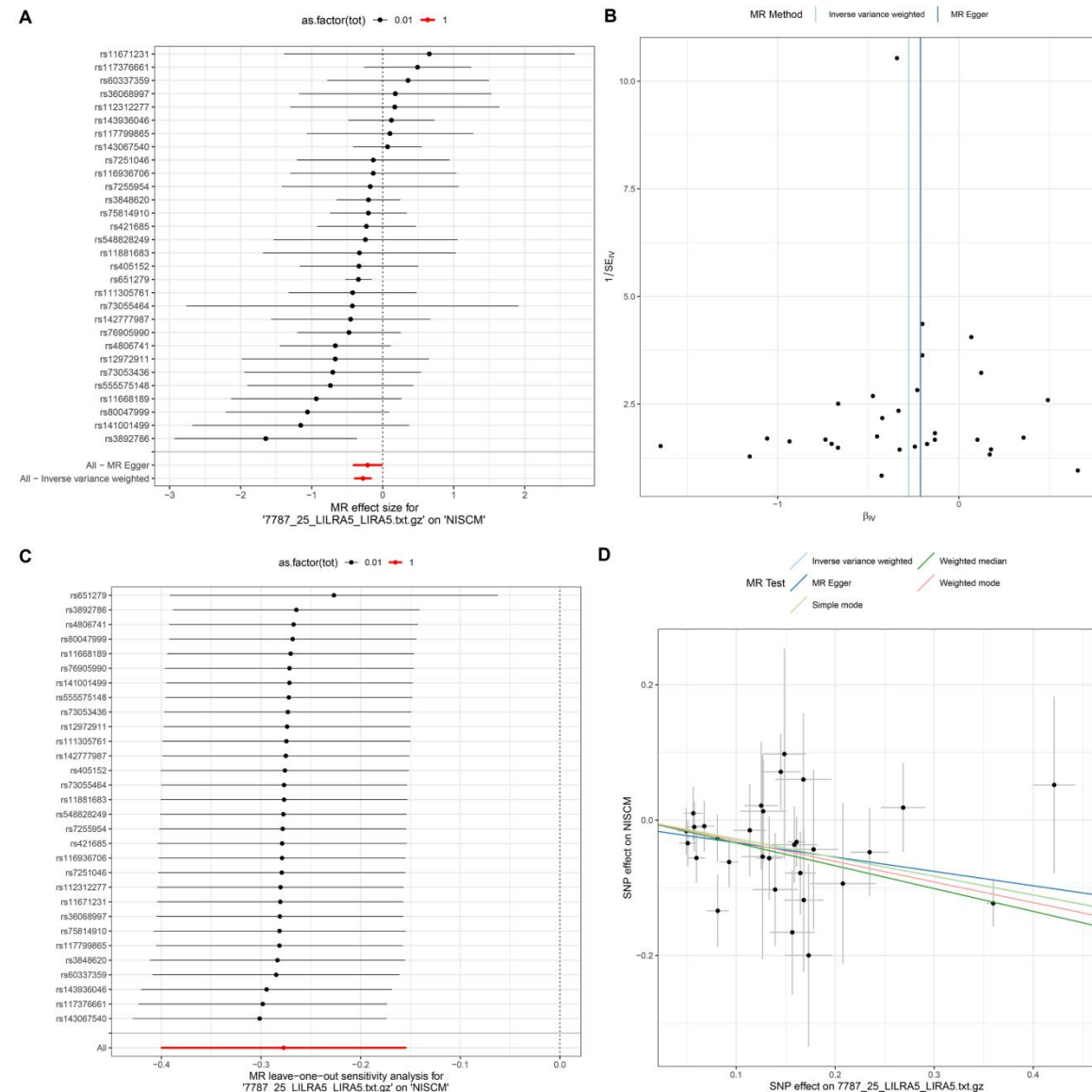
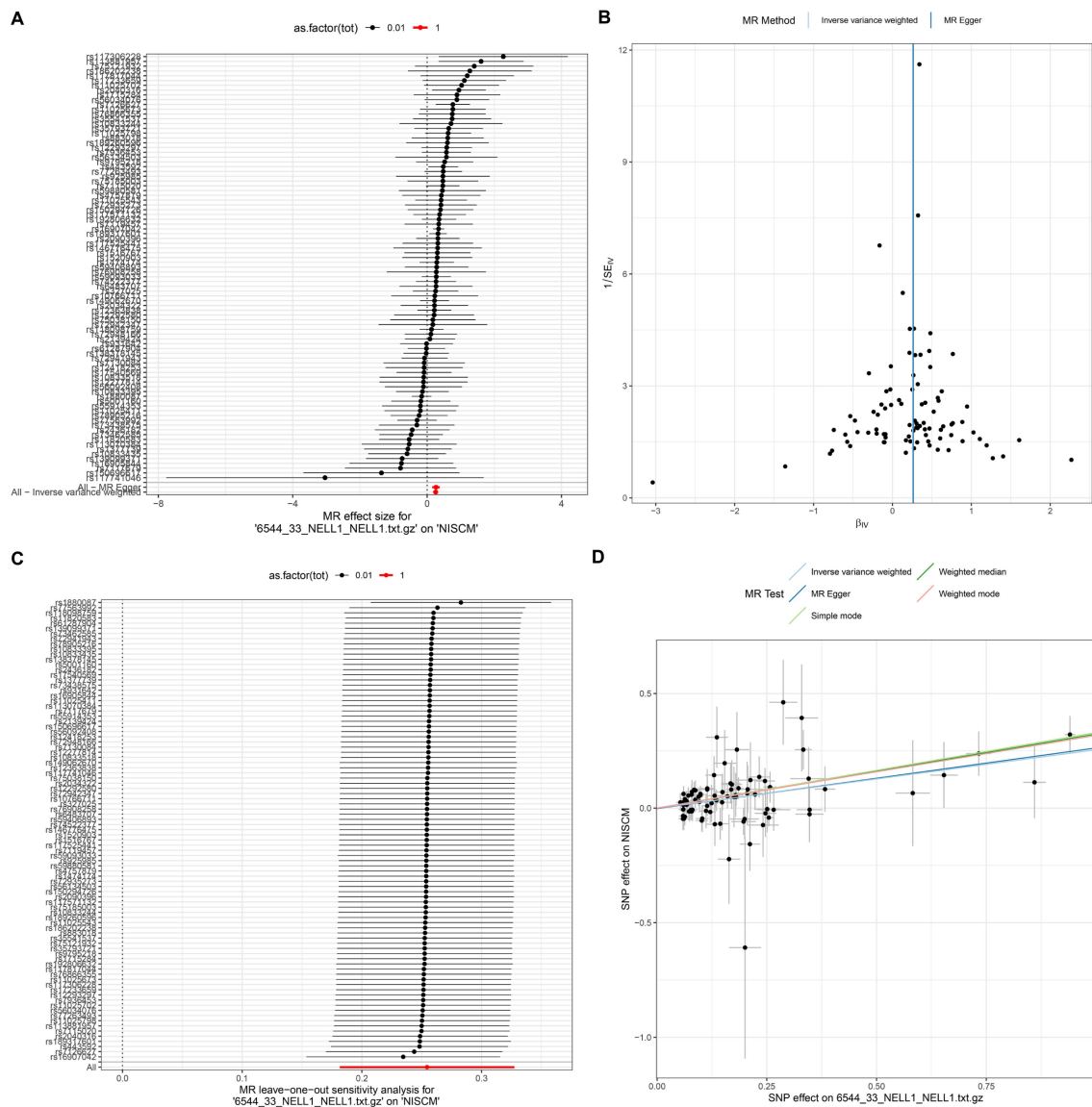


Figure S1. Heterogeneity or pleiotropy for the 16 identified plasma proteins.

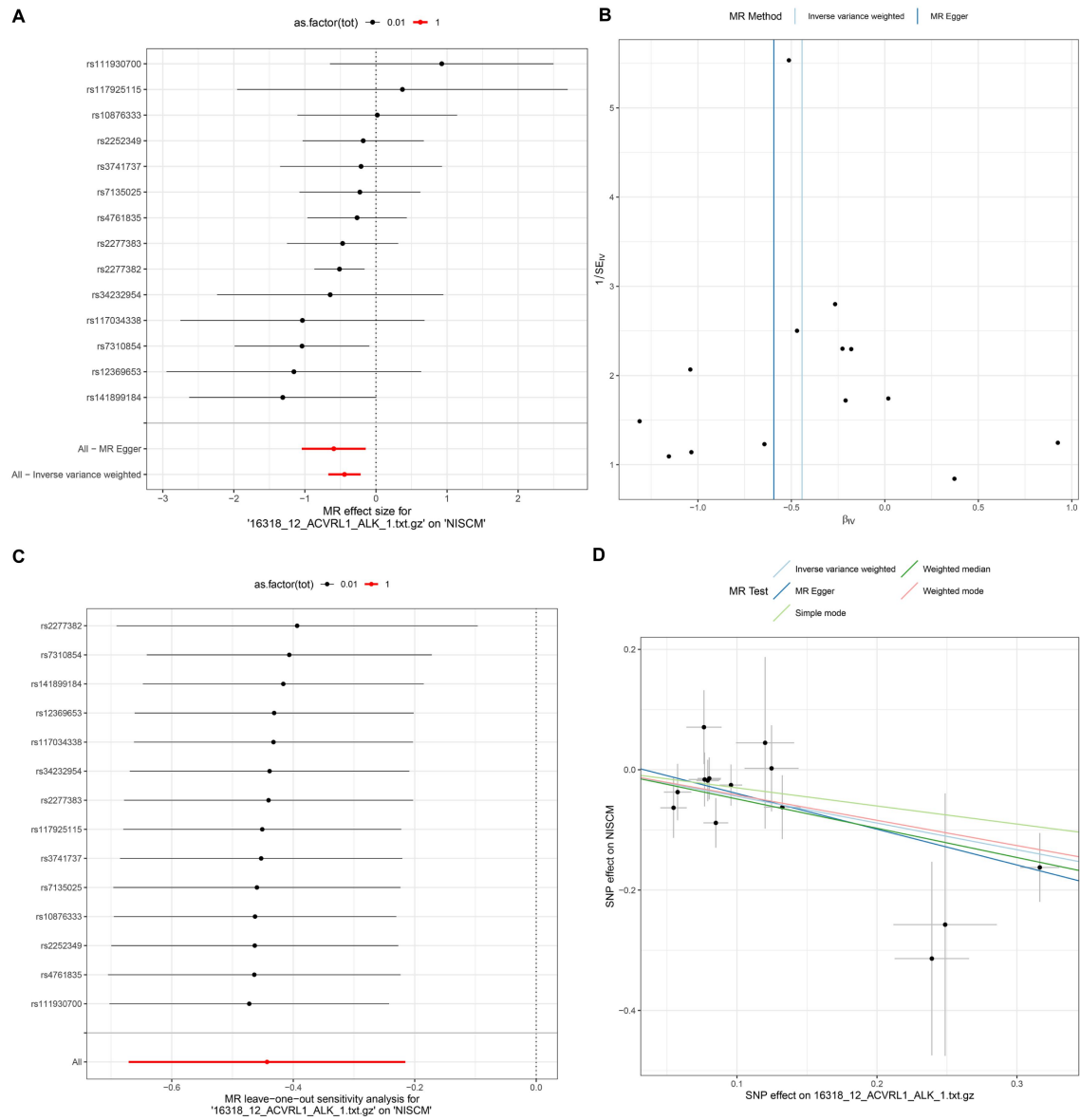
(A) Variant-specific inverse-variance forest plots were used to estimate the causal relationships between each of the 16 circulating plasma proteins and nonischemic cardiomyopathy (NISC) separately. (B) Funnel plot of causal association between each of the 16 circulating plasma proteins and NISC separately. (C) Leave-one-out plot was used to separately assess if a single variant is driving the association between each of the 16 circulating plasma proteins and NISC. (D) Mendelian Randomization sensitivity plots of the causal effect of each of the 16 circulating plasma proteins on NISC.



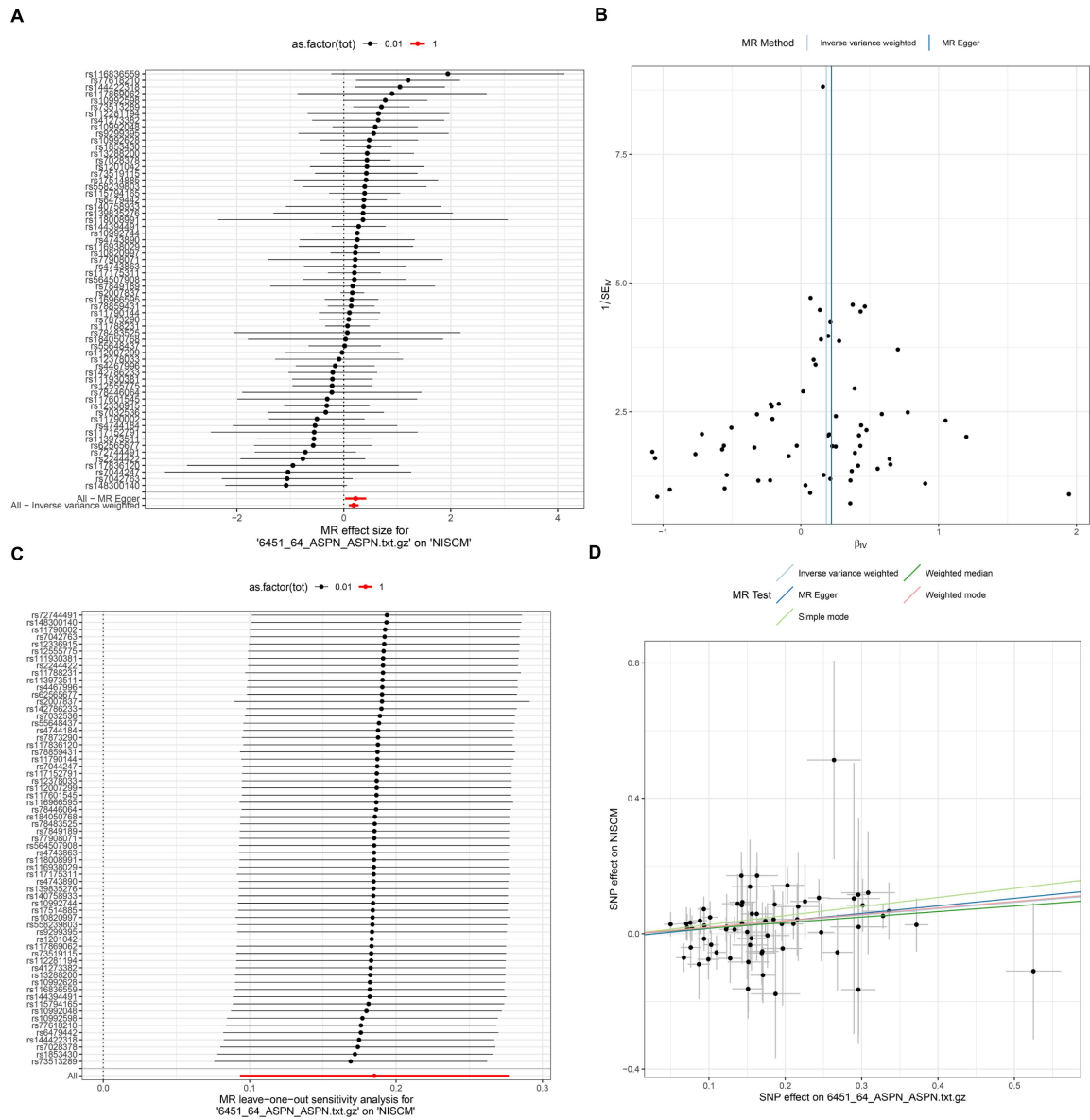
LILRA5



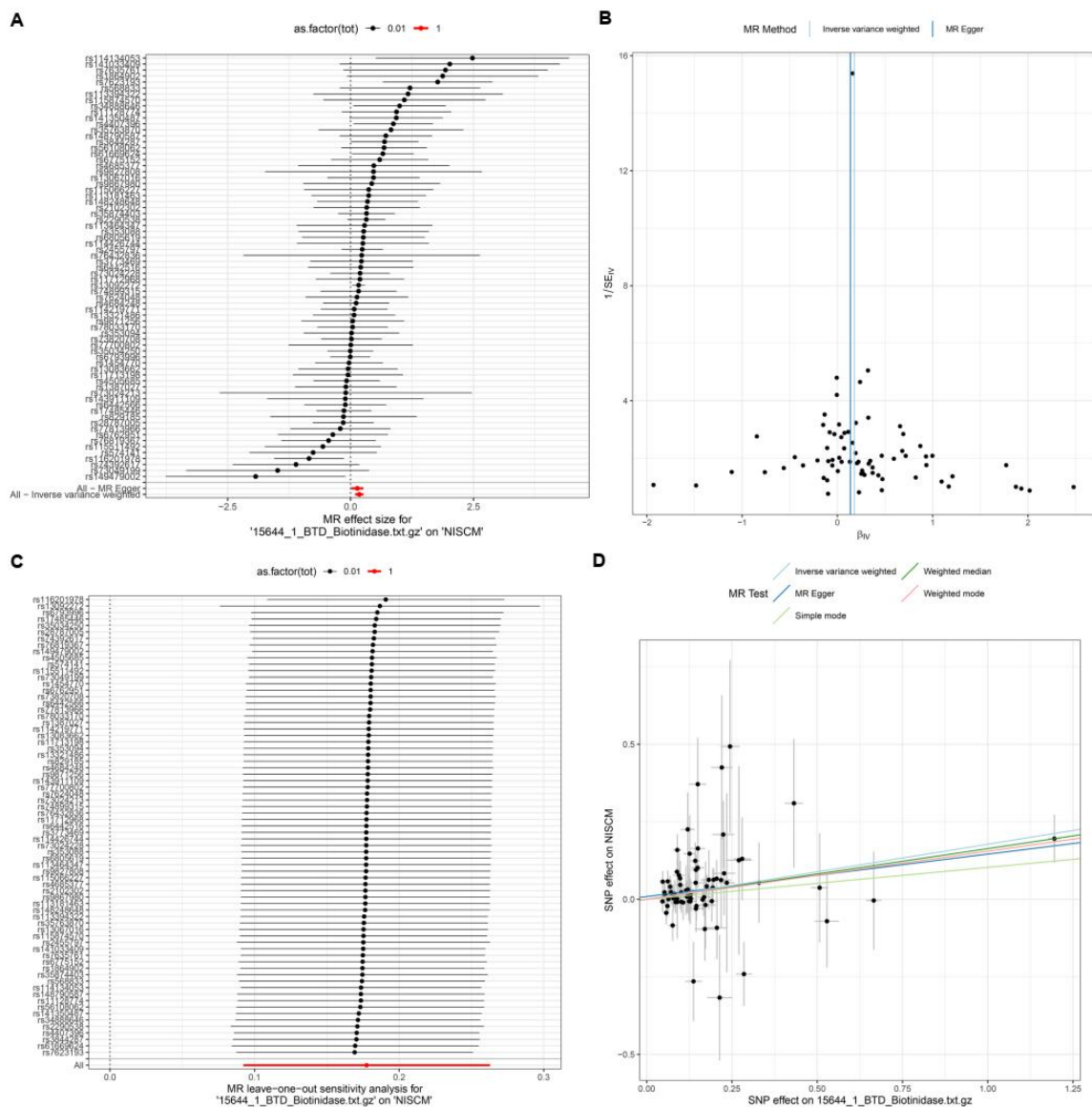
NELL1



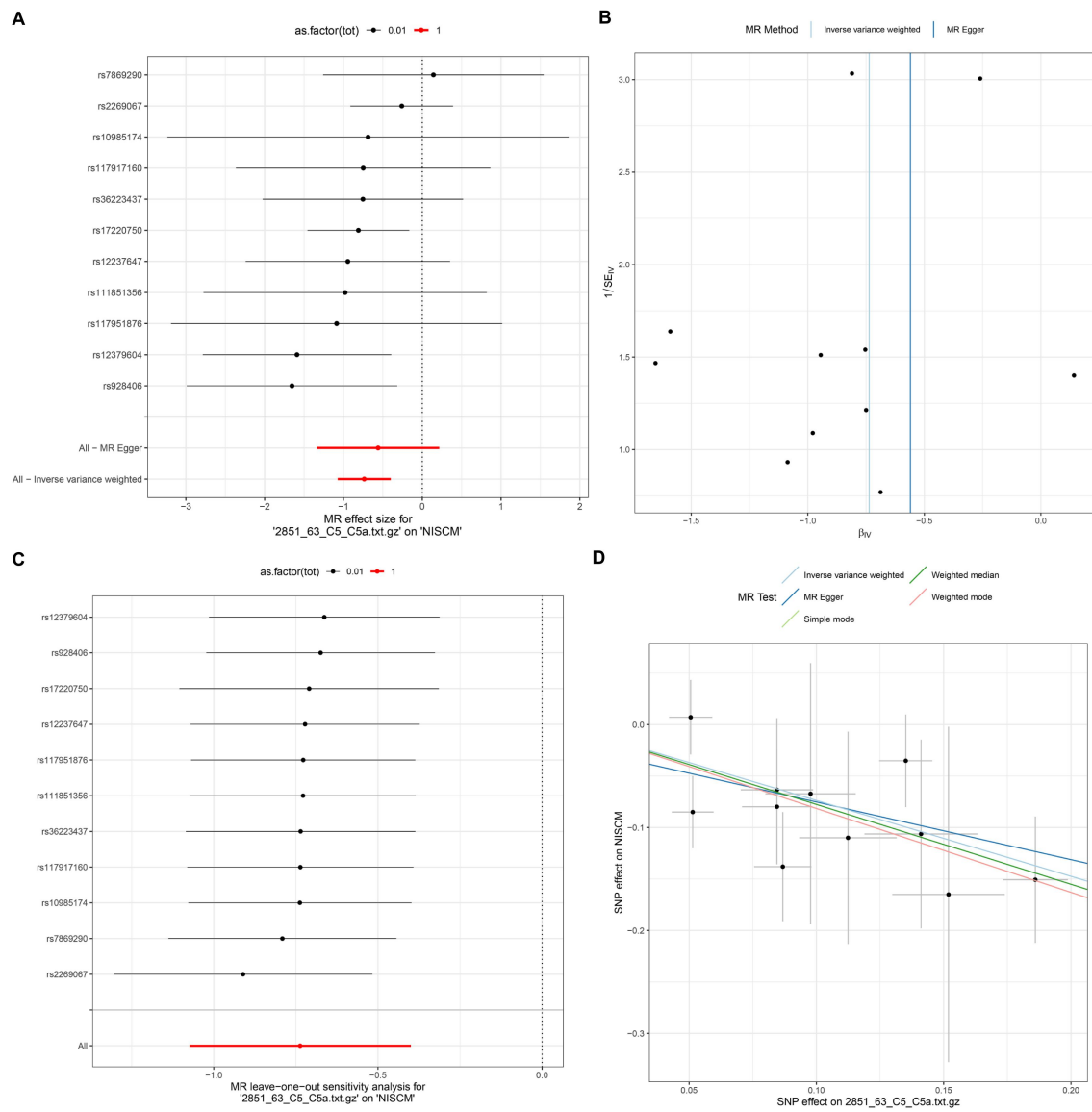
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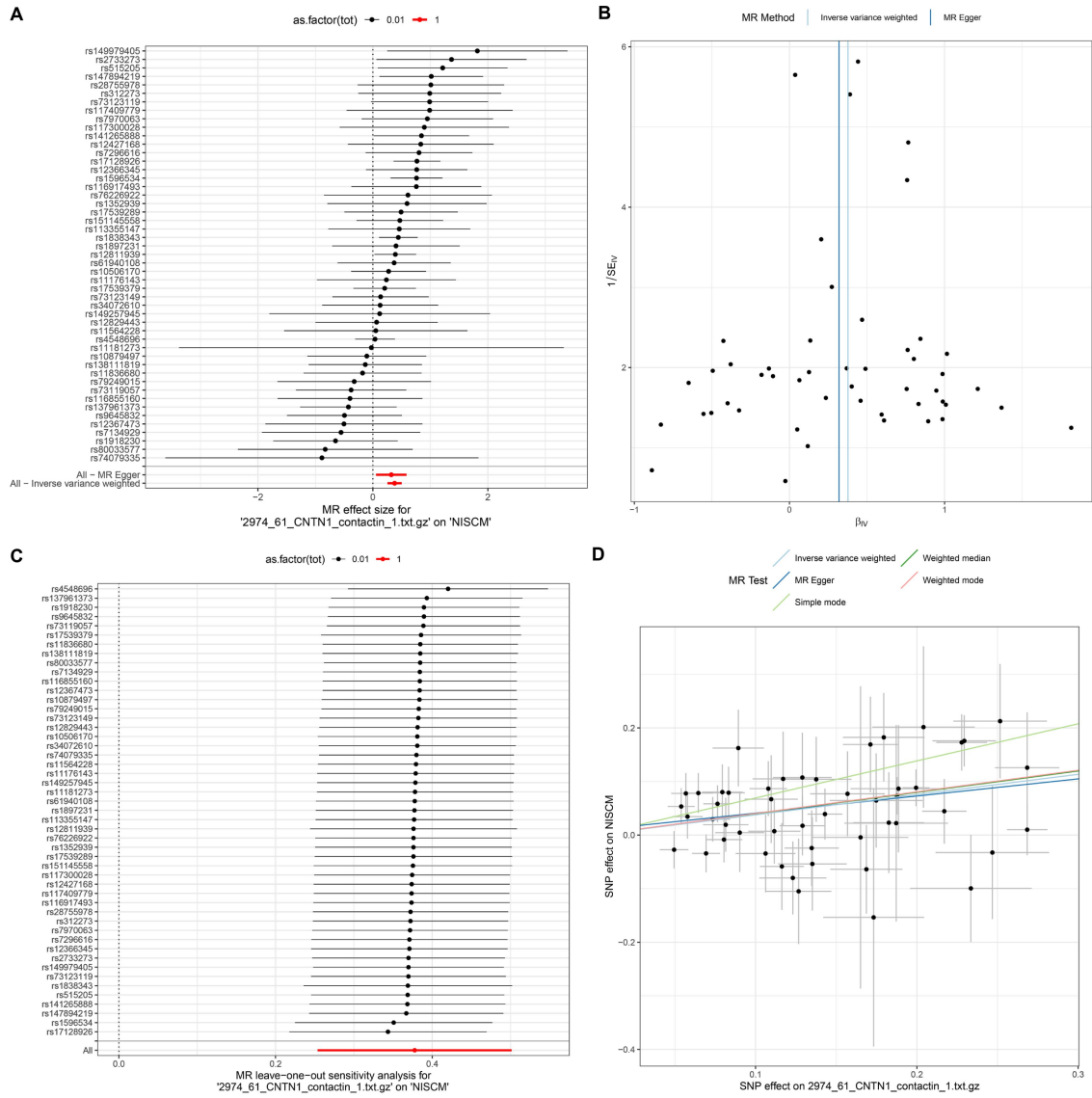
ASPN



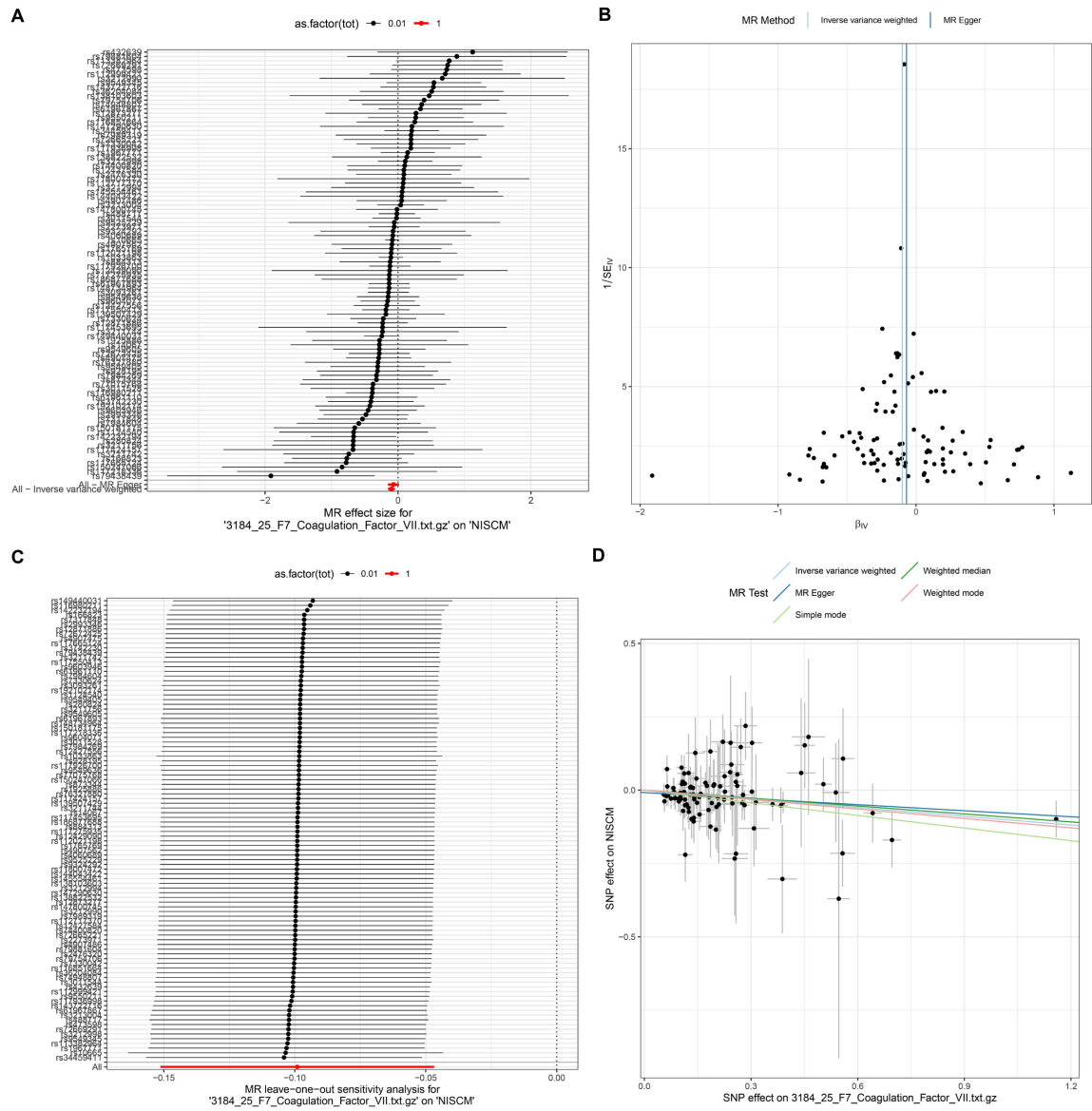
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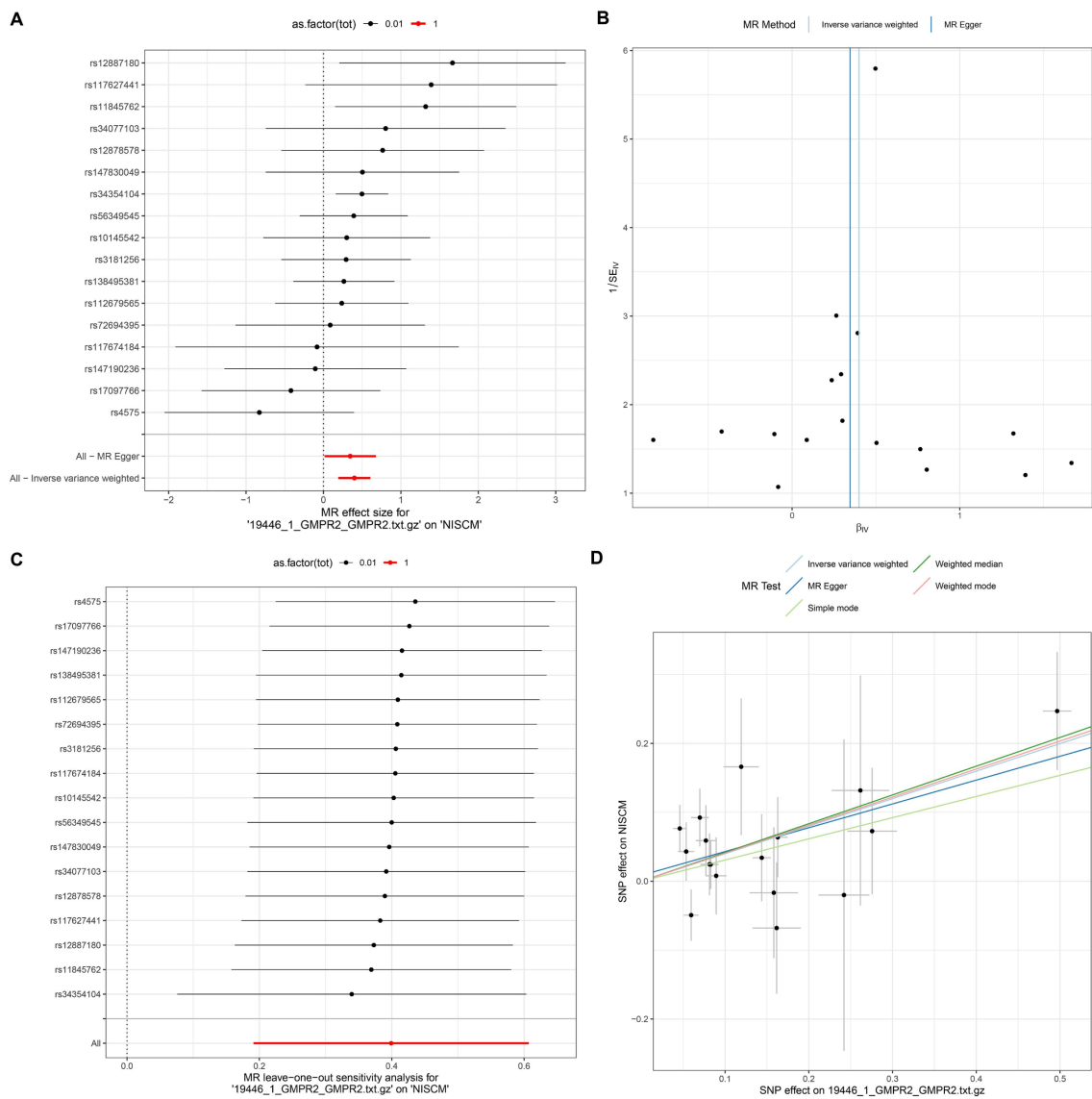
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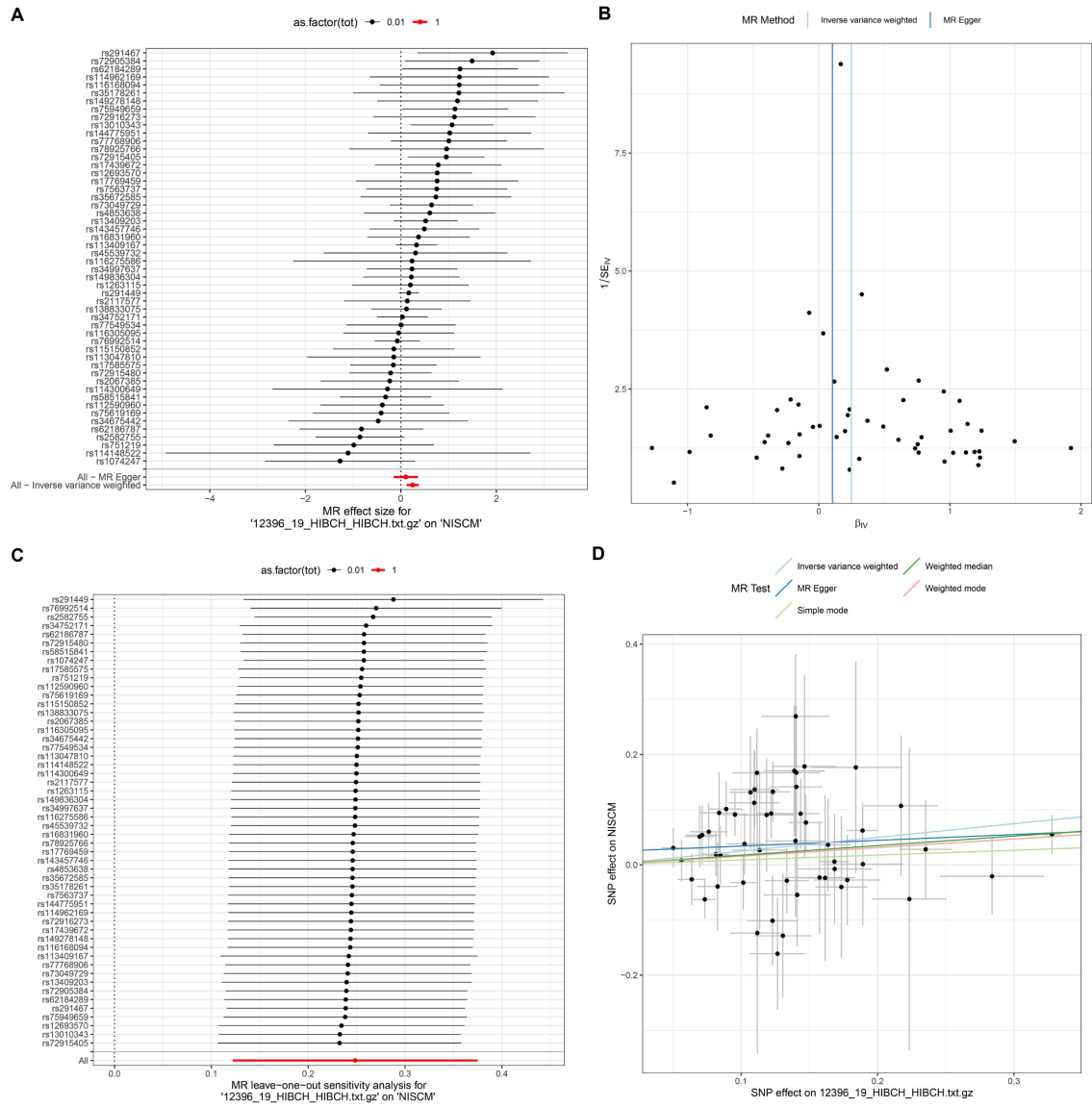
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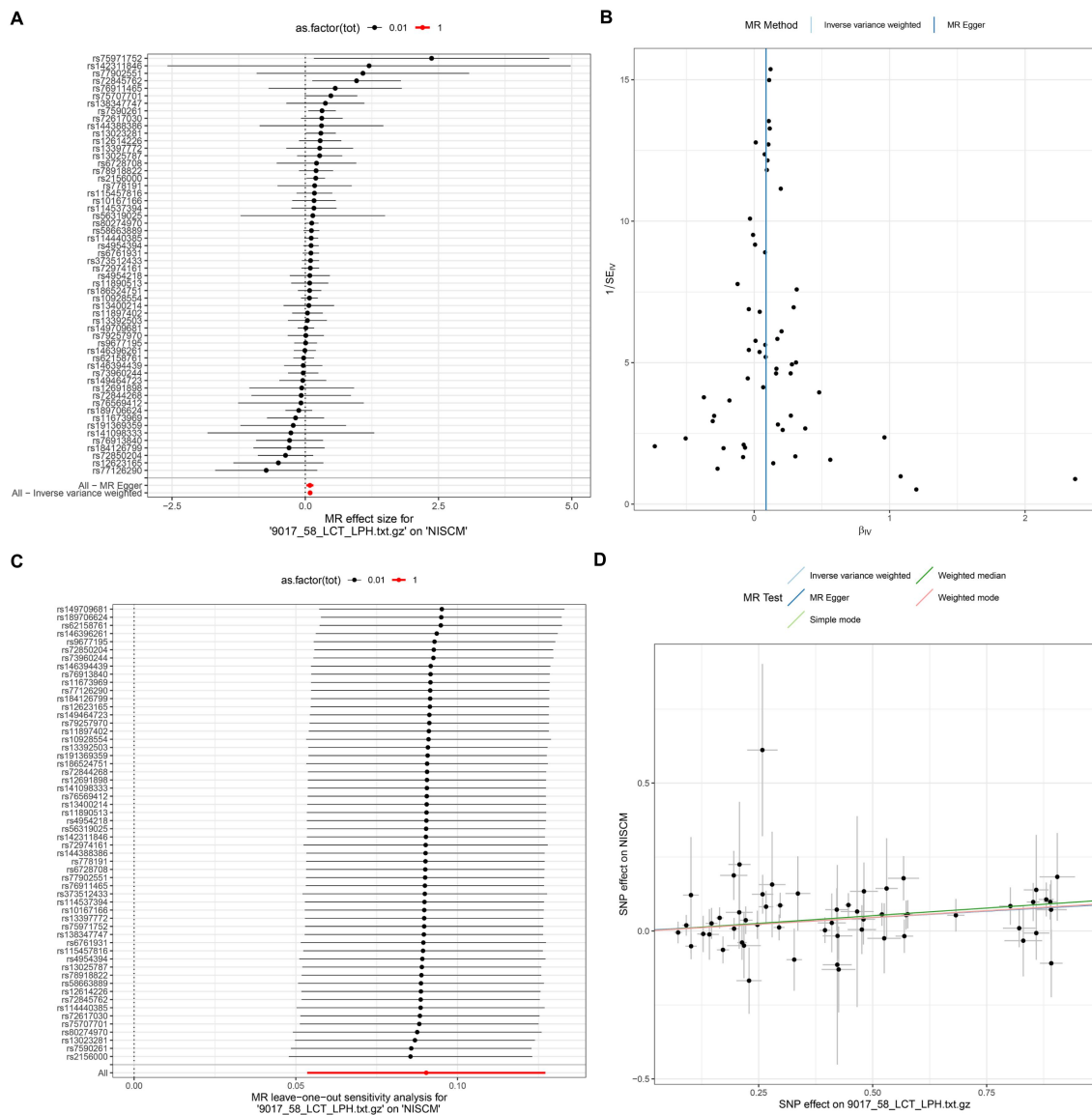
F7-Coagulation_Factor VII



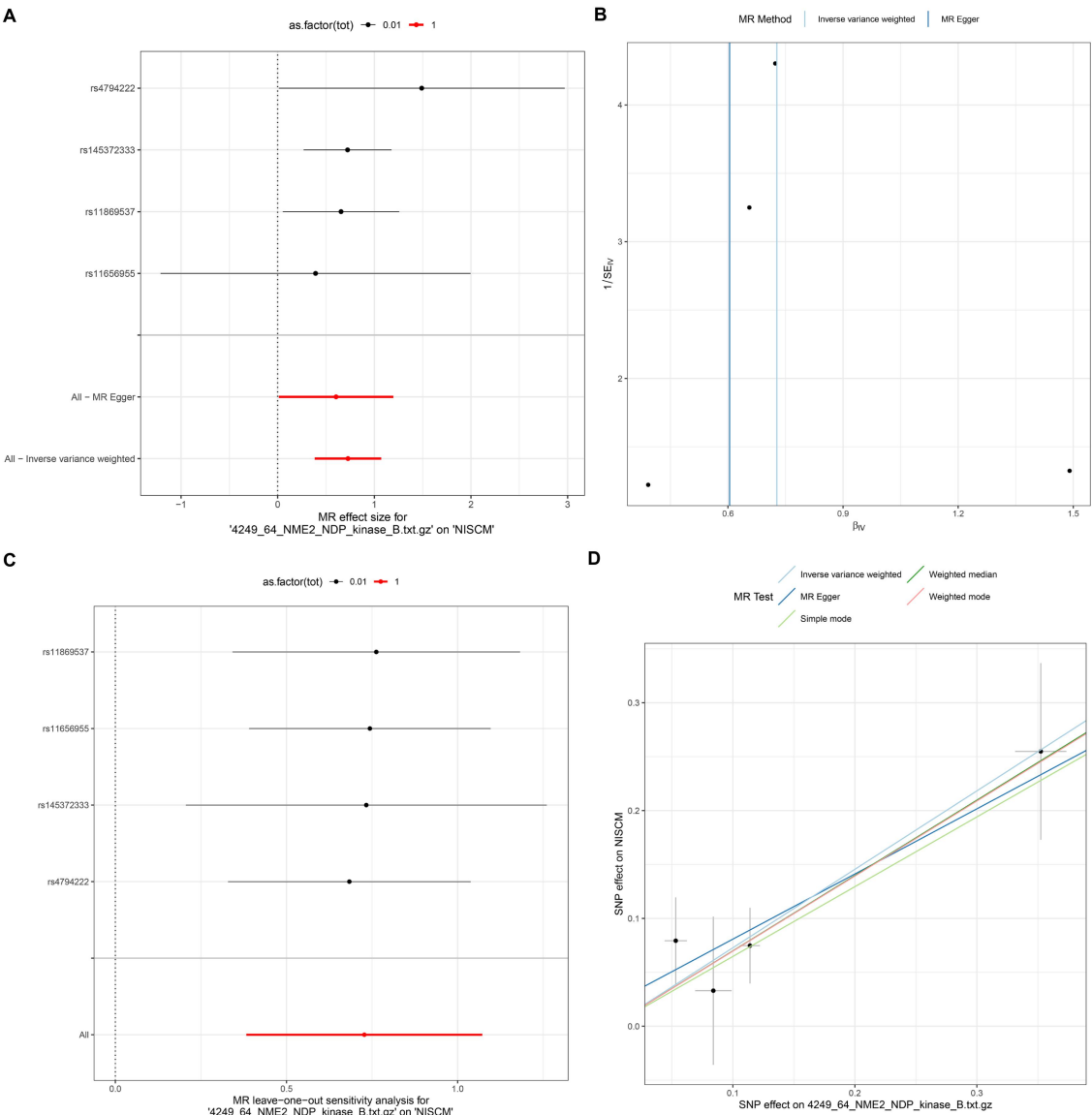
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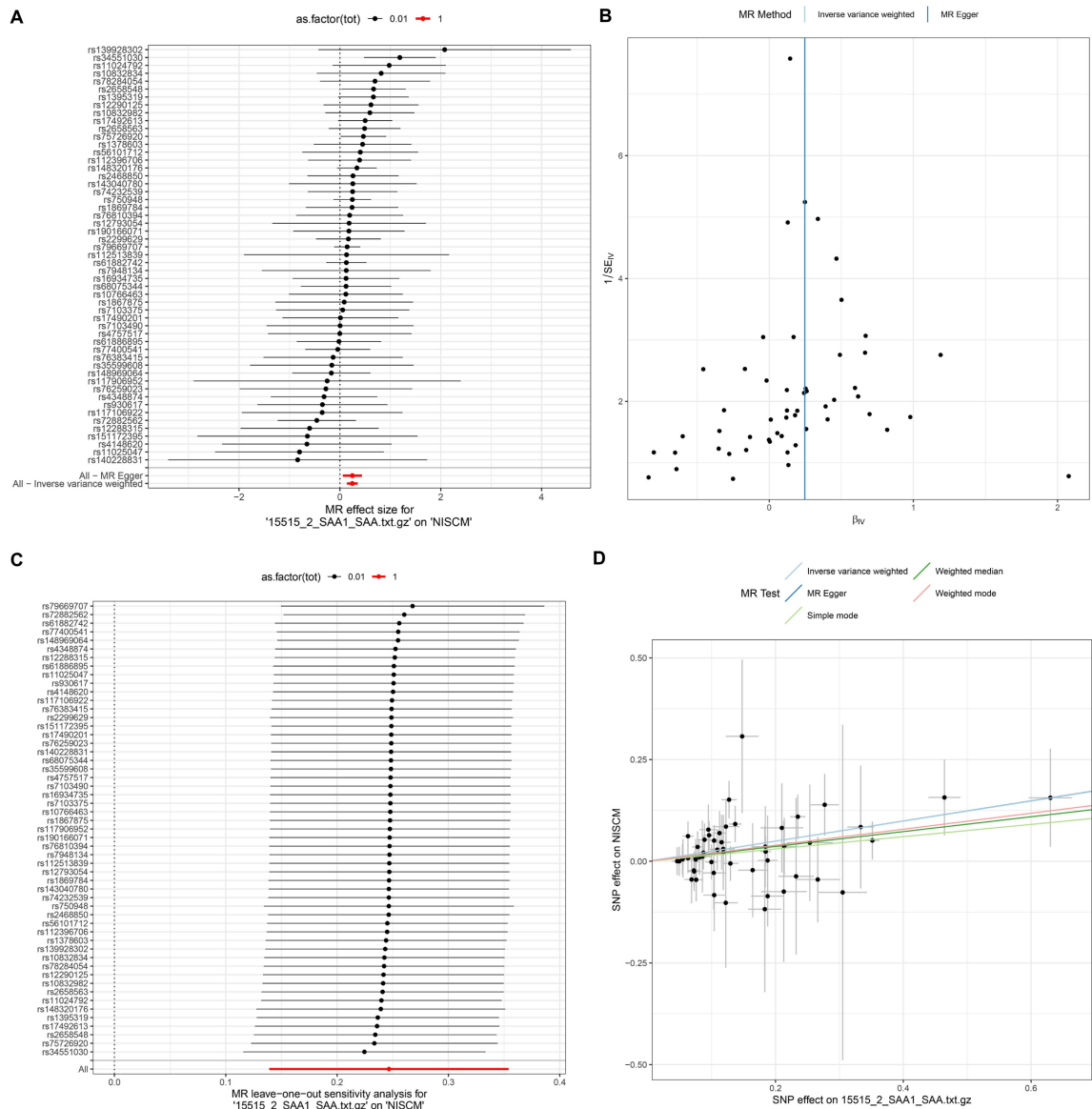
HIBCH



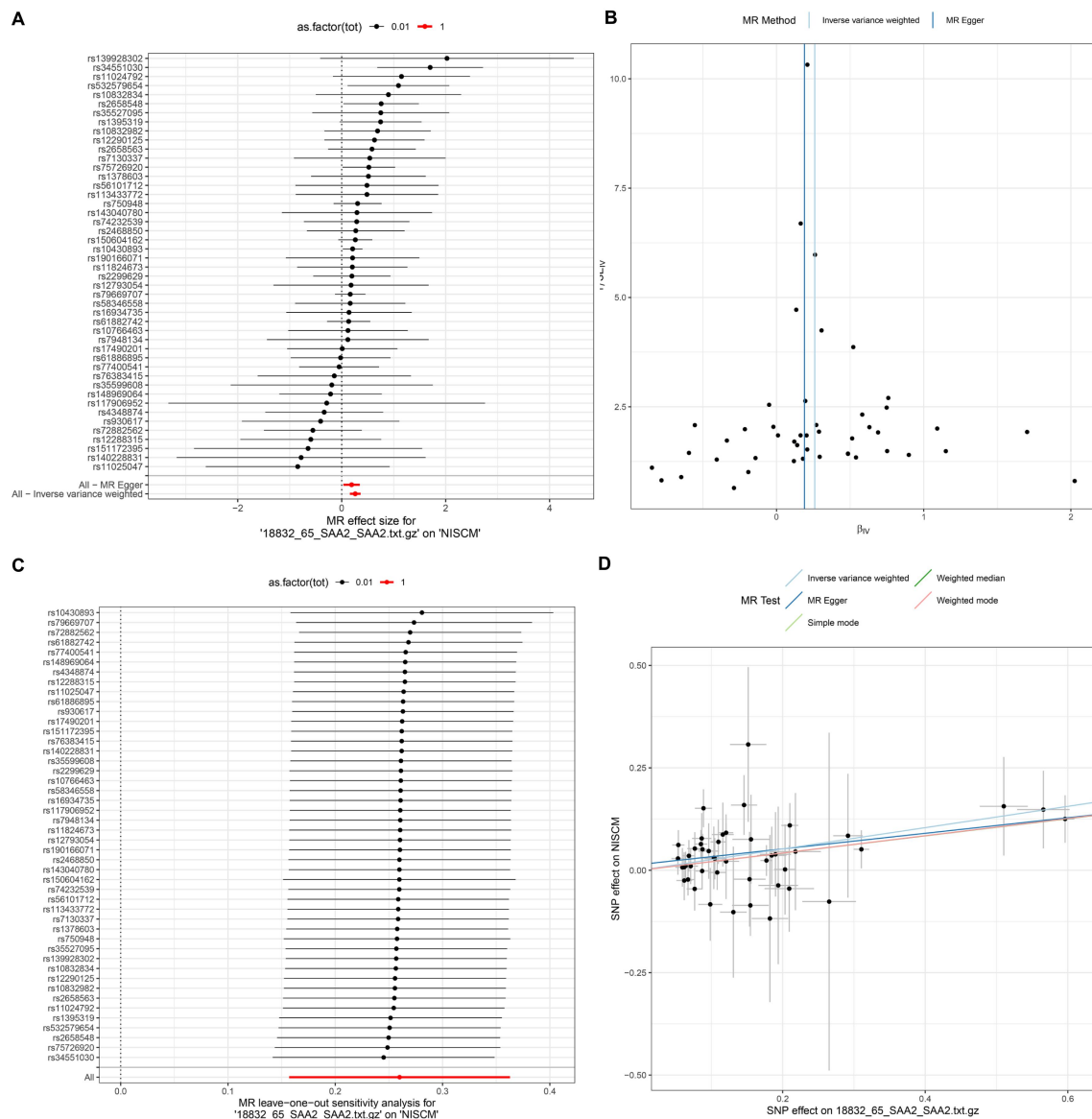
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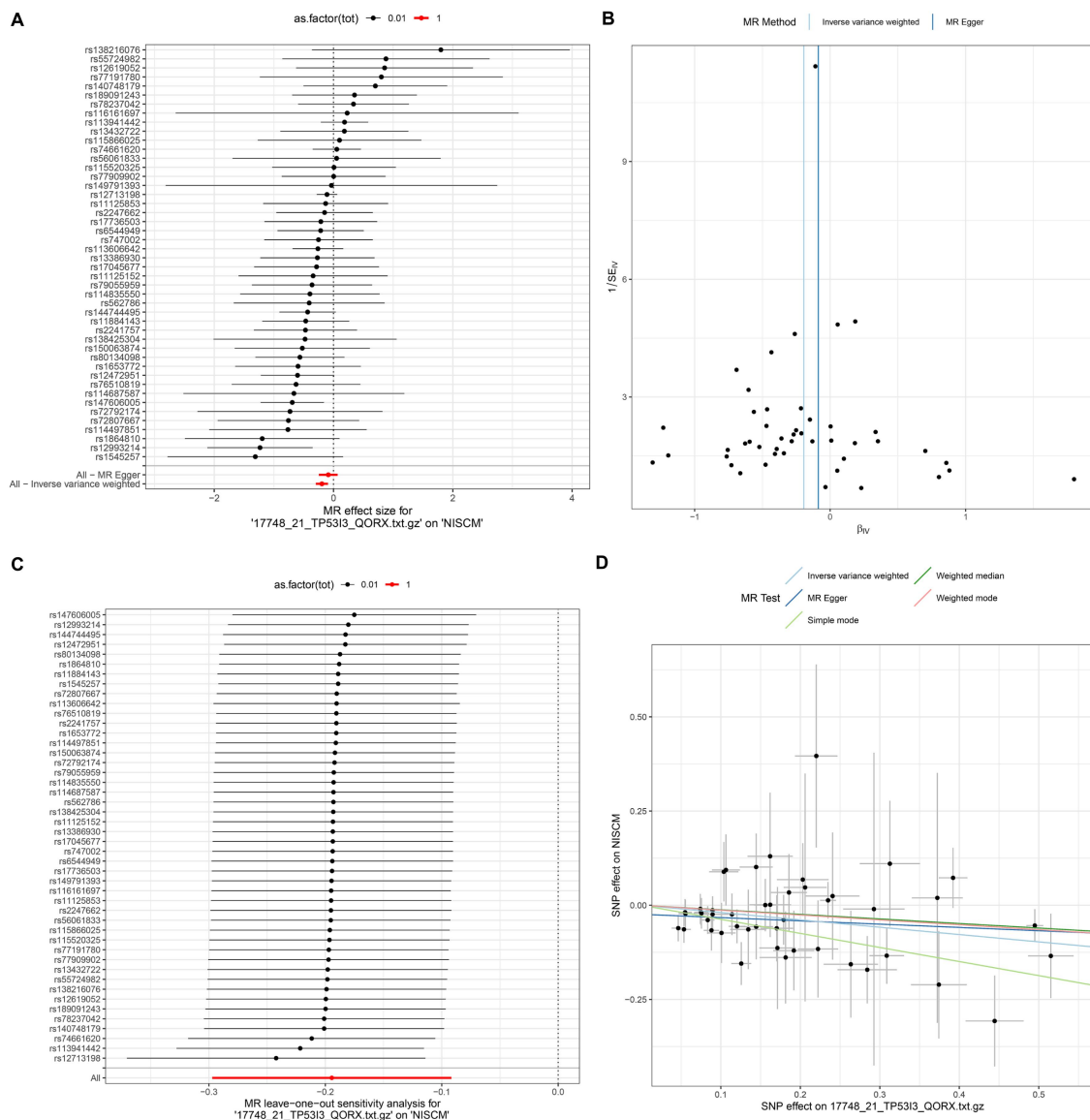
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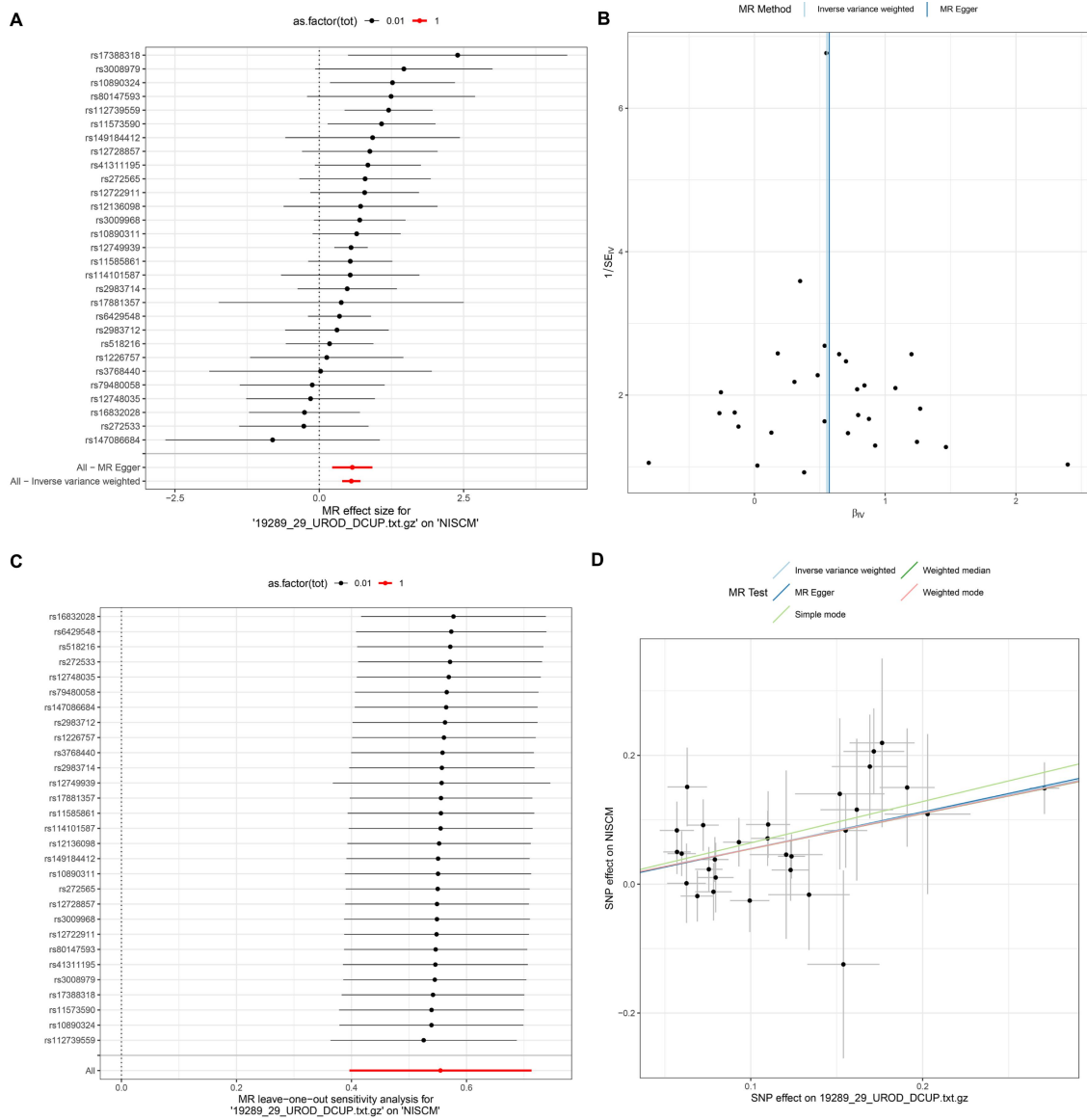
SAA1



SAA2



TP53l3



UROD