

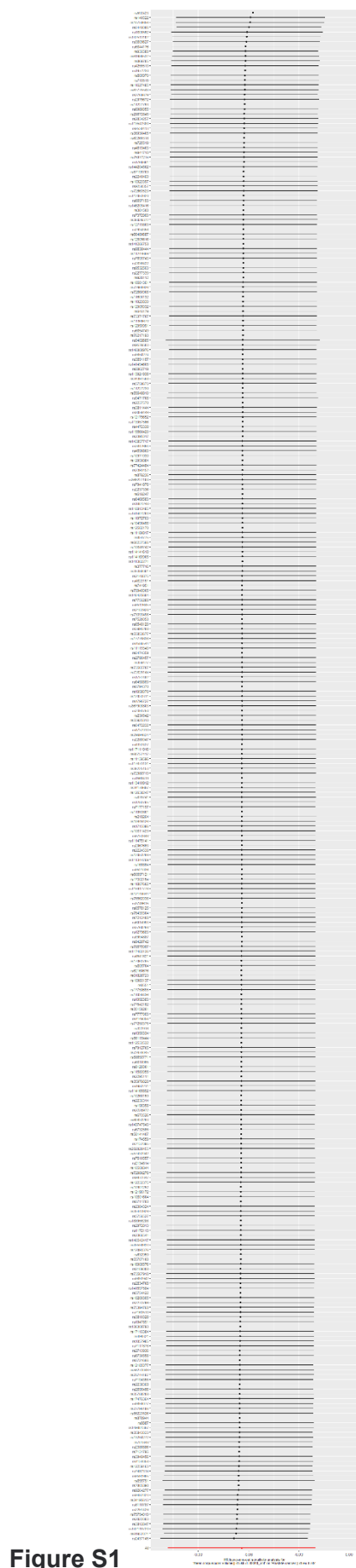


Figure S1

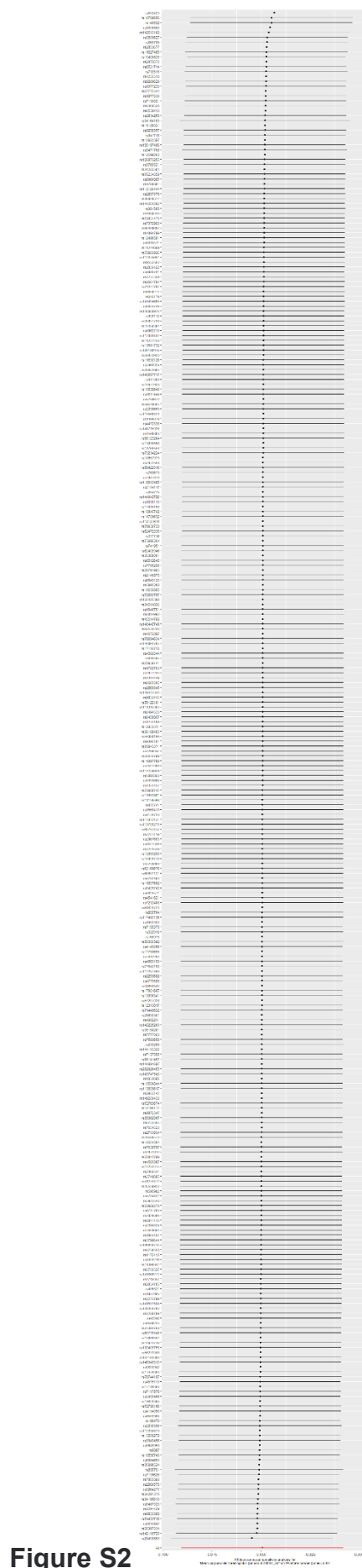


Figure S2

Figure S3

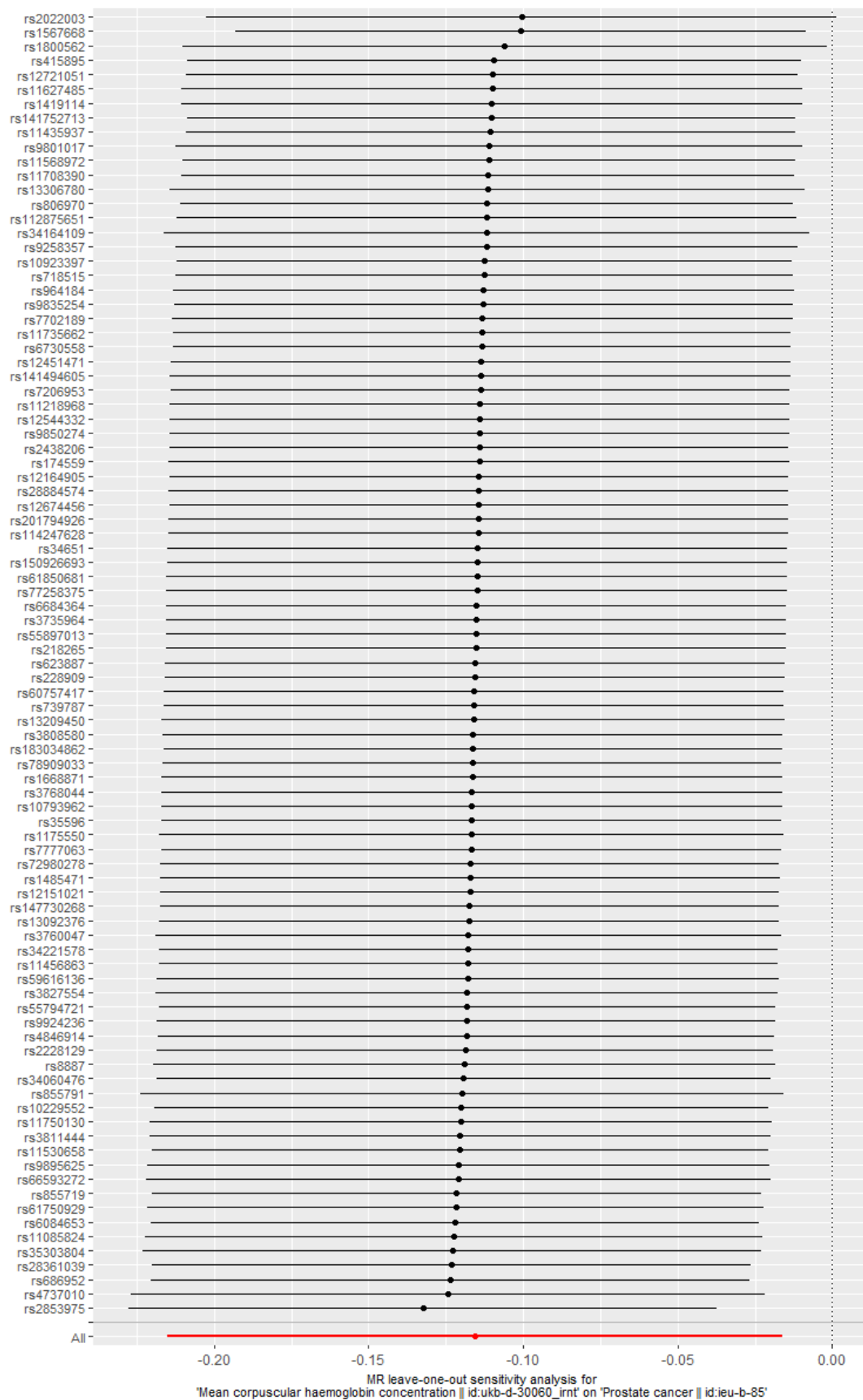


Figure S4

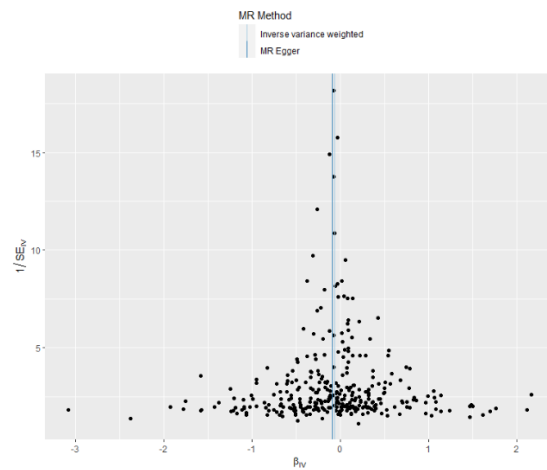


Figure S5

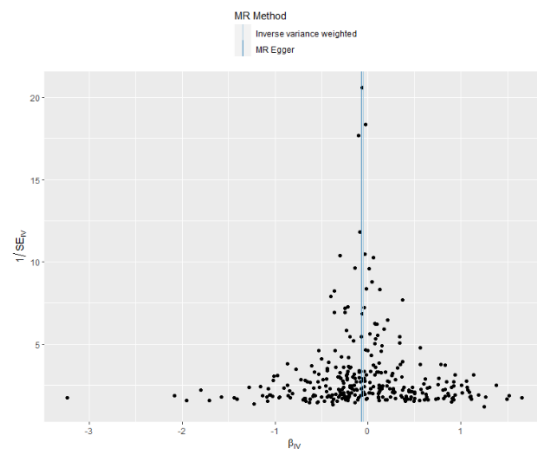
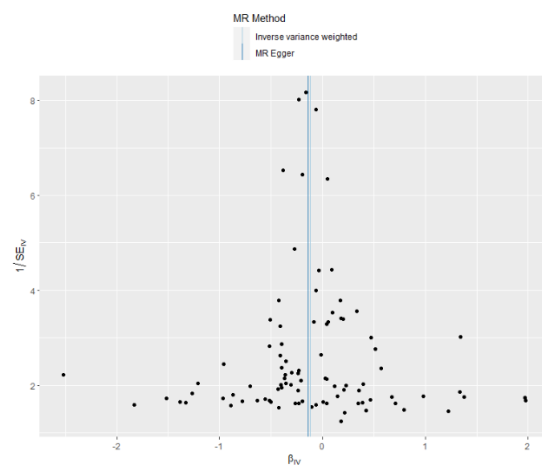


Figure S6



Supporting information Figure S1. “Leave - one - out” analysis to investigate whether the causal association between NCV and prostate cancer was driven by a unique single nucleotide polymorphism. MR, Mendelian randomization.

Supporting information Figure S2. “Leave - one - out” analysis to investigate

whether the causal association between MCH and prostate cancer was driven by a unique single nucleotide polymorphism. MR, Mendelian randomization.

Supporting information Figure S3. “Leave - one - out” analysis to investigate whether the causal association between MCHC and prostate cancer was driven by a unique single nucleotide polymorphism. MR, Mendelian randomization.

Supporting information Figure S4. Funnel plot from genetically predicted MCV on PCa.

Supporting information Figure S5. Funnel plot from genetically predicted MCH on PCa.

Supporting information Figure S6. Funnel plot from genetically predicted MCHC on PCa.