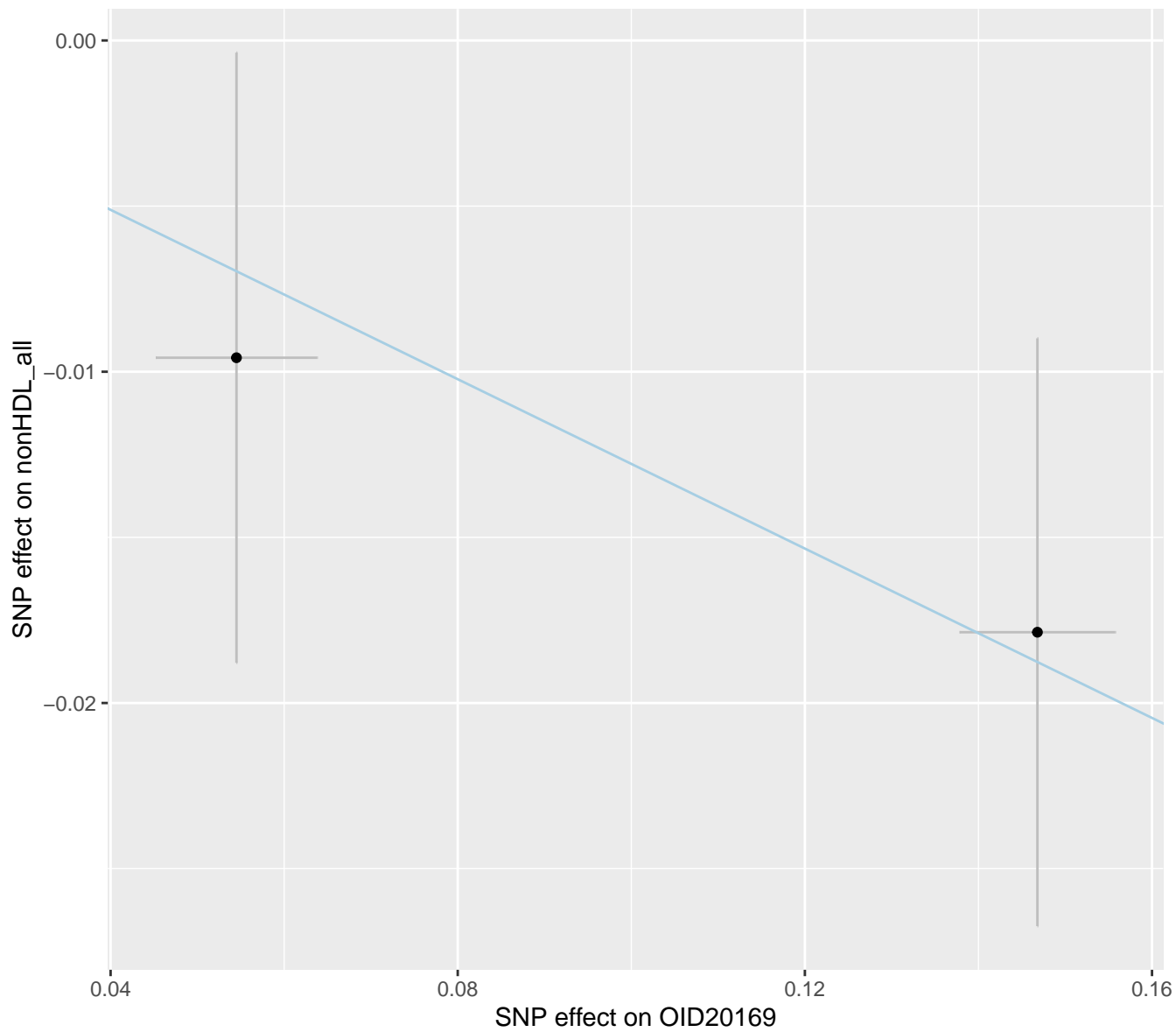


# MR Estimate

Inverse variance weighted



Insufficient number of SNPs