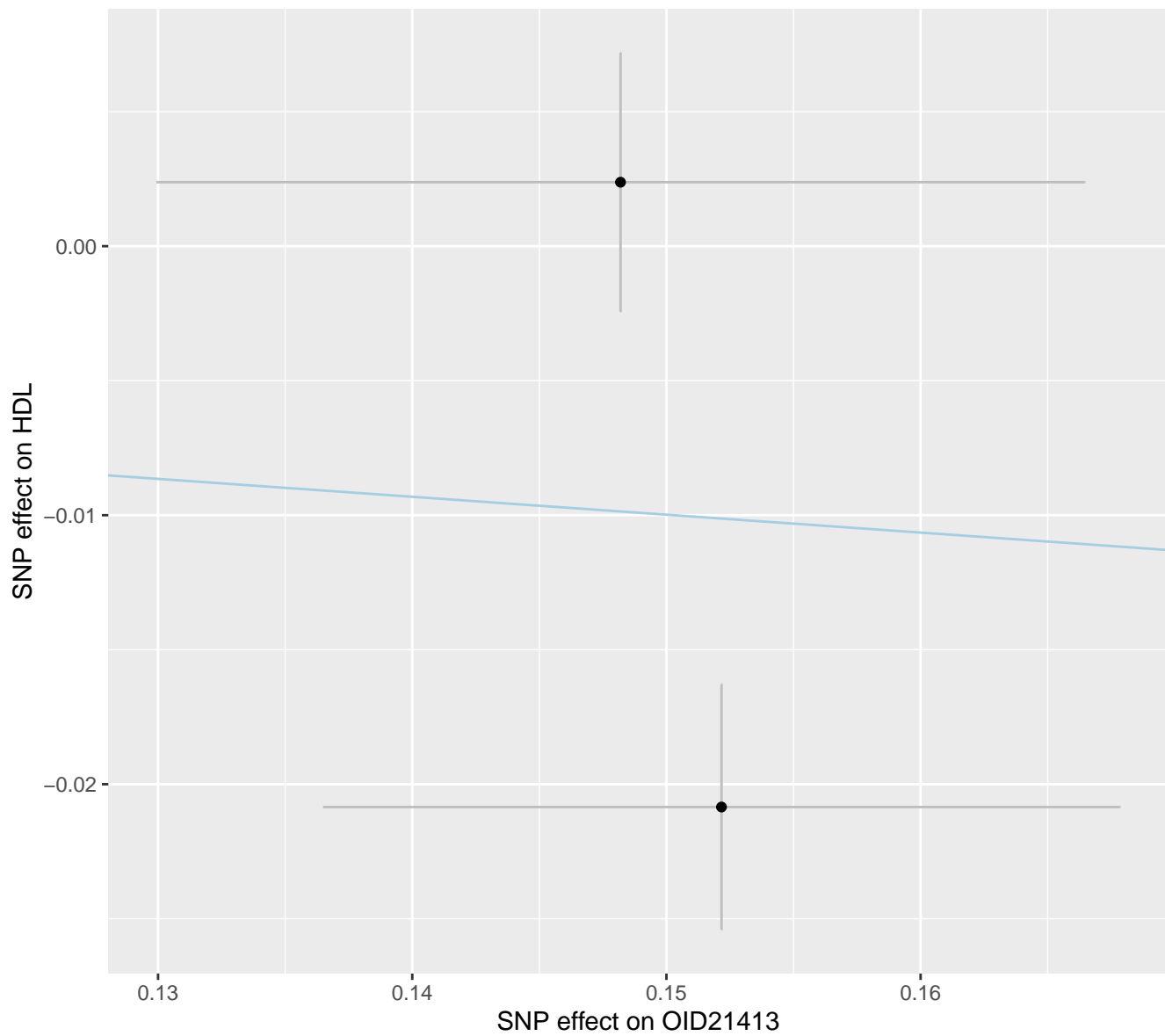


# MR Estimate

 Inverse variance weighted



Insufficient number of SNPs