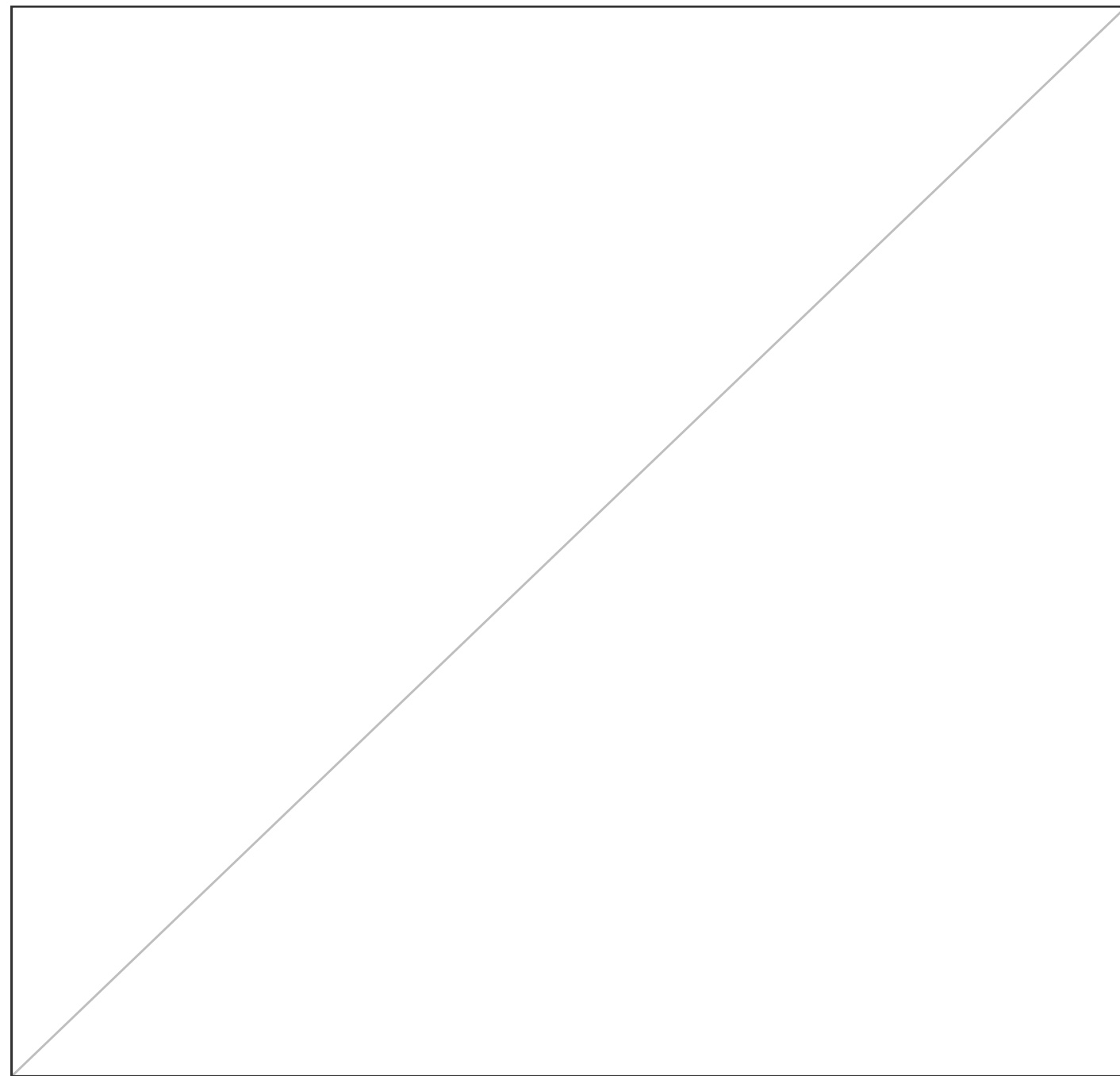


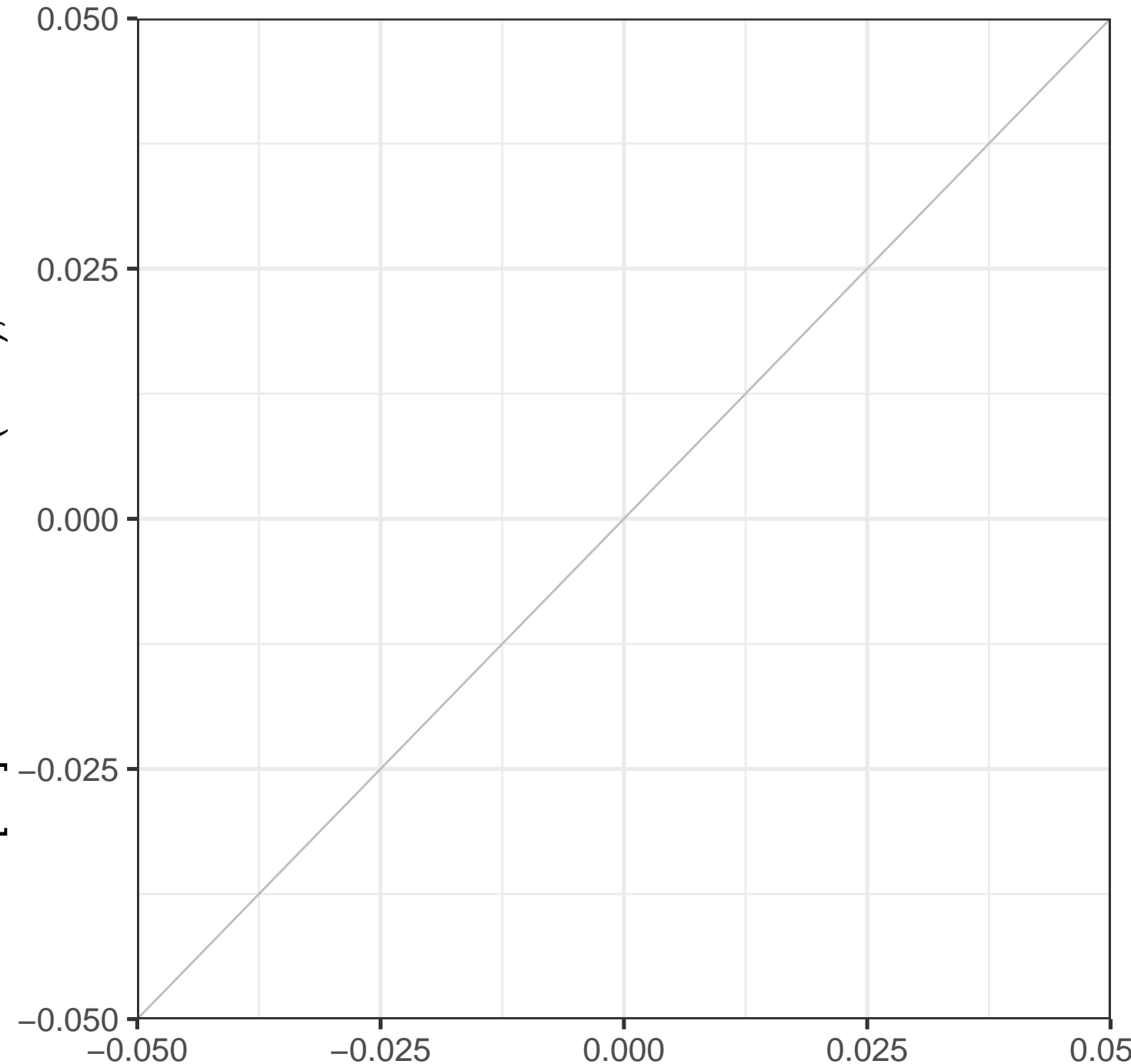
Comparison of PRS–PheWAS associations (B1, eur_leave_23andme, clump p1: 1e−3)

−log₁₀(P) of scale(PRS), without chr3 region



−log₁₀(P) of scale(PRS), with chr3 region

BETA [SE] estimate of scale(PRS), without chr3



BETA [SE] estimate of scale(PRS), with chr3