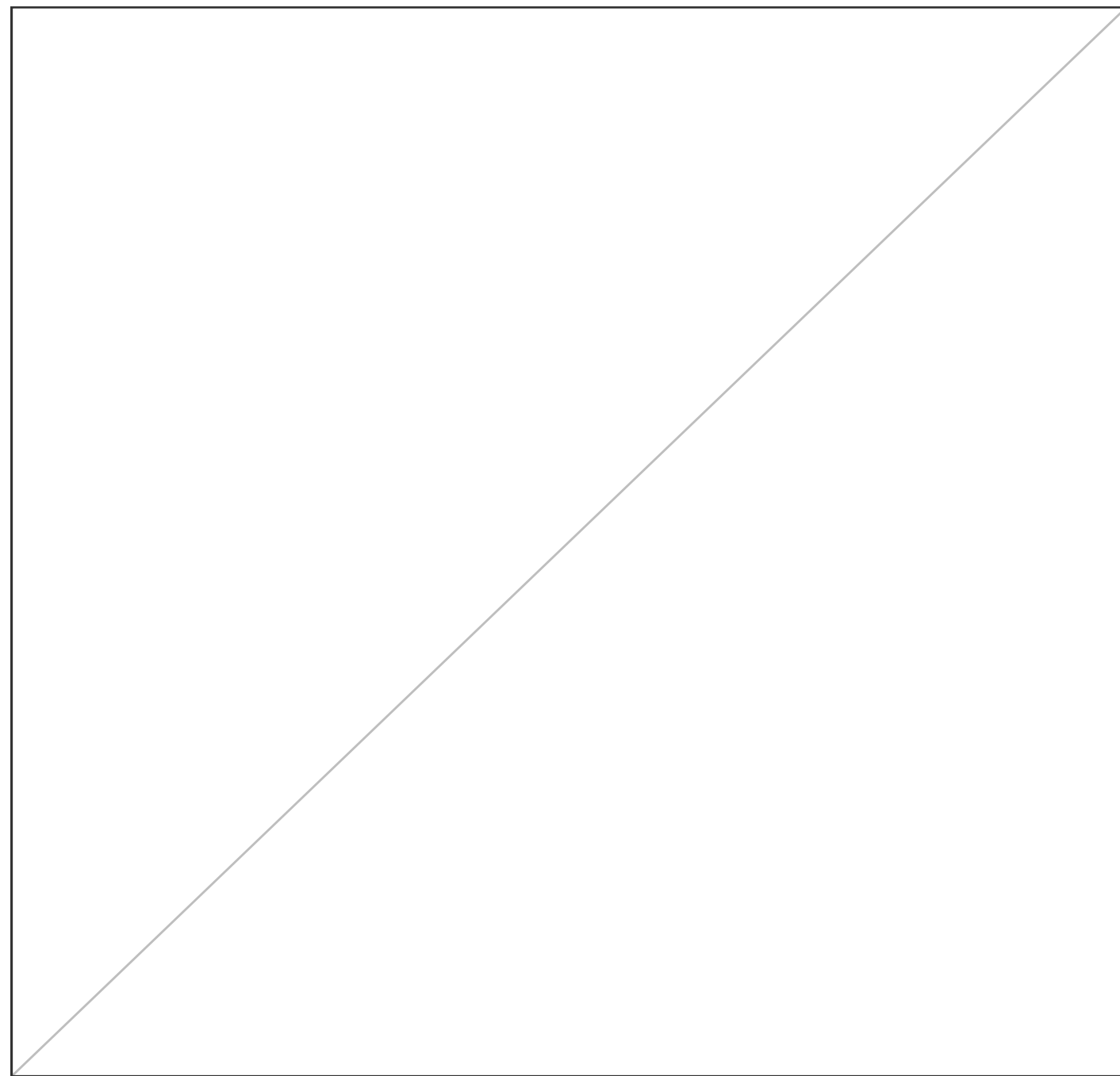


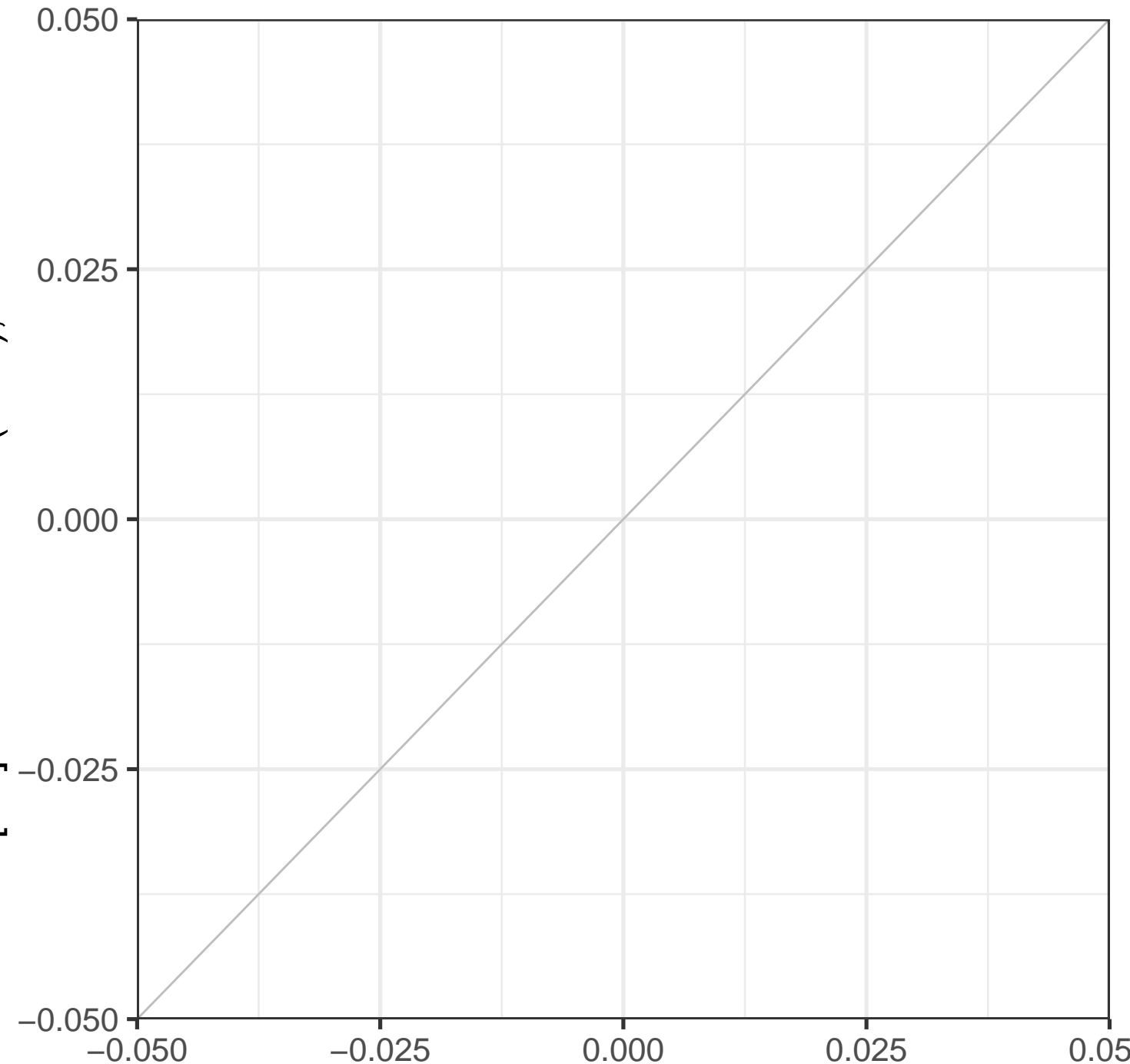
Comparison of PRS–PheWAS associations (B1, eur_leave_23andme, clump p1: .001)

$-\log_{10}(P)$ of scale(PRS), without chr3 region



$-\log_{10}(P)$ of scale(PRS), with chr3 region

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



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