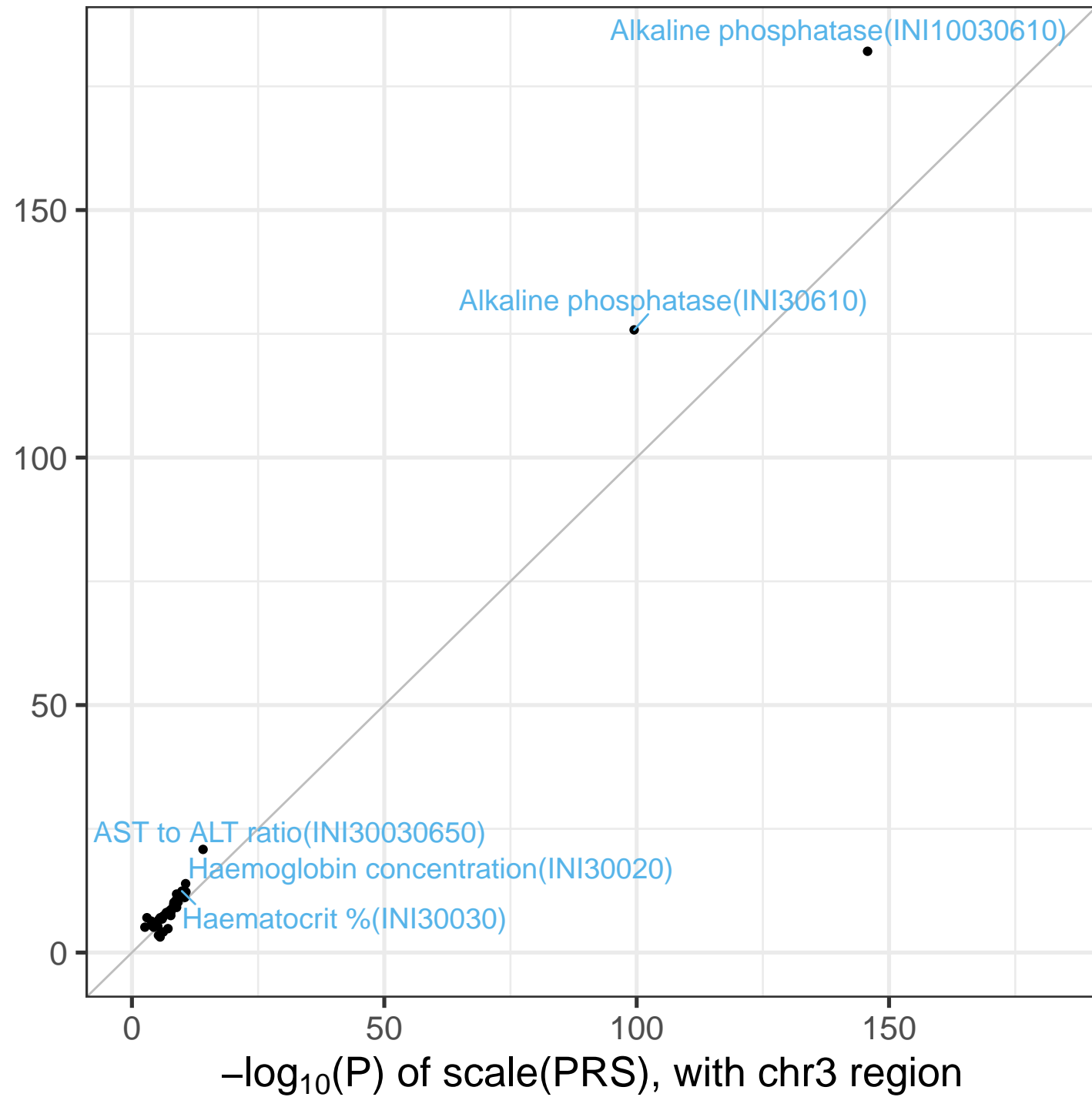


Comparison of PRS–PheWAS associations (C2, eur_leave_23andme, clump p1: 1e−4)

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



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

