Comparison of PRS-PheWAS associations (B1, eur_leave_23andme, clump p1: 1e-5) 0.12 -10.0 -BETA [SE] estimate of scale(PRS), without chr3 0.08 7.5 -Malabsorption/coeliac disease(HC308 0.04 -5.0 -2.5 -Monocyte count(INI30130) 0.00 -Malabsorption/coeliac disease(HC303) Monocyte %(INI30190) Monocyte %(INI30190). 0.0 -Monocyte count(INI30130) -0.040.0 2.5 5.0 7.5 10.0 0.04 0.12 0.00 0.08 -0.04-log₁₀(P) of scale(PRS), with chr3 region BETA [SE] estimate of scale(PRS), with chr3

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