Comparison of PRS-PheWAS associations (B1, eur_leave_ukbb_23andme, clump p1: 1e-5) 0.00 Monocyte count(INI30130) ♦ BETA [SE] estimate of scale(PRS), without chr3 -0.01Monocyte %(INI30190) -0.02Monocyte %(INI30190)• -0.03 **-**Monocyte count(INI30130) -0.03-0.02 -0.010.00

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

0 -

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

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