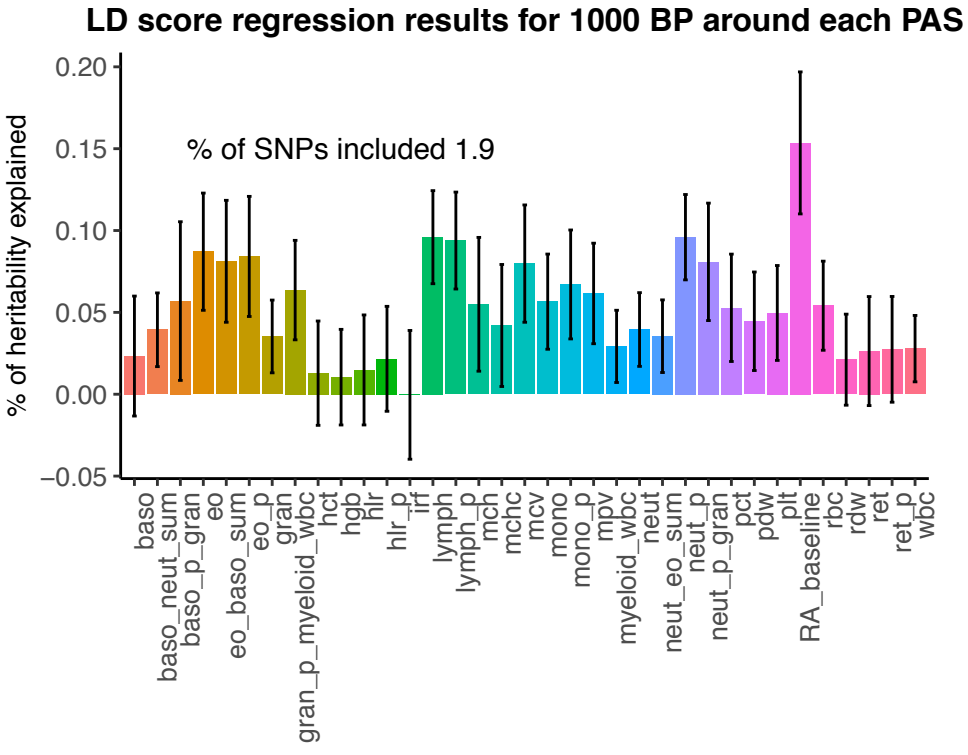
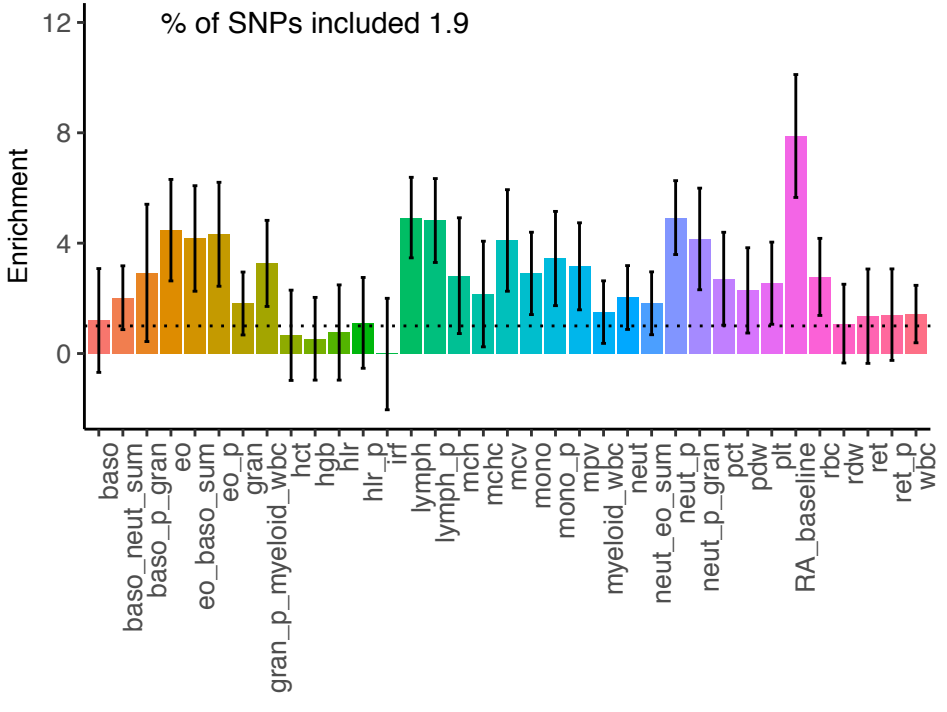
A


LD score regression results for 1000 BP around each PAS
% of SNPs included 1.9
% of heritability explained
0.20
0.15
0.10
0.05
0.00
−0.05
baso
baso_neut_sum
baso_p_gran
eo
eo_baso_sum
eo_p
gran
gran_p_myeloid_wbc
hct
hgb
hlr
hlr_p
irf
lymph
lymph_p
mch
mchc
mcv
mono
mono_p
mpv
myeloid_wbc
neut
neut_eo_sum
neut_p
neut_p_gran
pct
pdw
plt
RA_baseline
rbc
rdw
ret
ret_p
wbc

B


LD score regression results for 1000 BP around each PAS
% of SNPs included 1.9
Enrichment
12
8
4
0
baso
baso_neut_sum
baso_p_gran
eo
eo_baso_sum
eo_p
gran
gran_p_myeloid_wbc
hct
hgb
hlr
hlr_p
irf
lymph
lymph_p
mch
mchc
mcv
mono
mono_p
mpv
myeloid_wbc
neut
neut_eo_sum
neut_p
neut_p_gran
pct
pdw
plt
RA_baseline
rbc
rdw
ret
ret_p
wbc