

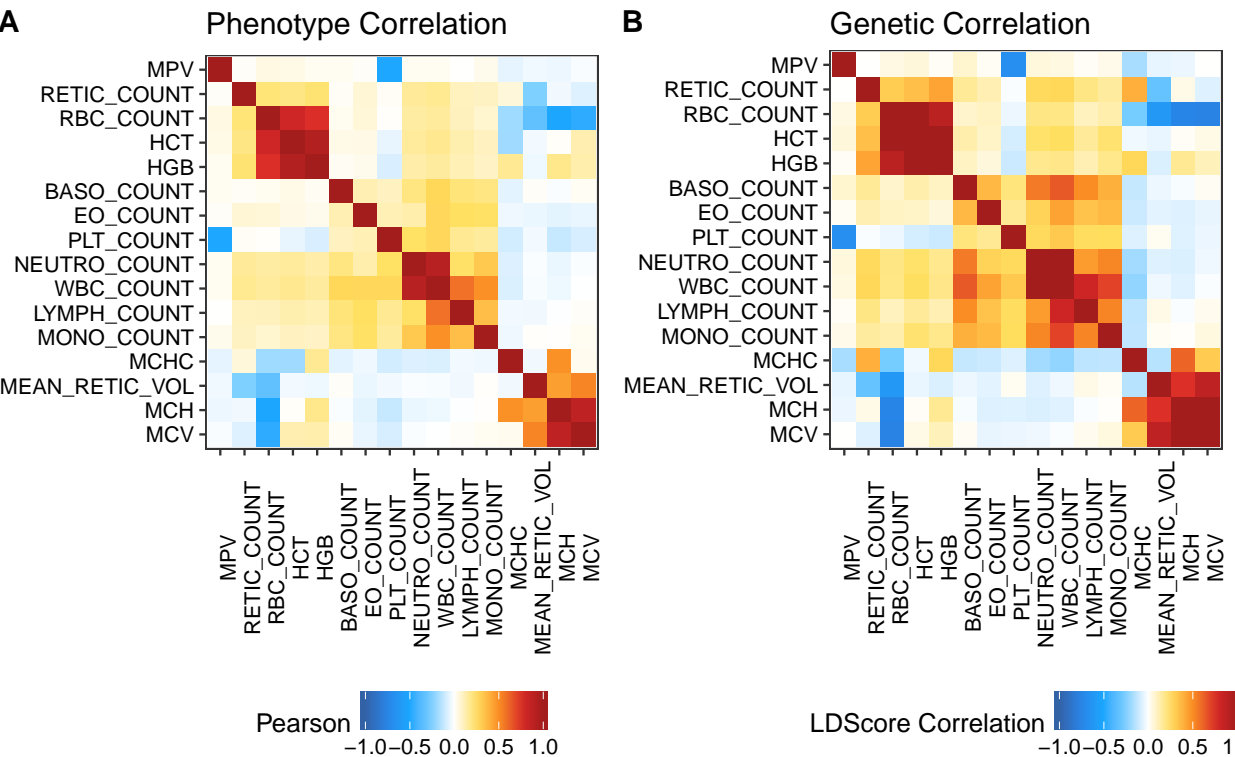
Interrogation of human hematopoietic traits at single-cell and single-variant resolution

(Caleb, Erik, Jacob), Will?, Hilary, Joel, (Martin, Jason, Vijay)

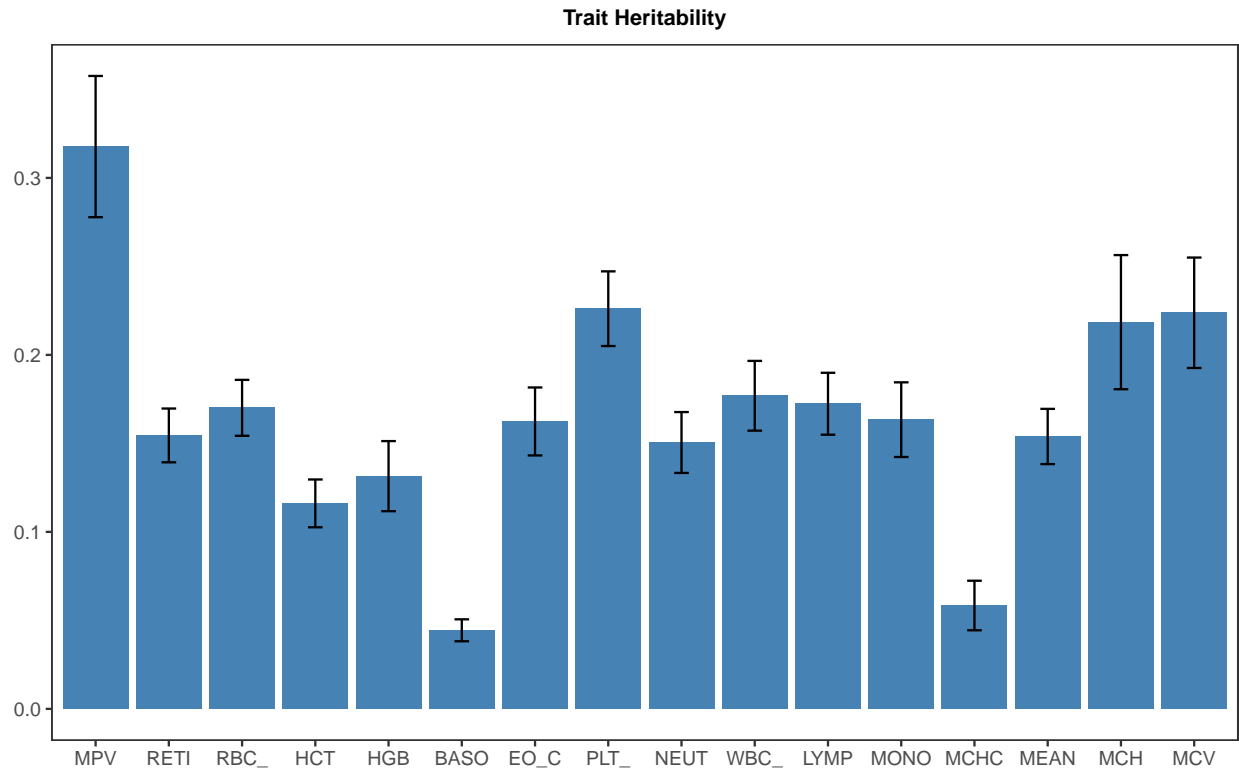
Supplemental Information

Overview of UK Biobank data

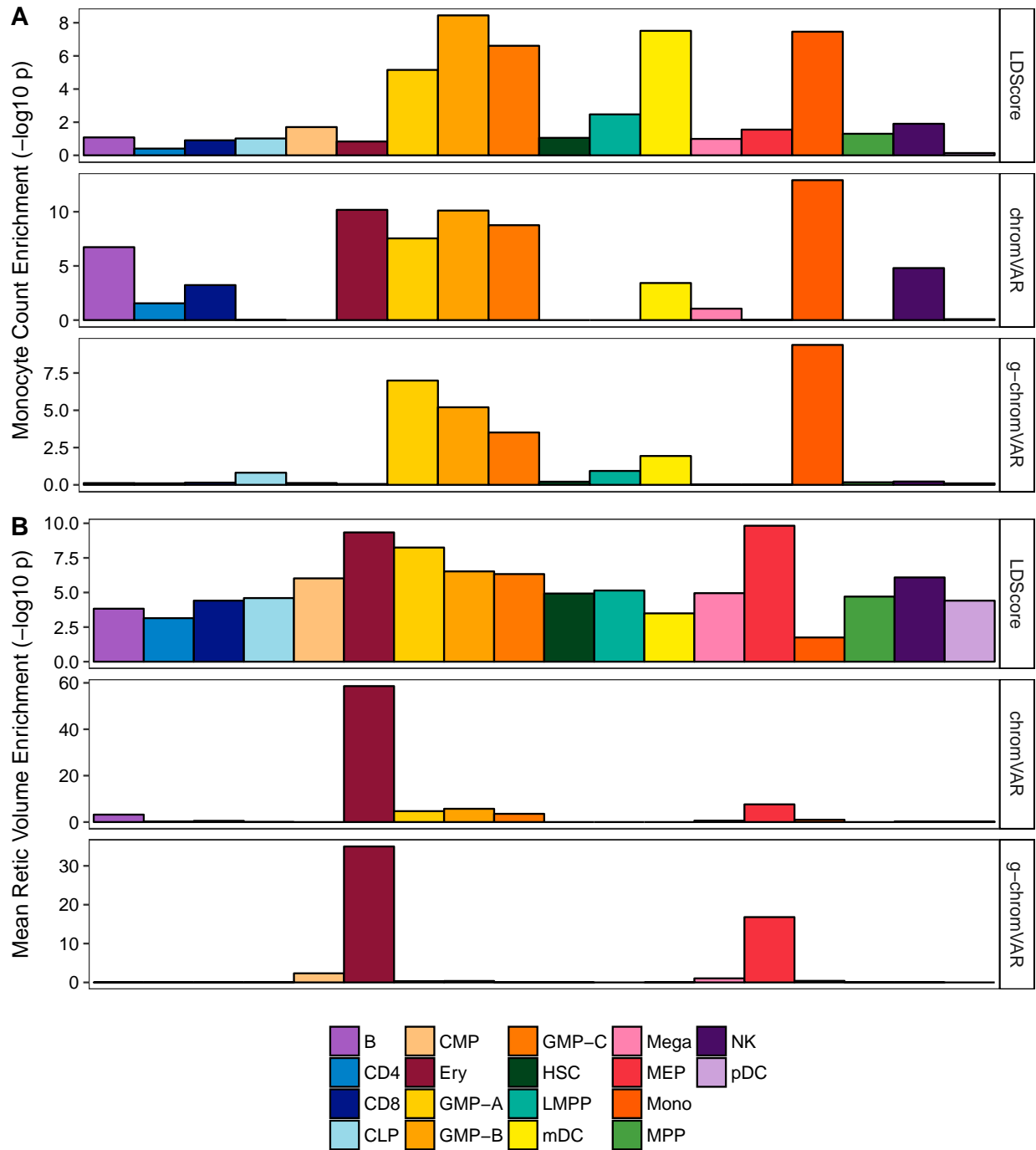
(A) a



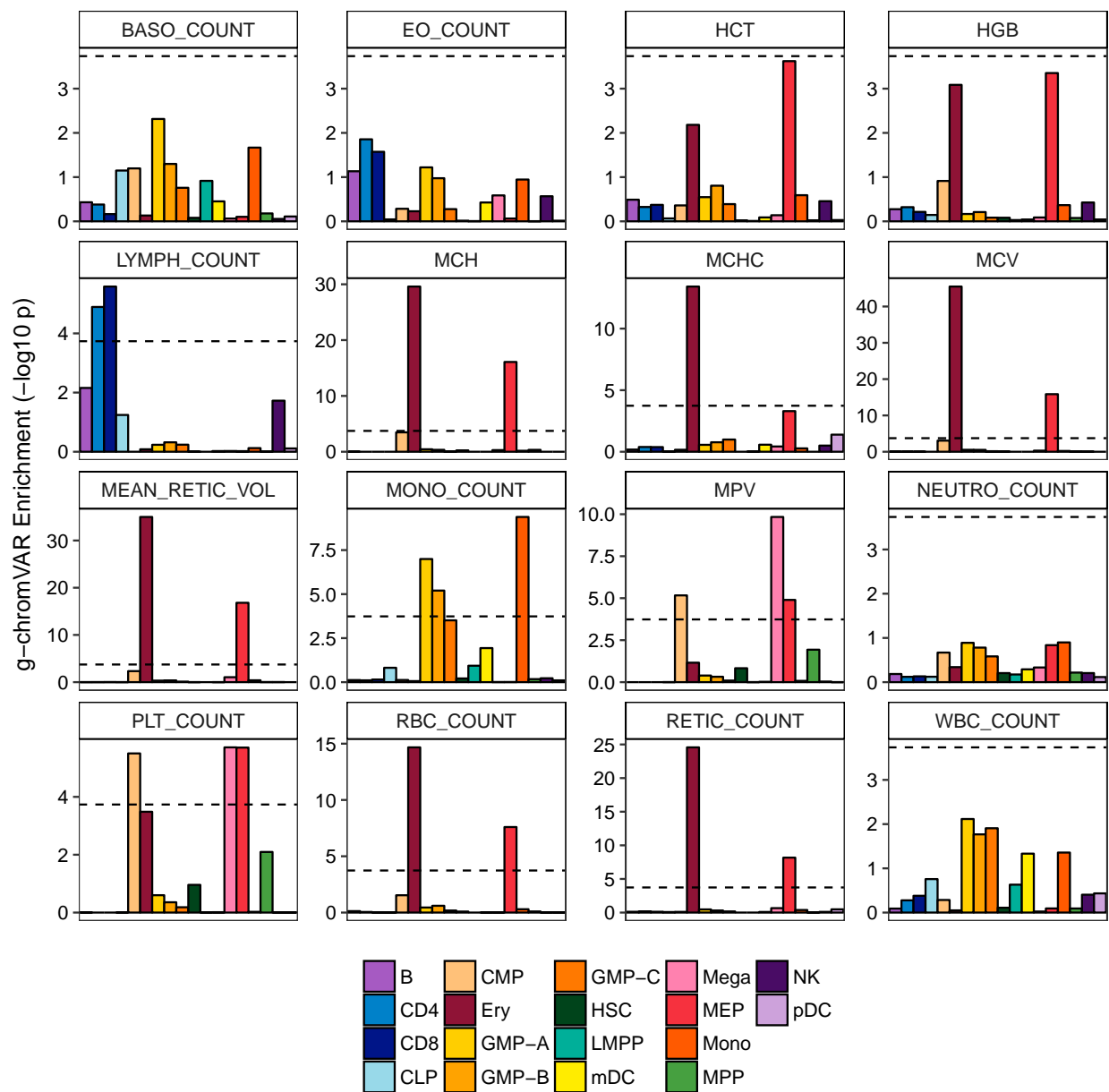
Supplemental Figure 1: Phenotypic and genetic correlations across the 16 traits examined.



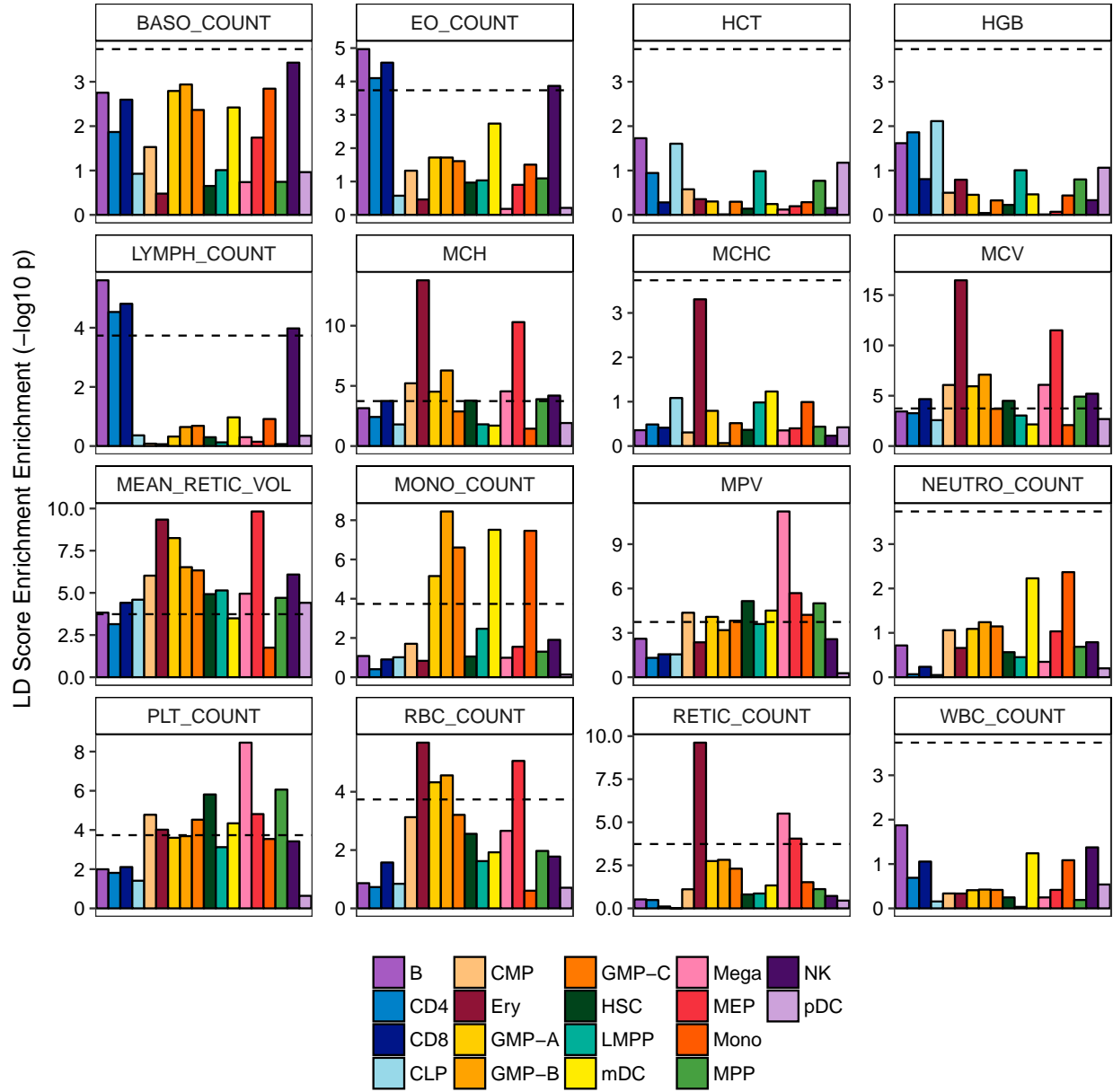
Supplemental Figure 2: Heritability estimates from LD Score Regression across 16 hematopoietic traits. The estimates of the narrow-sense SNP heritabilities are plotted with their corresponding standard errors.



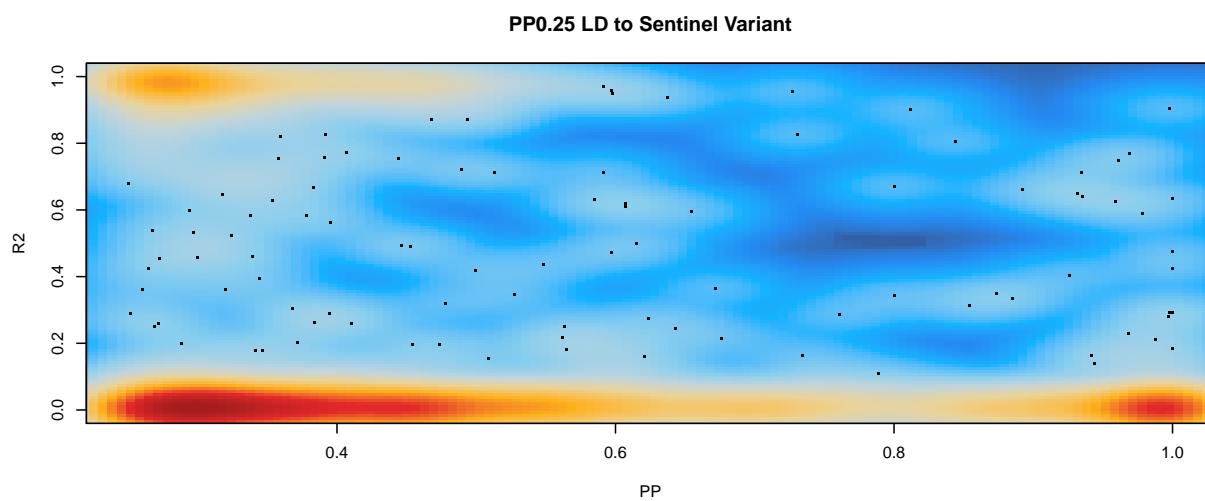
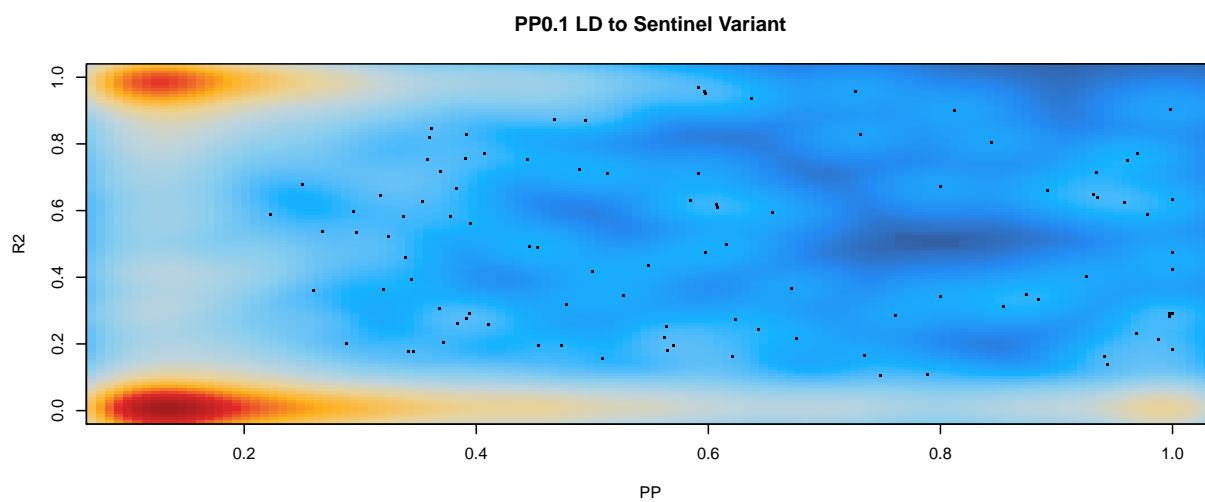
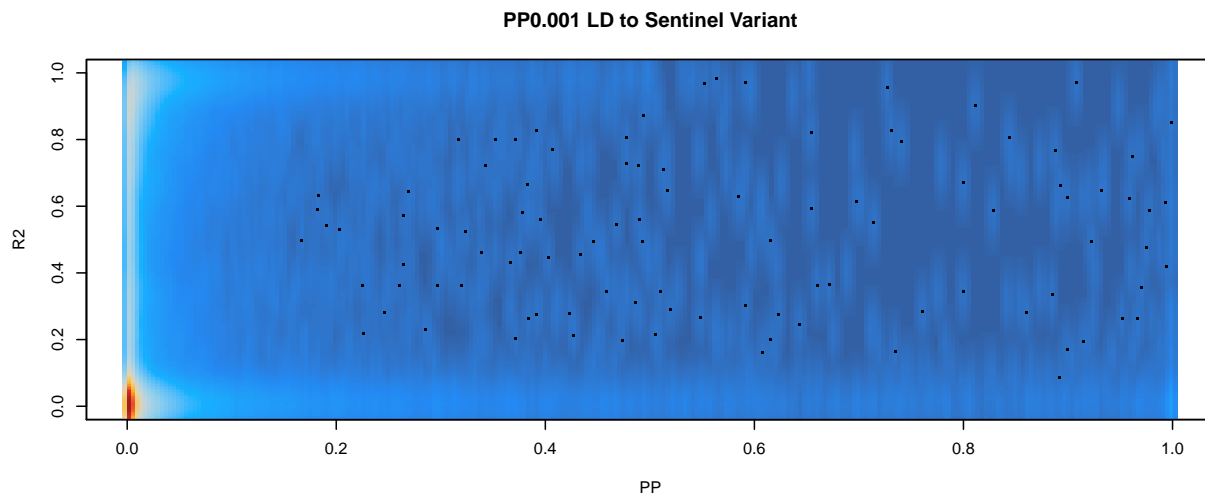
Supplemental Figure 3: Hematopoietic cell type enrichments for Mean Retic Volume and Monocyte count using various methods.



Supplemental Figure 4: All enrichments from g-chromVAR. The horizontal line shows a Bonferroni multiple testing adjusted threshold for statistical significance of enrichment.



Supplemental Figure 5: All enrichments from LD Score. The horizontal line shows a Bonferonni multiple testing adjusted threshold for statistical significance of enrichment.



Supplemental Figure 6: Sample embedding of figure in document.

(B)