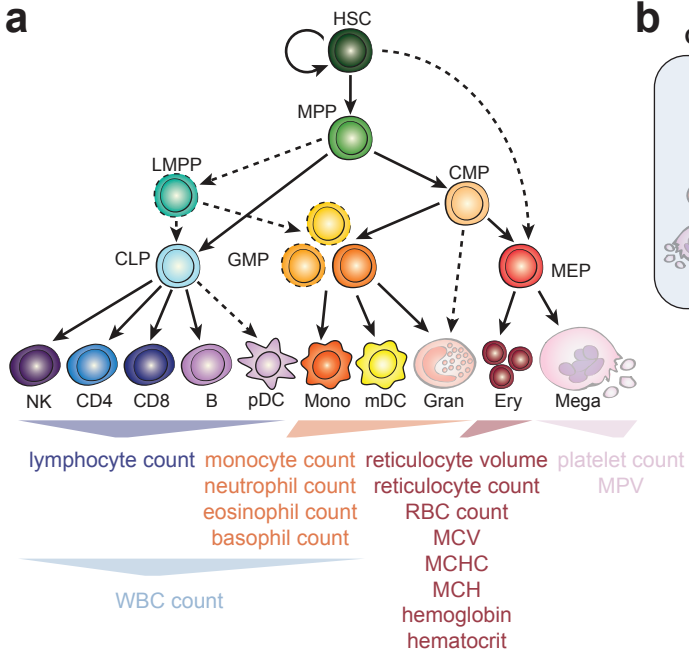
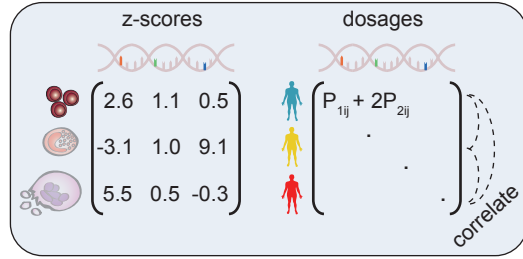


**a**

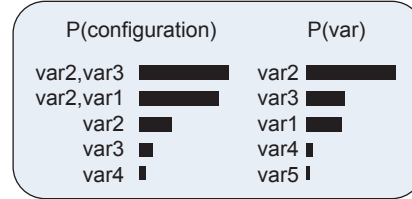


**b**

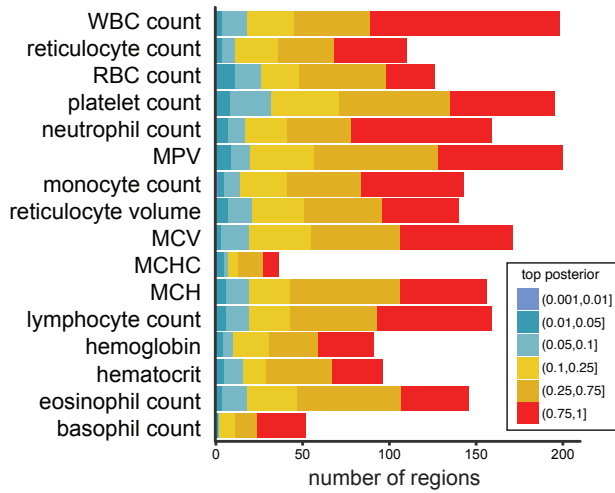
GWAS on UKBB (~113K) for 16 blood cell traits



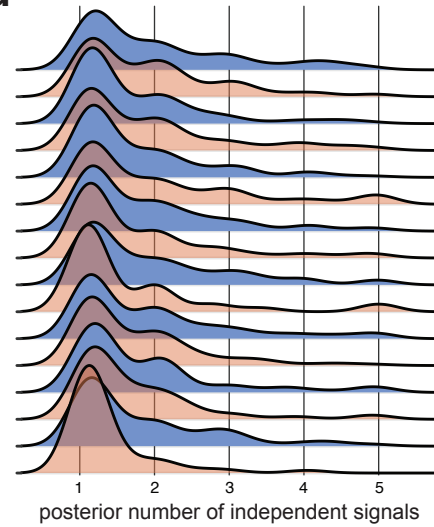
Finemap 2,178 GWAS regions (3MB) allowing multiple "causal" signals



**c**



**d**



**e**

