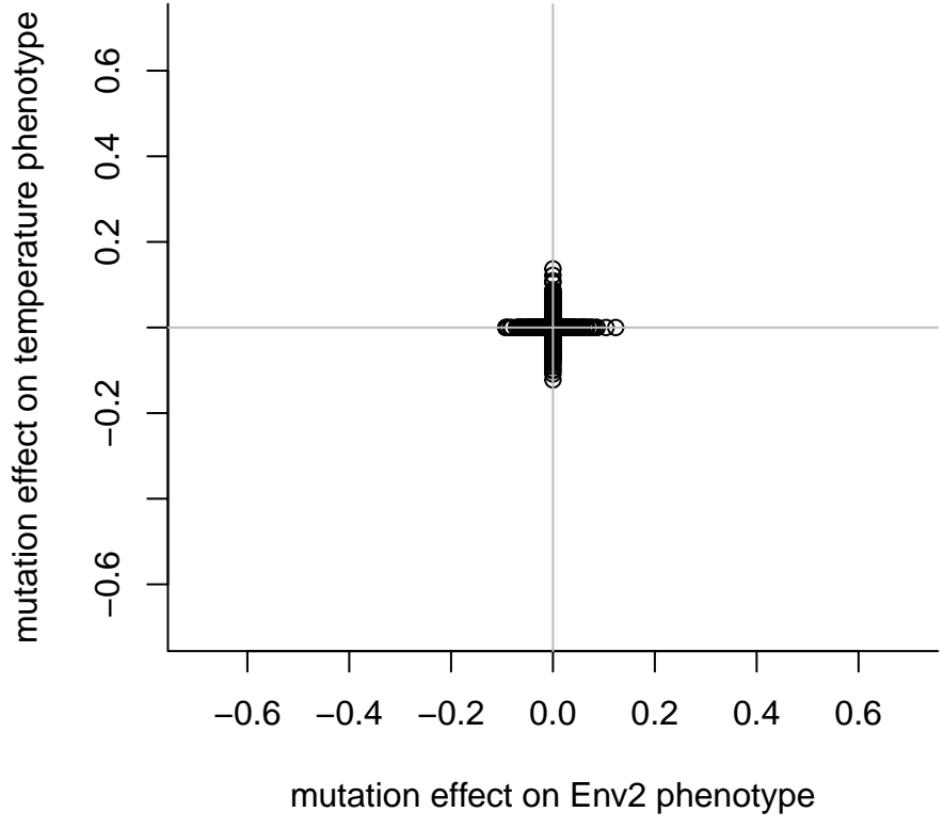
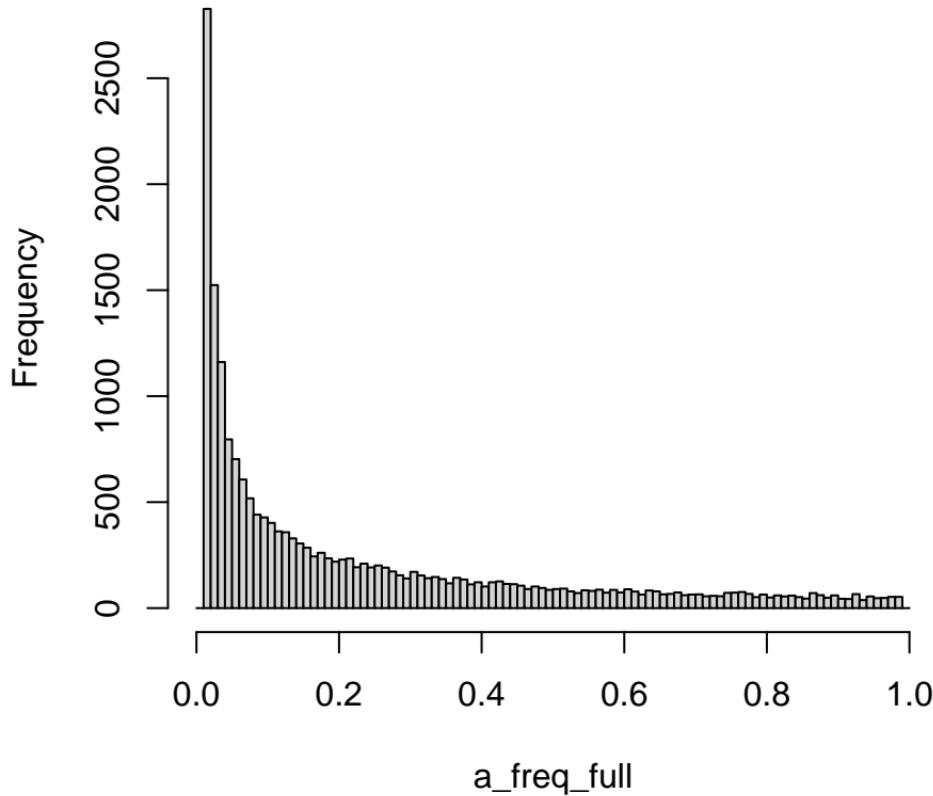


highly-polygenic\_2-trait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-center-to-edge\_m-constant  
1231129



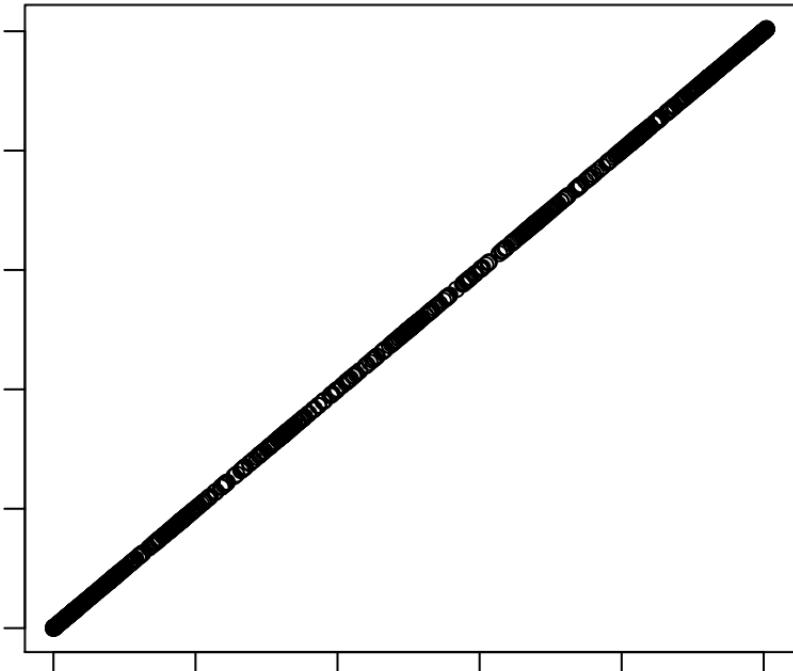
## Histogram of a\_freq\_full

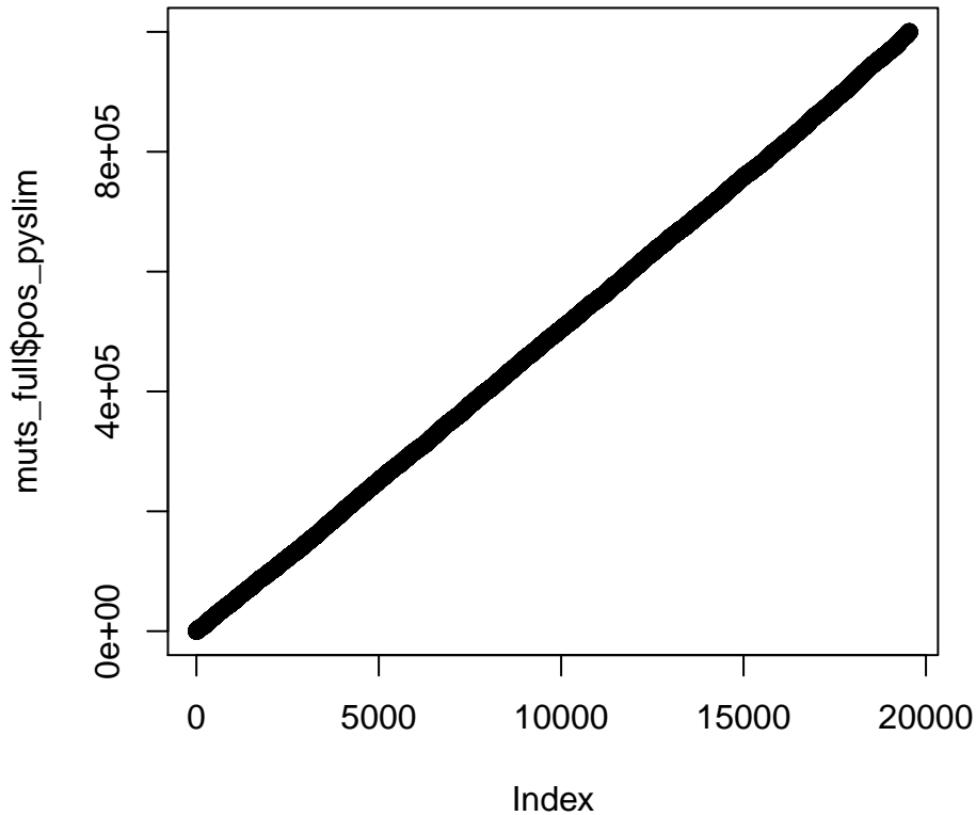


indPhen\_df\$indID[which(indPhen\_df\$subset)]

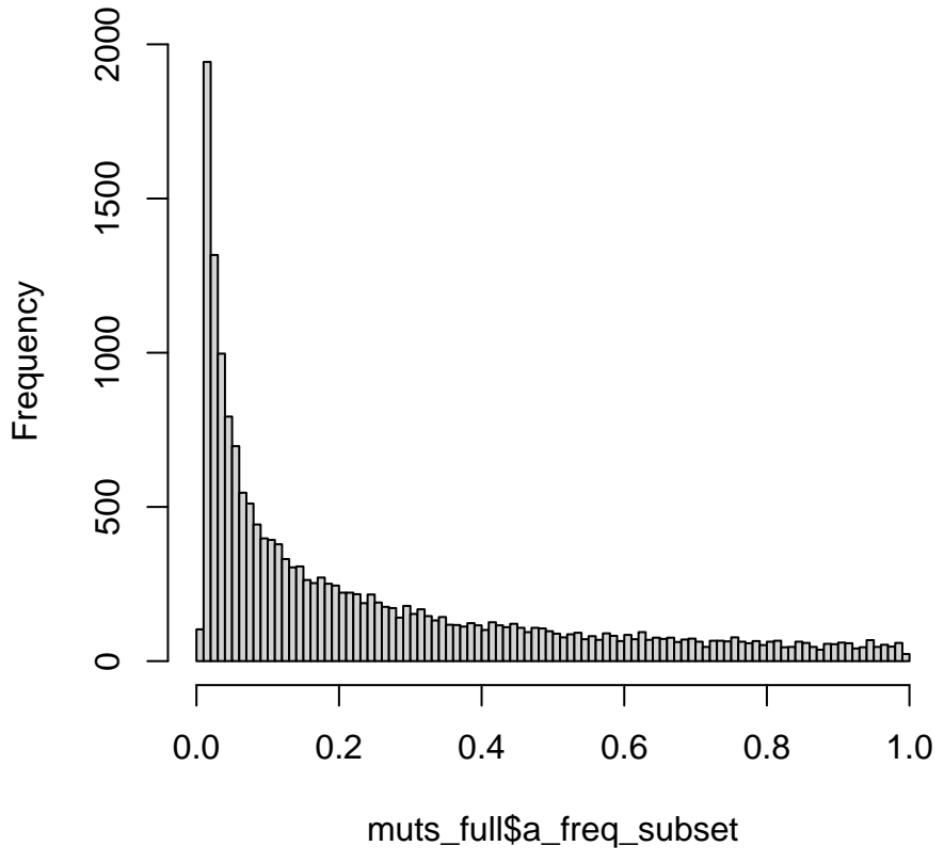
0 2000 4000 6000 8000 10000

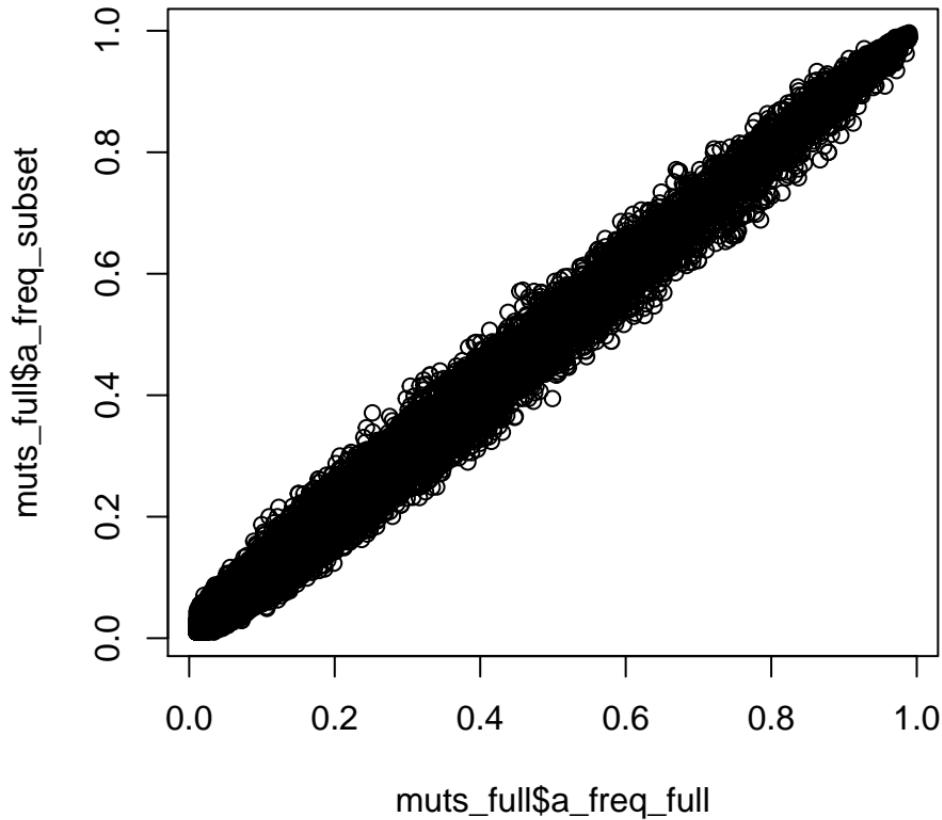
colnames(G\_full\_subset)



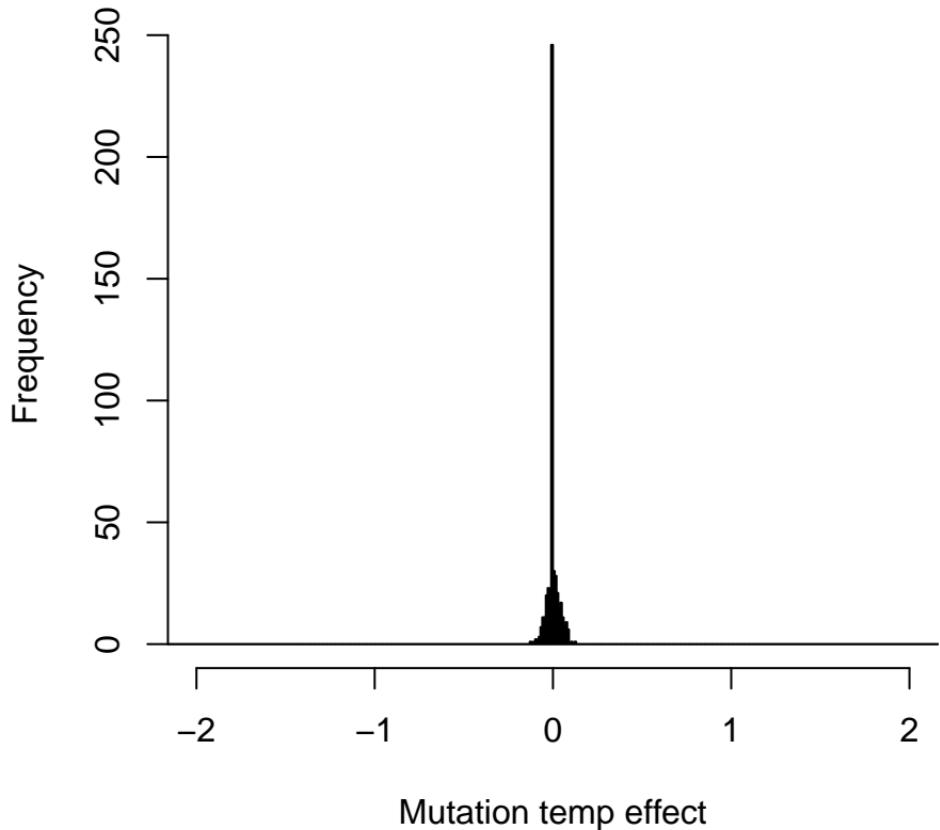


## Histogram of muts\_full\$a\_freq\_subset

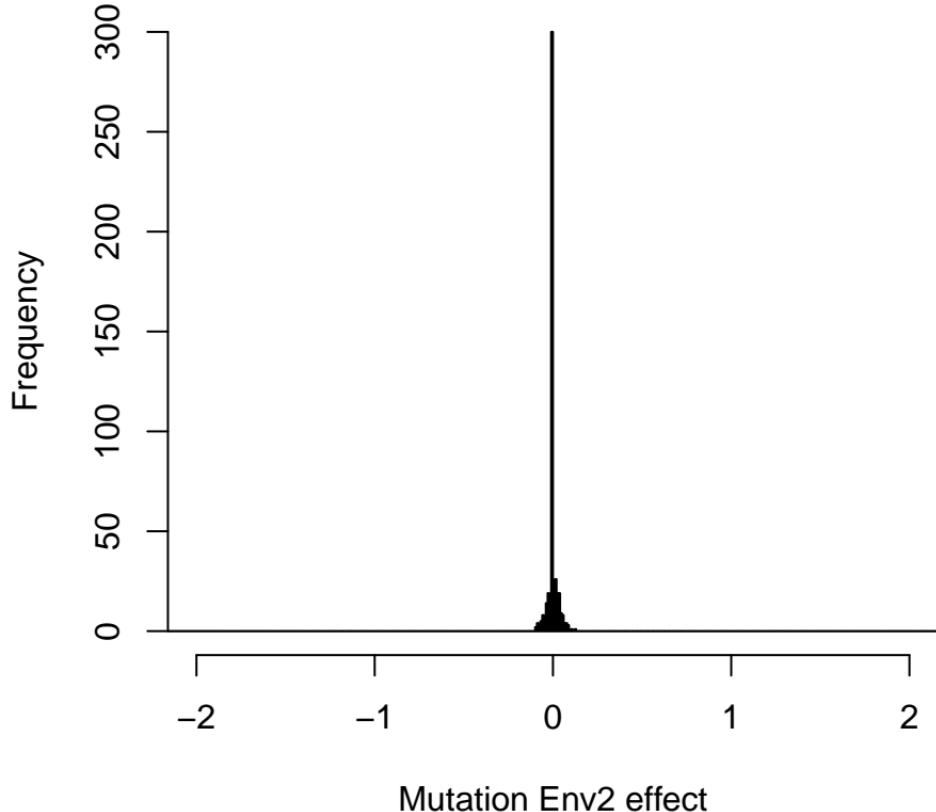




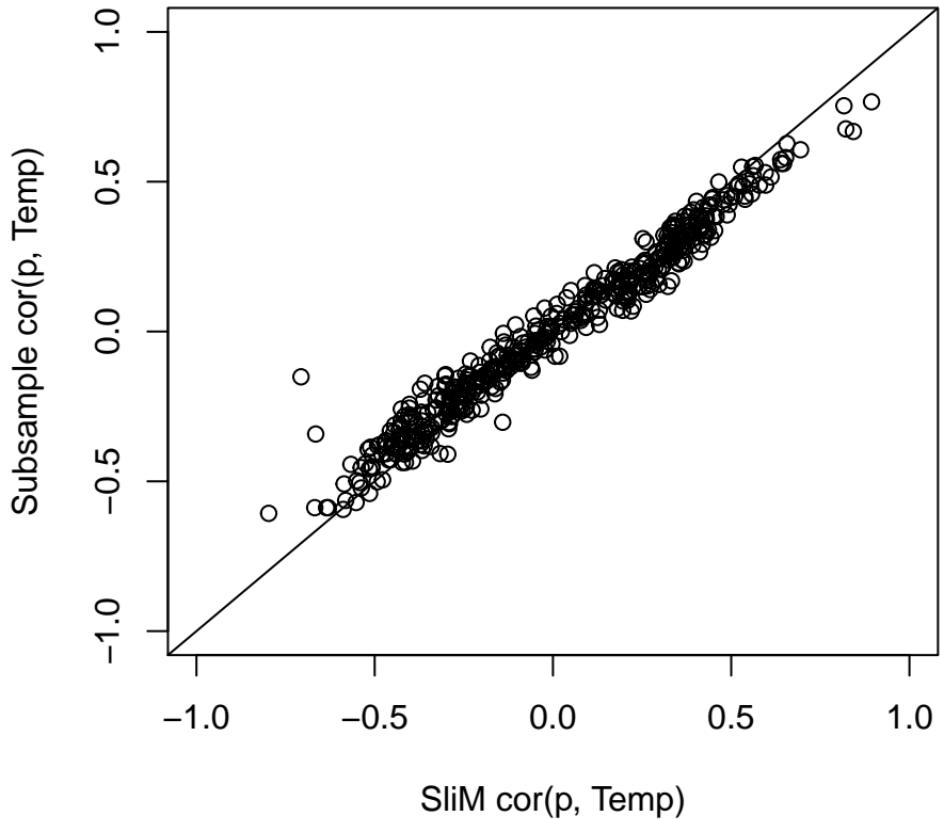
highly-polygenic\_2-trait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-center-to-edge\_m-constant  
1231129



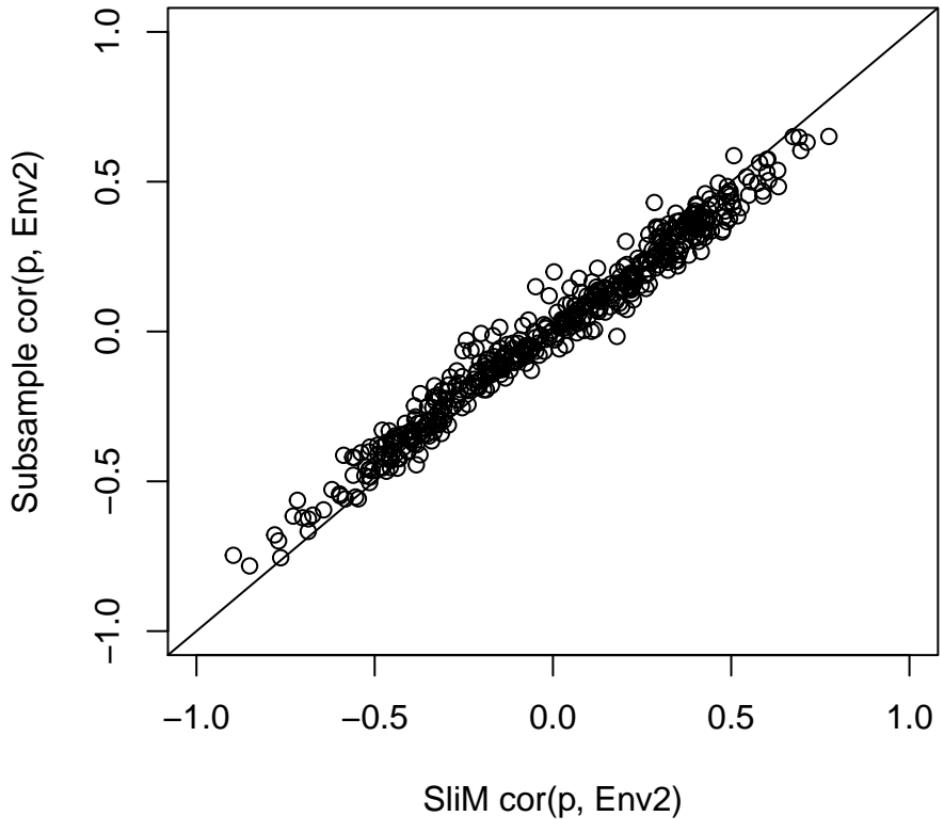
highly-polygenic\_2-trait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-center-to-edge\_m-constant  
1231129



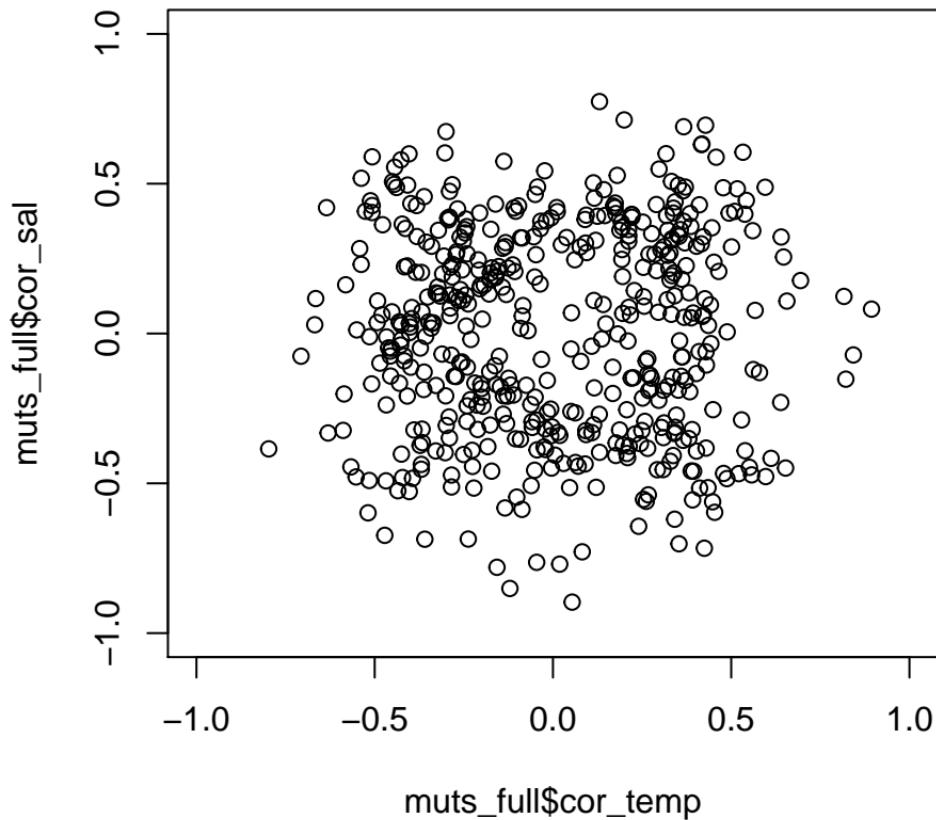
ait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-ce  
1231129



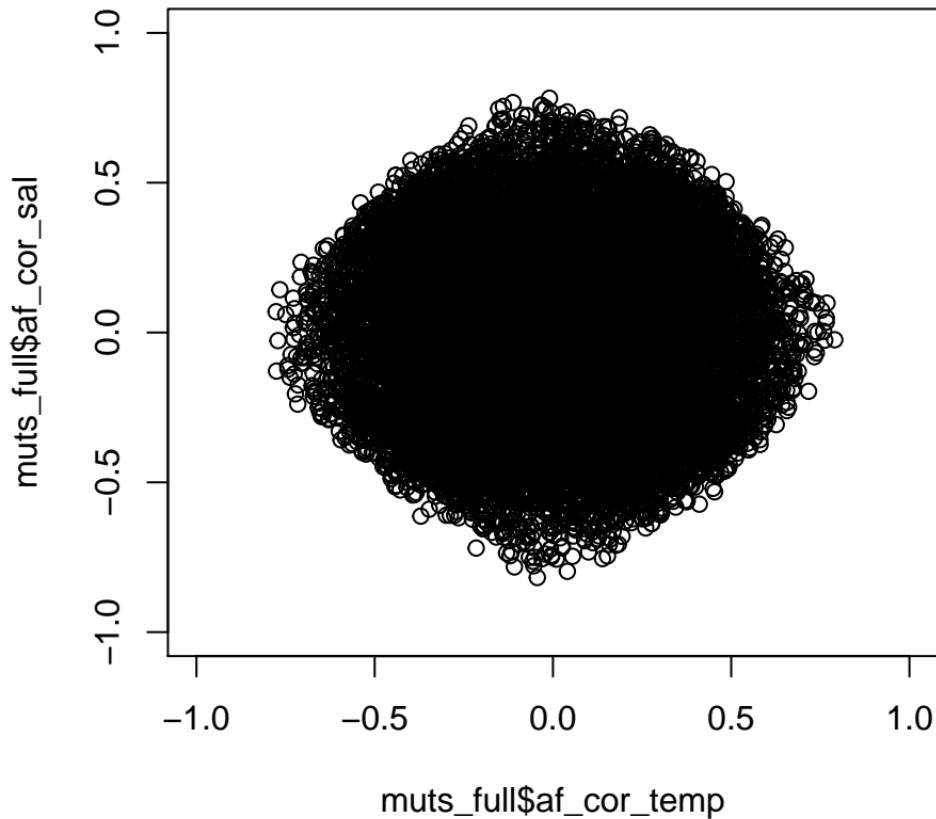
ait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-ce  
1231129



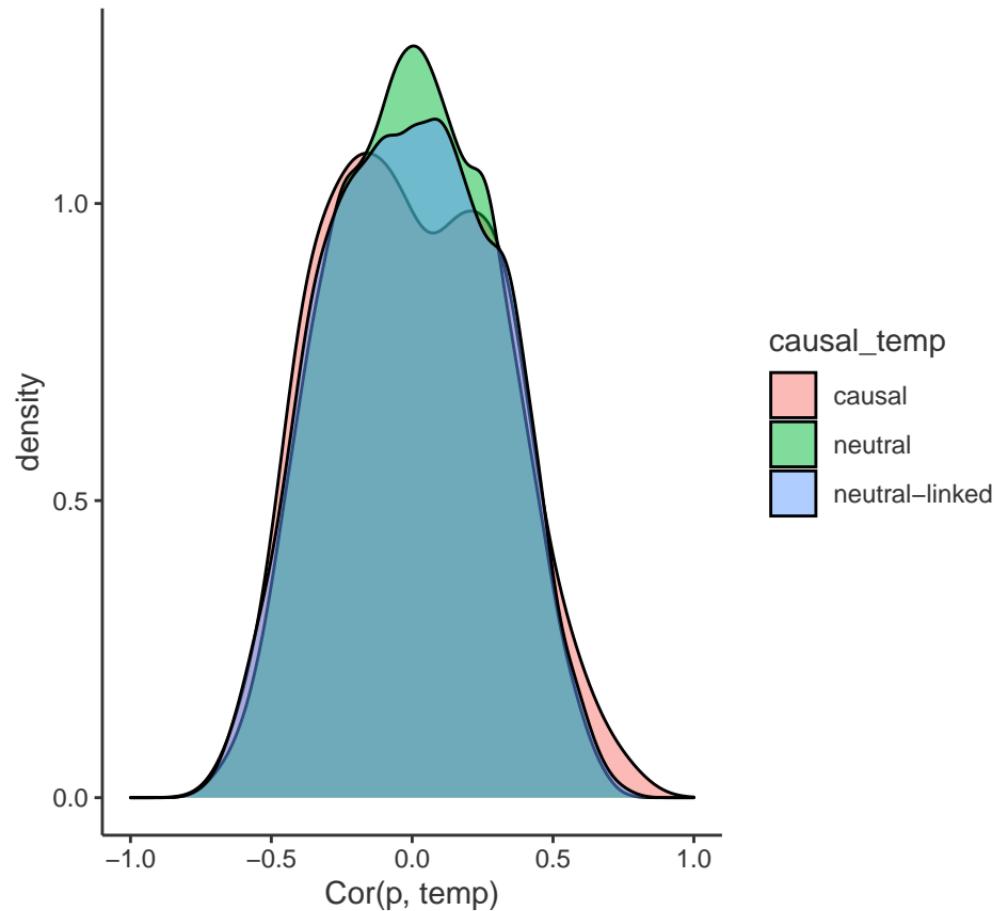
## SiM Correlation (AF, ENV) all samples



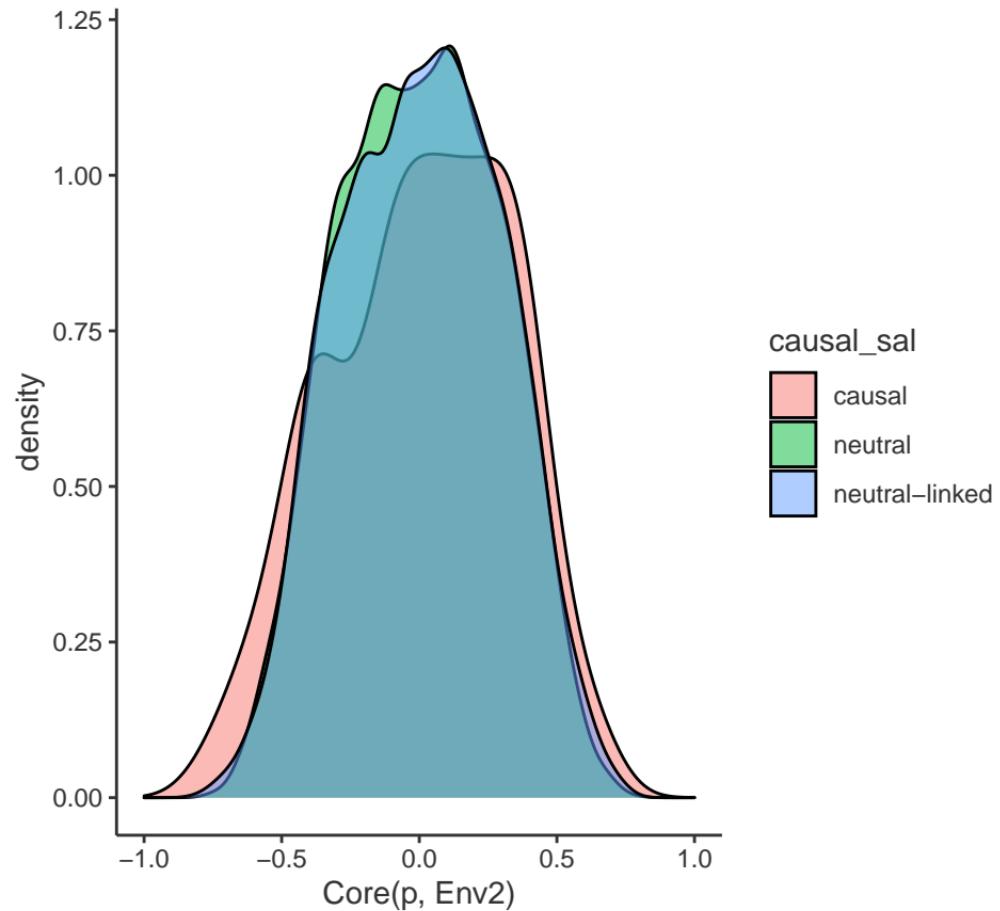
## Subsample Correlation (AF, ENV) all samples



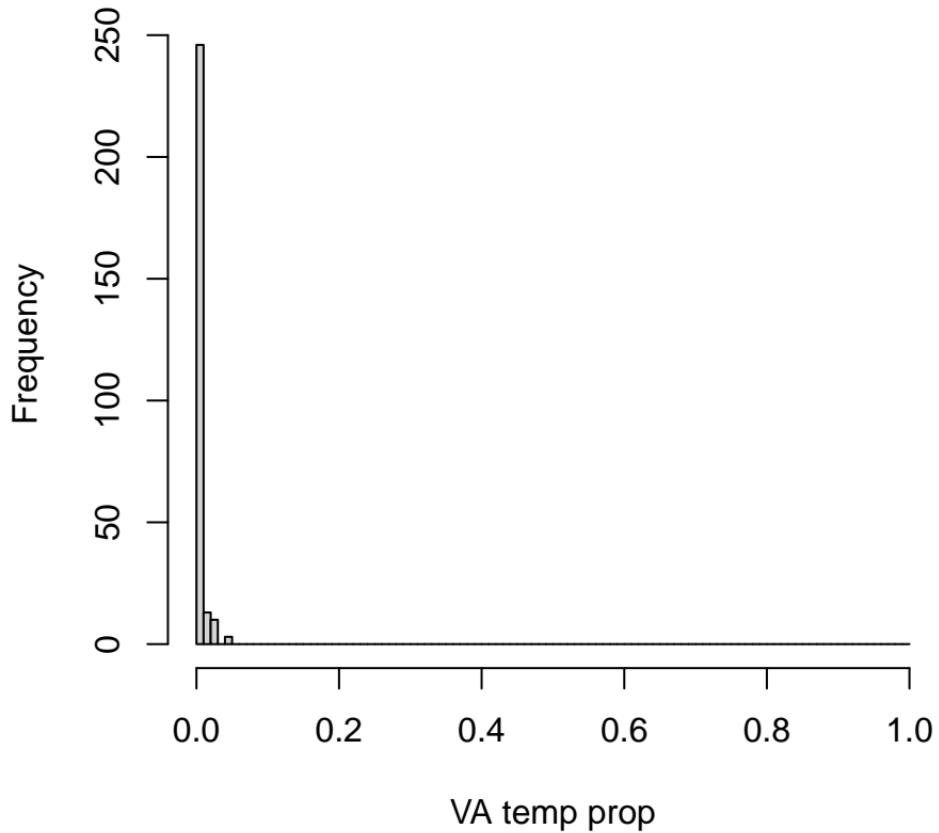
highly-polygenic\_2-trait-no-pleiotropy-unequal-S\_SS  
1231129



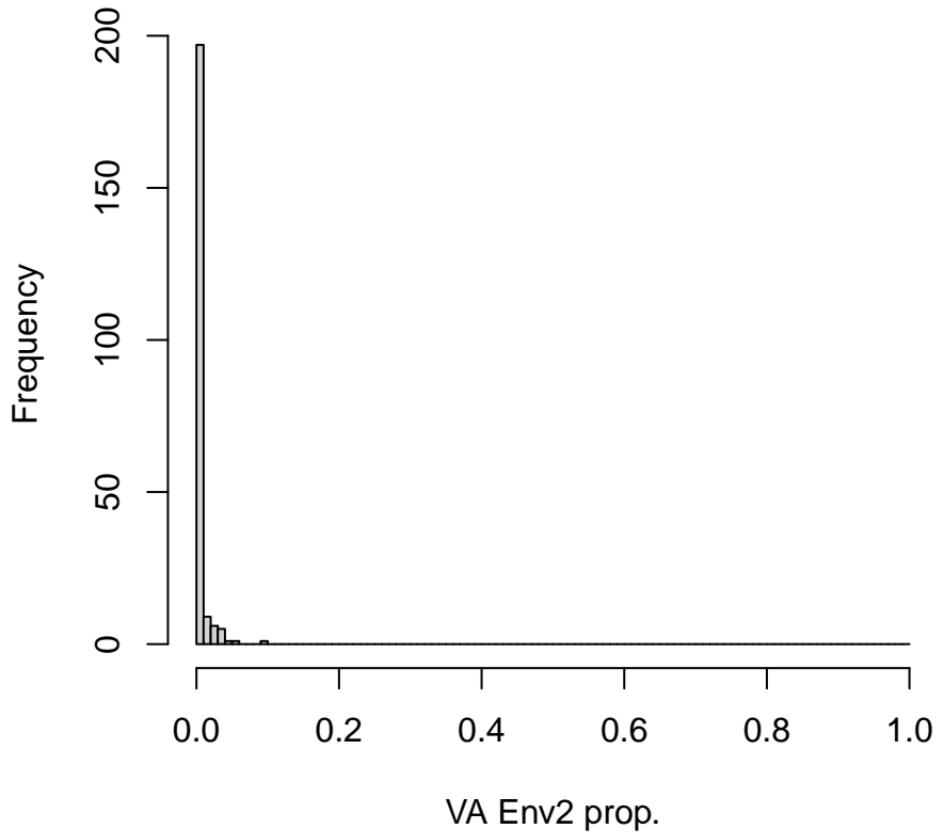
highly-polygenic\_2-trait-no-pleiotropy-unequal-S\_S  
1231129



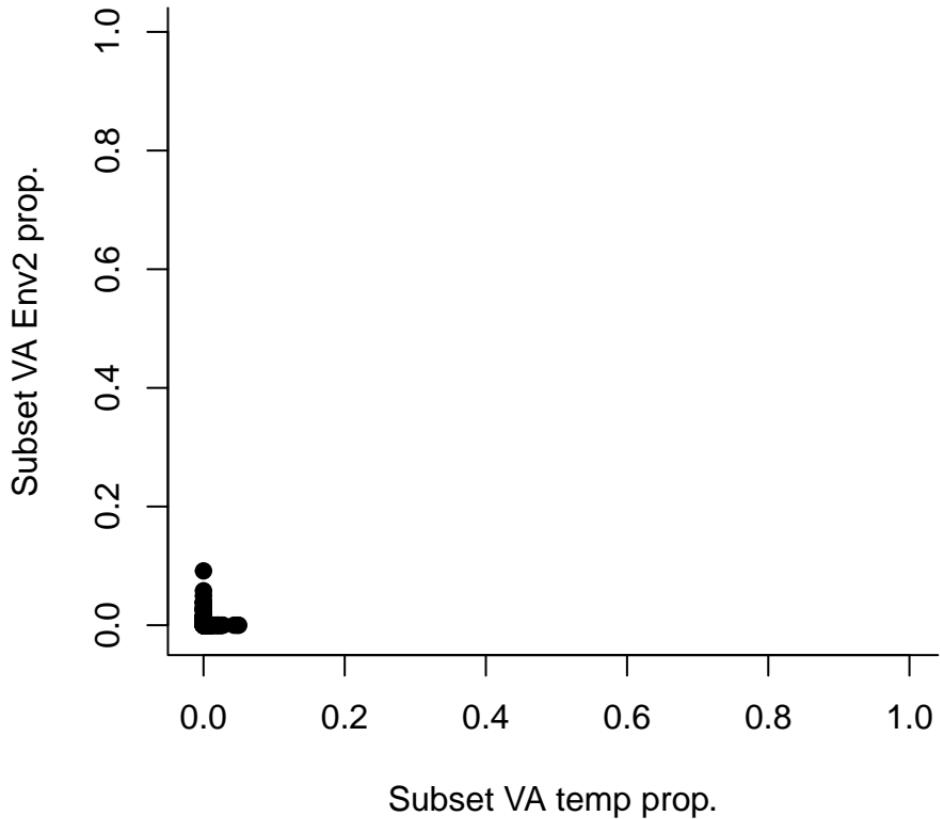
ait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-ce  
1231129



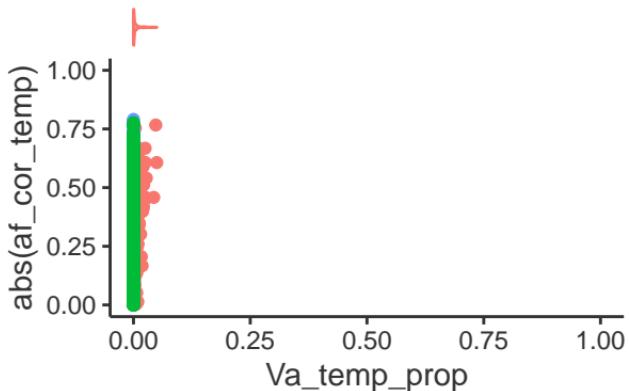
ait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-ce  
1231129



ait-no-pleiotropy-unequal-S\_SS-Clines\_N-cline-ce  
1231129



highly-polygenic\_2-trait-no-pl  
1231129

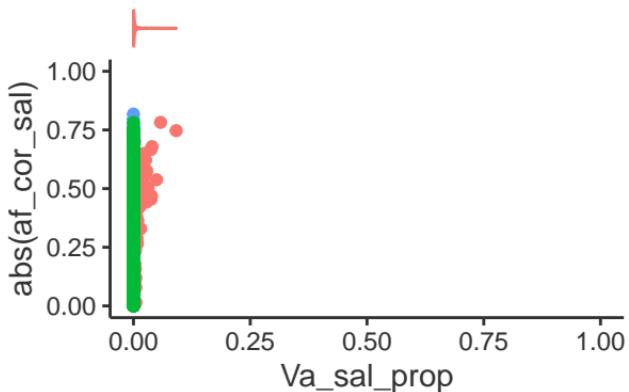


causal\_temp

- causal
- neutral
- neutral-linked



highly-polygenic\_2-trait-no-pl  
1231129



causal\_sal

- causal
- neutral
- neutral-linked

