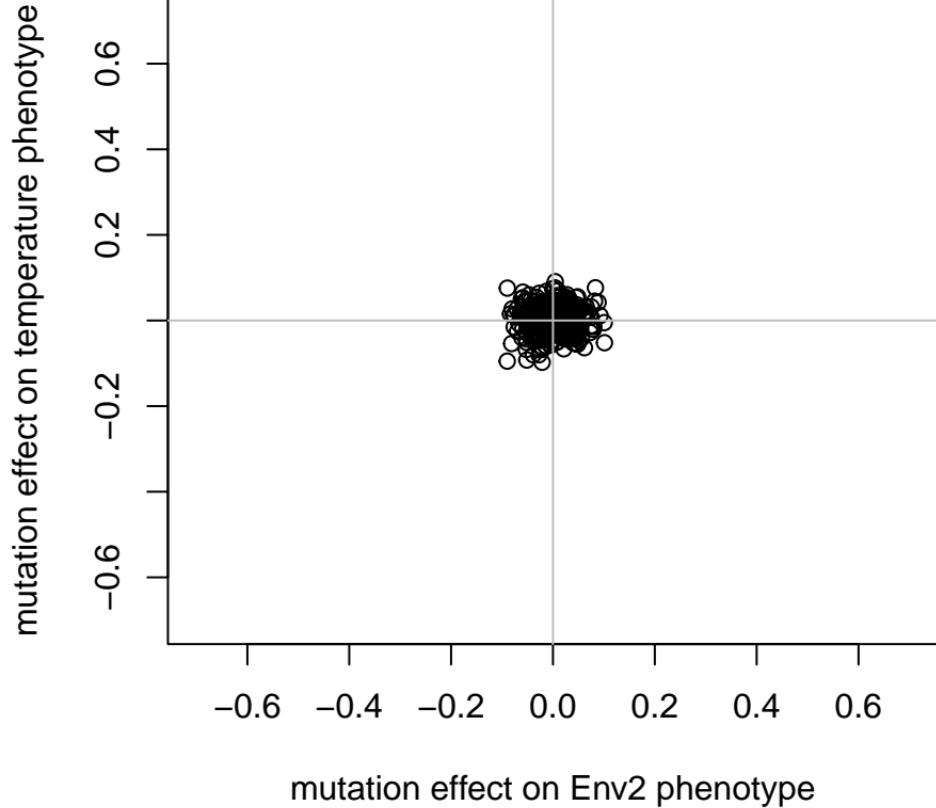
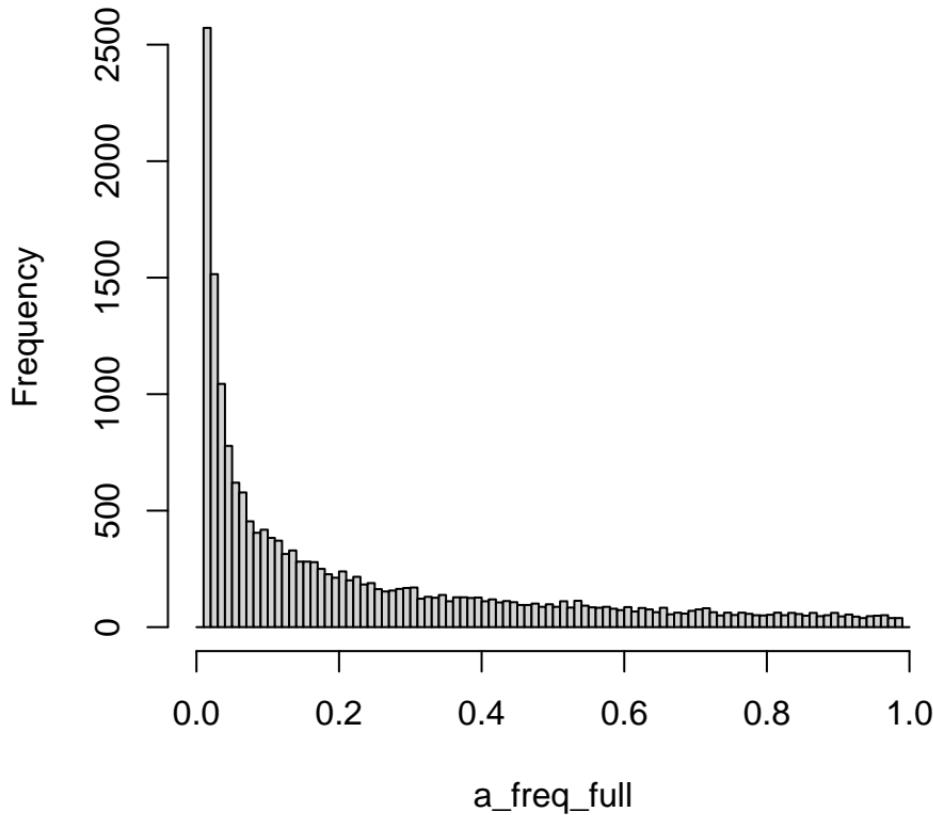


polygenic\_2-trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to-edge\_m-constant  
1231219



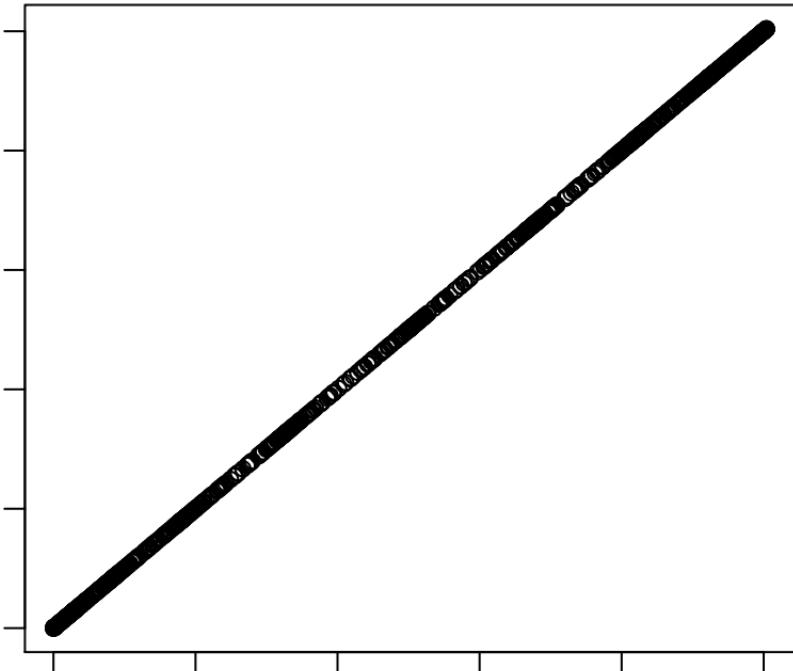
# Histogram of a\_freq\_full

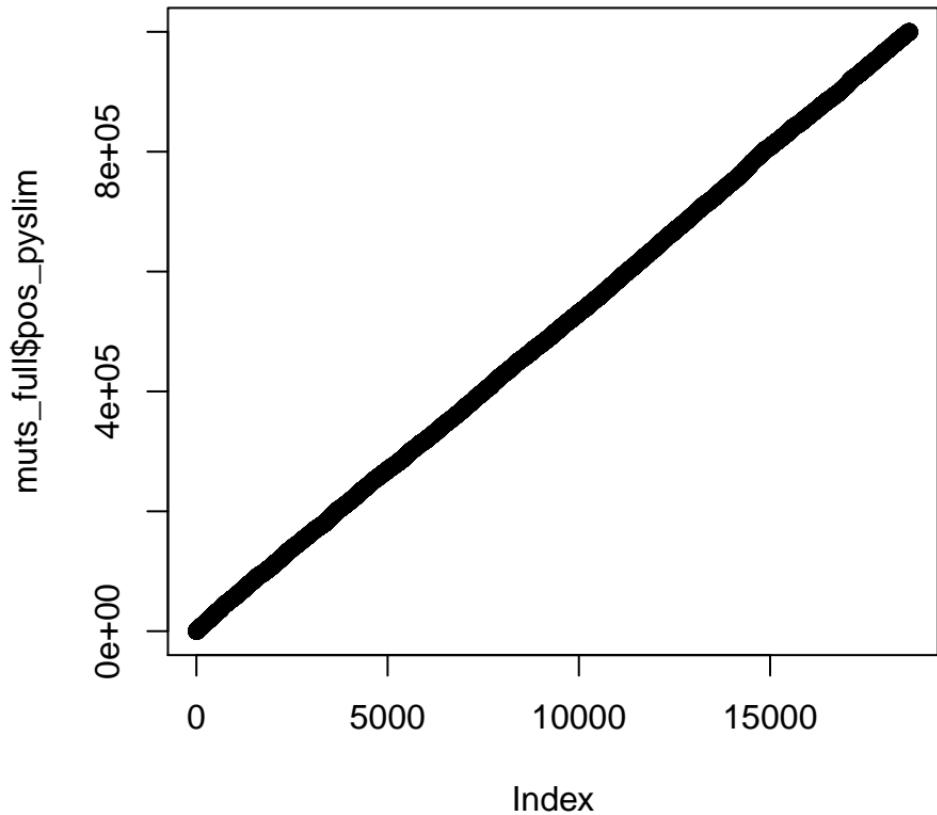


indPhen\_df\$indID[which(indPhen\_df\$subset)]

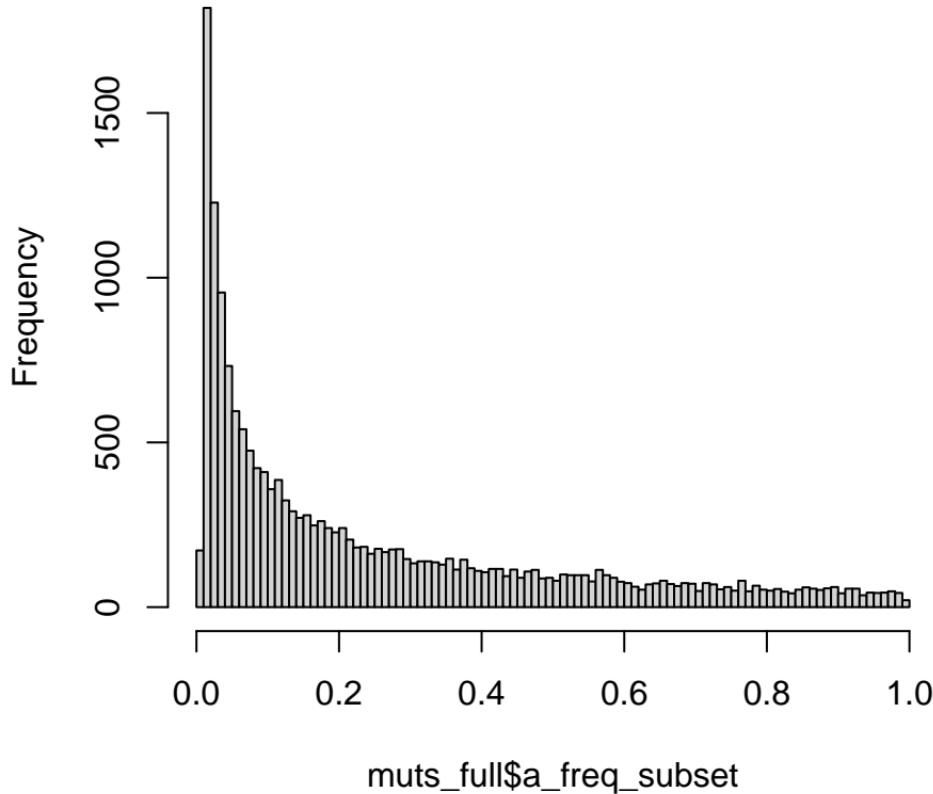
0 2000 4000 6000 8000 10000

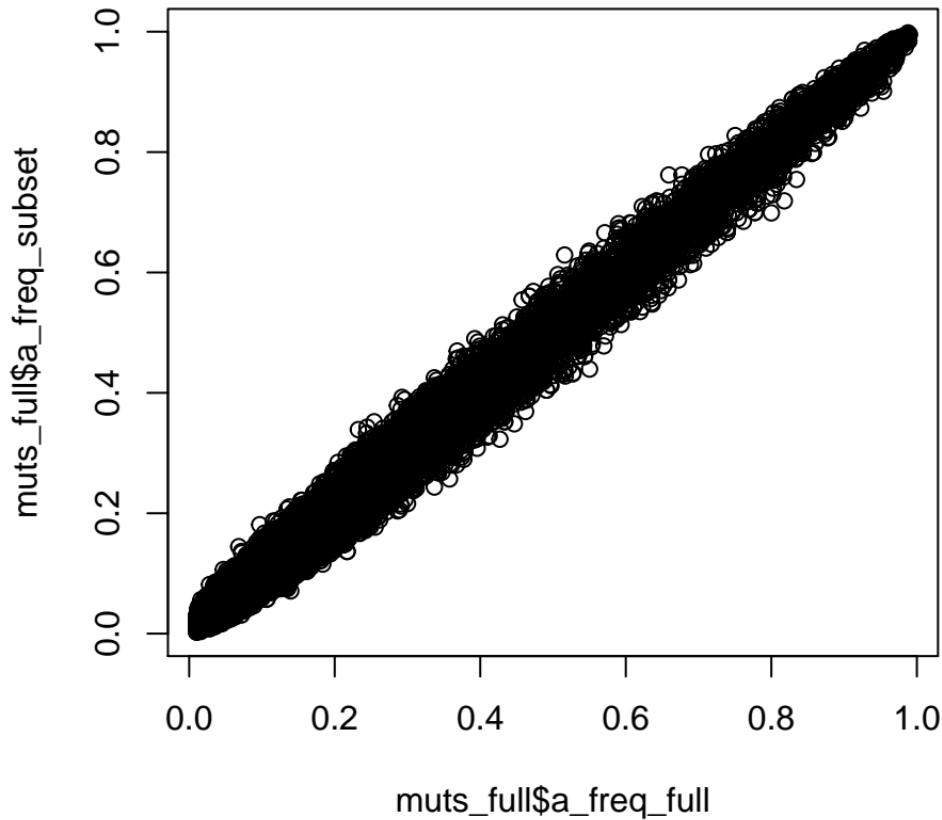
colnames(G\_full\_subset)



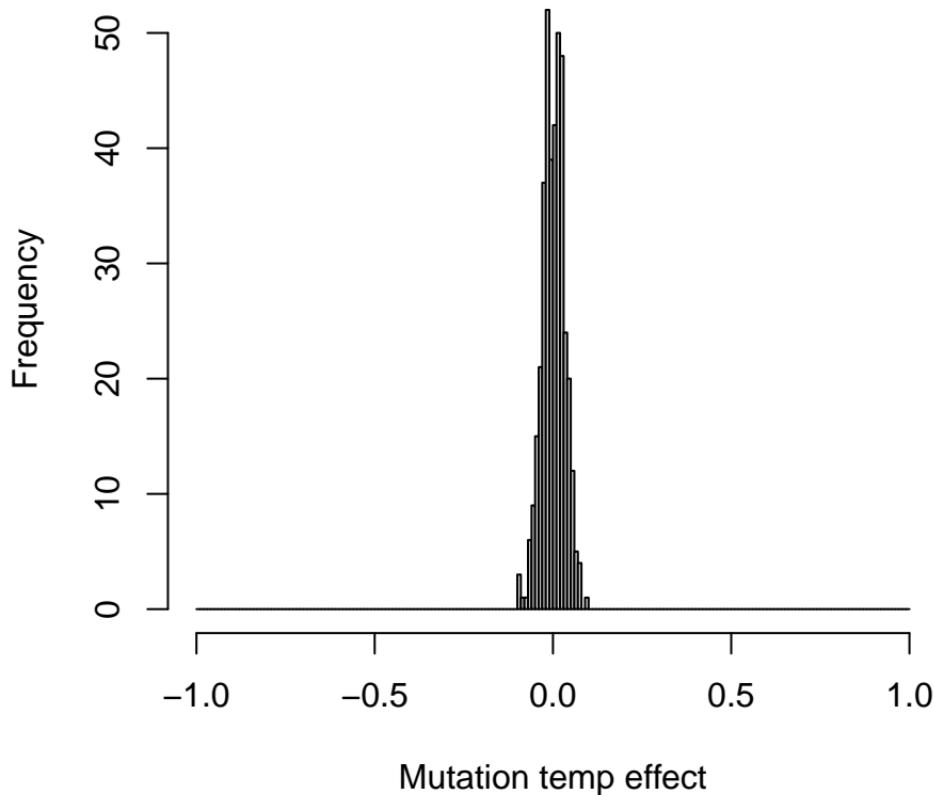


## Histogram of muts\_full\$a\_freq\_subset

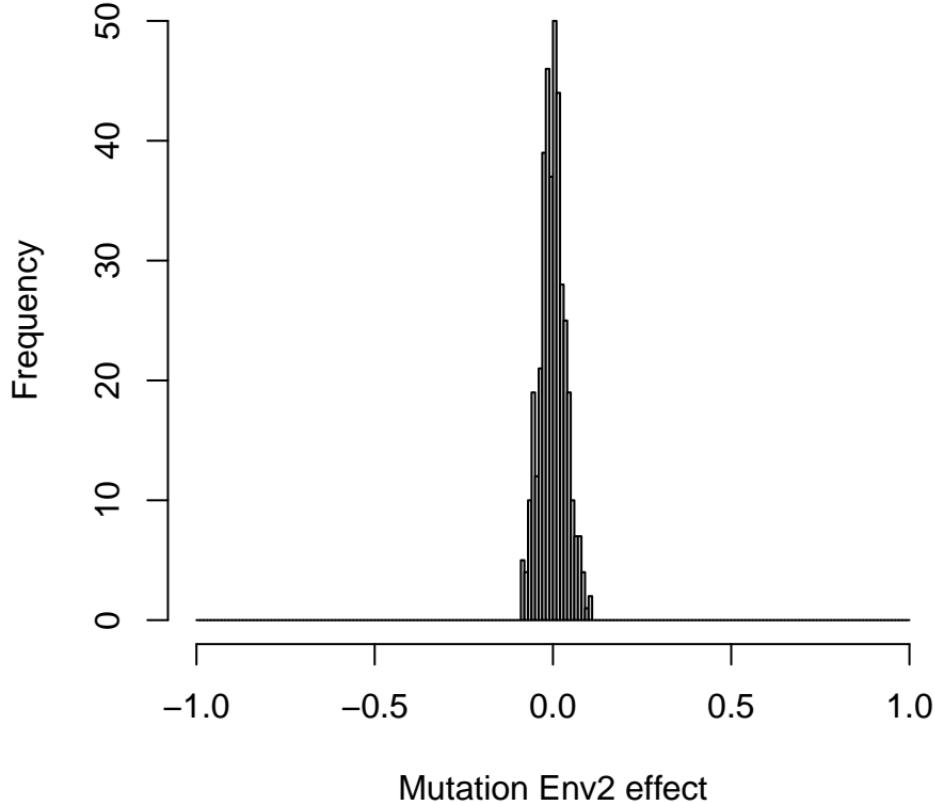




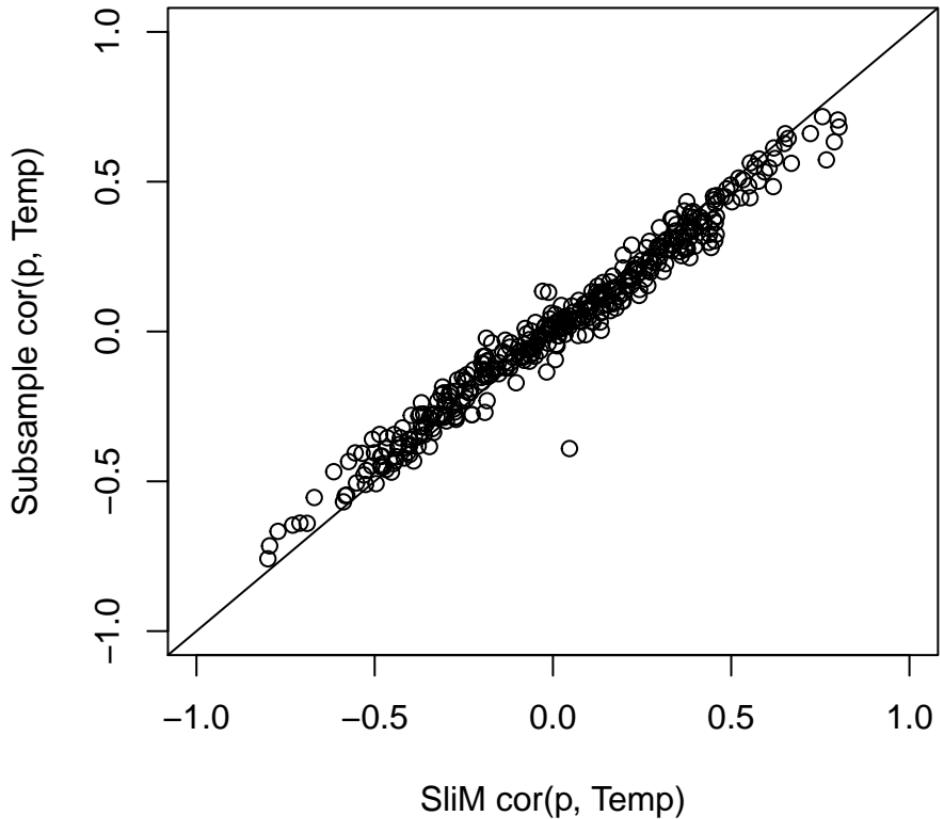
polygenic\_2-trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to-edge\_m-constant  
1231219



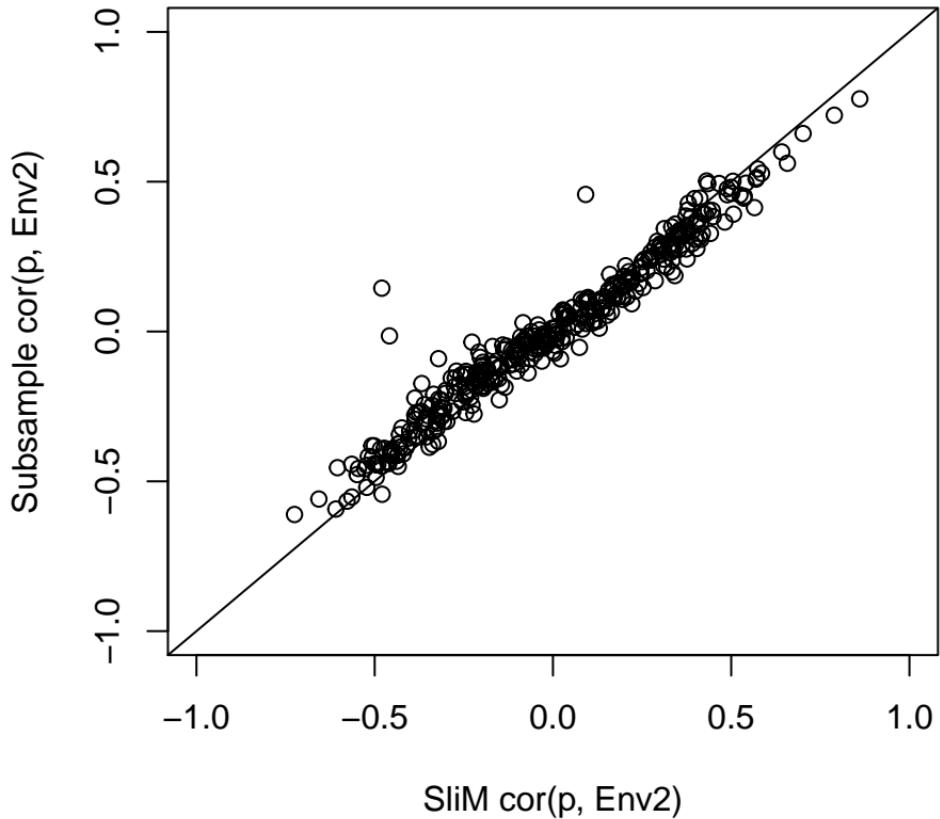
polygenic\_2-trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to-edge\_m-constant  
1231219



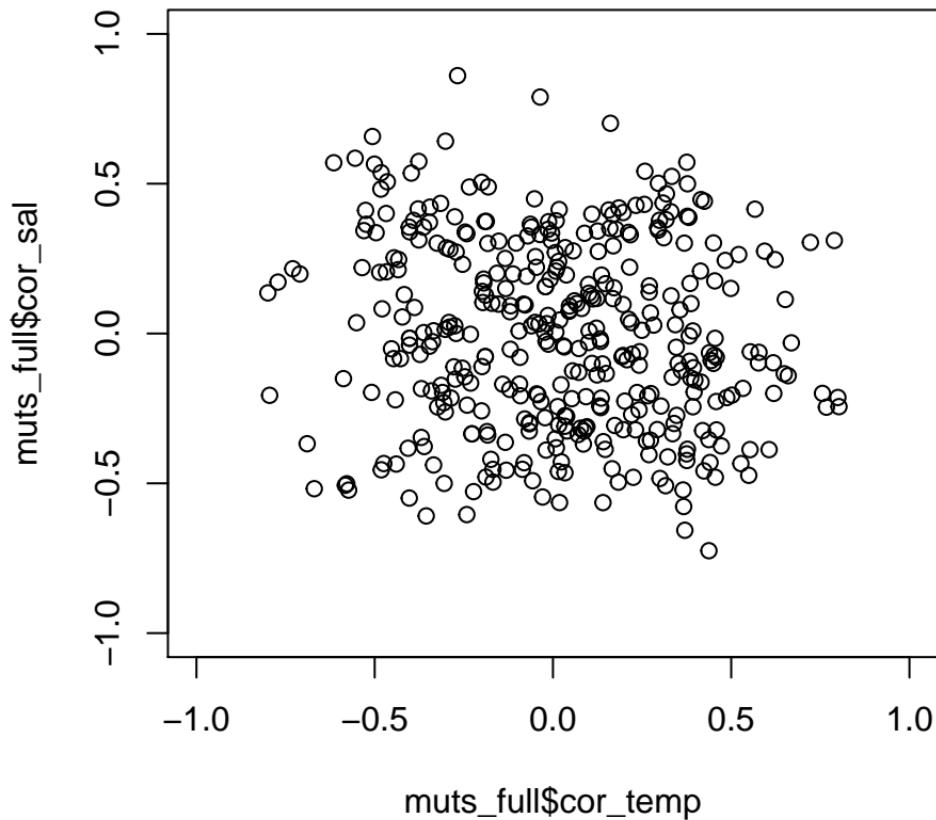
trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to  
1231219



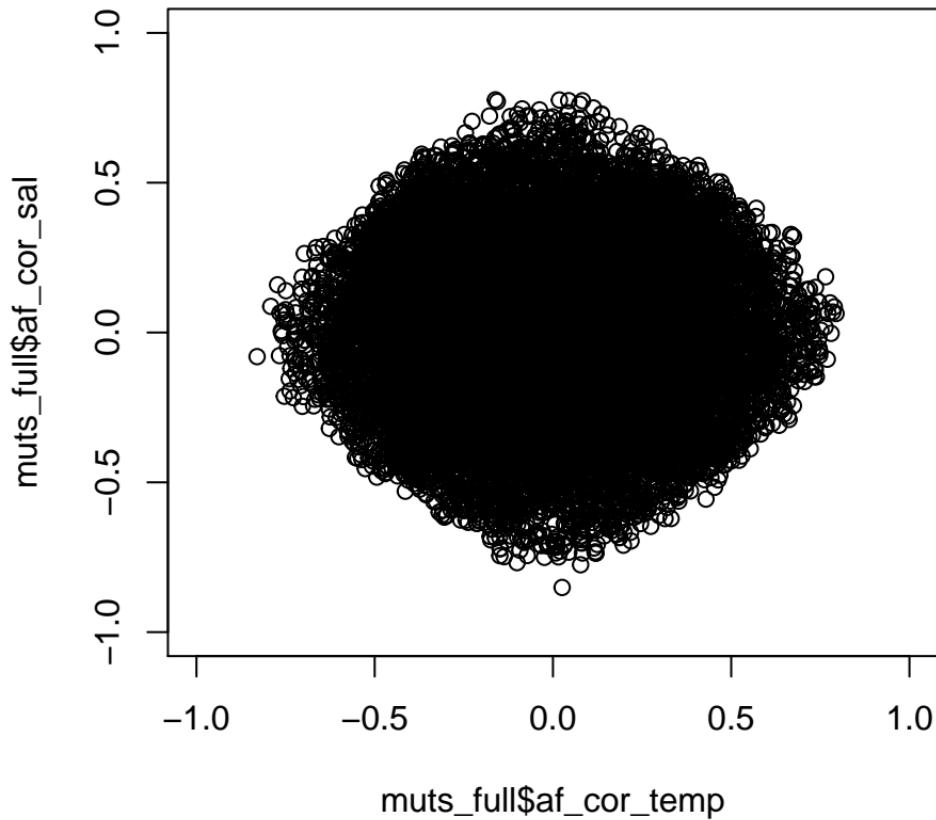
trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to  
1231219



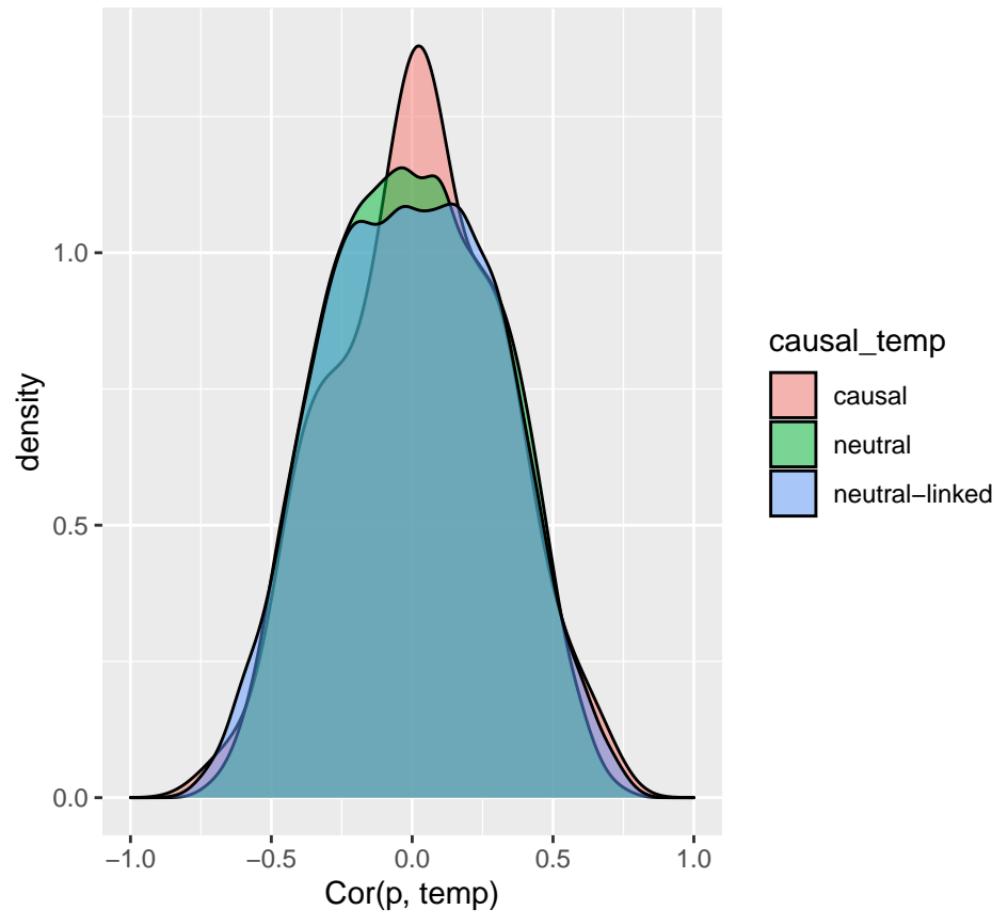
## SiM Correlation (AF, ENV) all samples



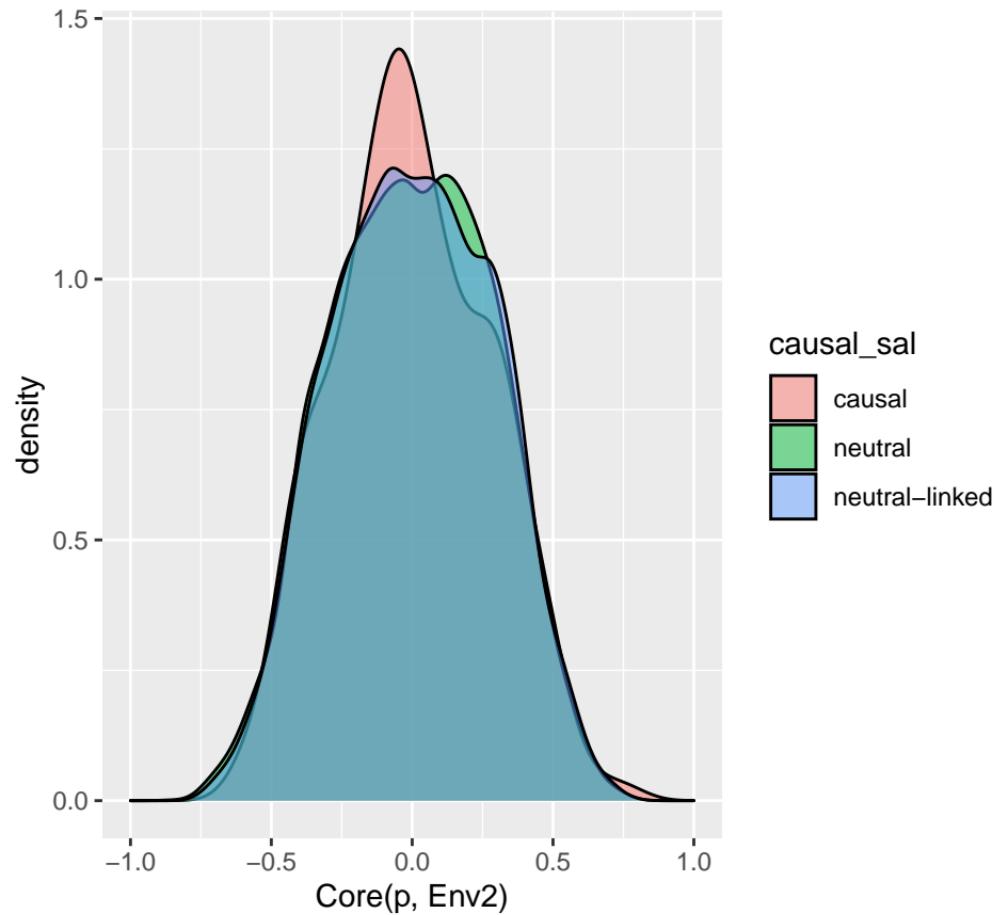
## Subsample Correlation (AF, ENV) all samples



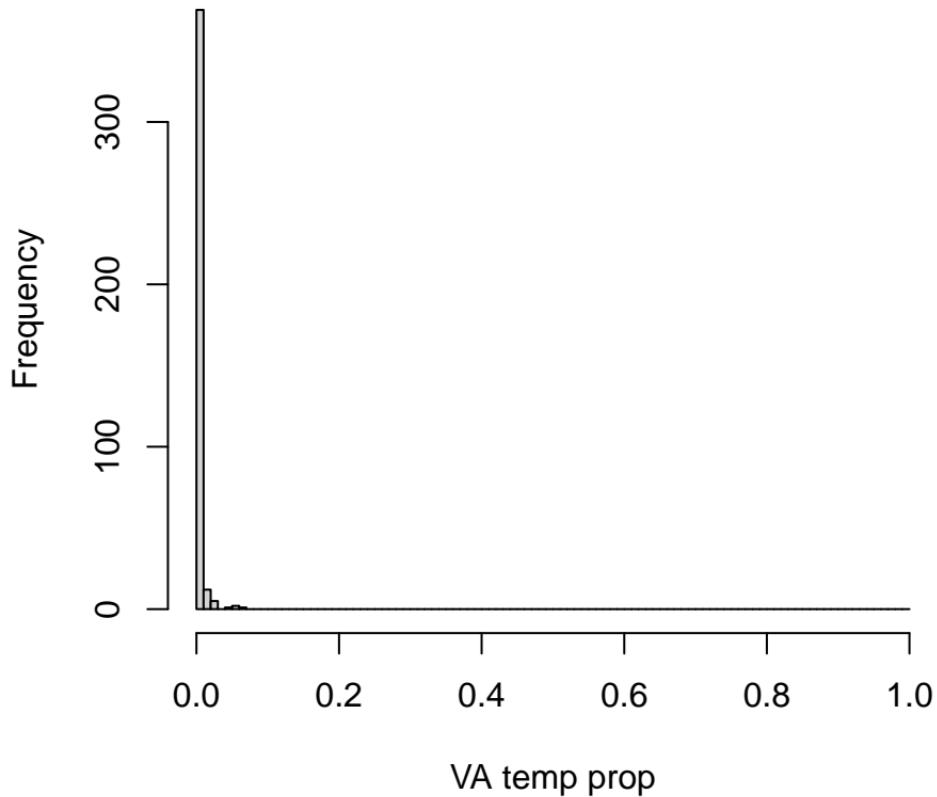
polygenic\_2-trait-pleiotropy-equal-S\_SS-Clines\_N-c  
1231219



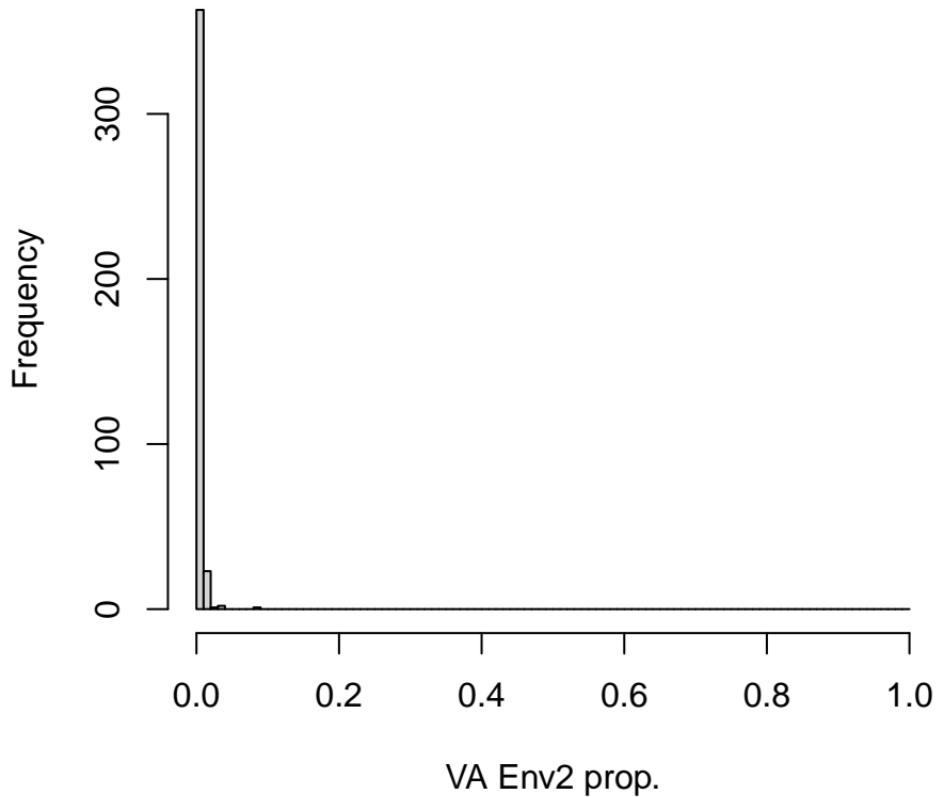
polygenic\_2-trait-pleiotropy-equal-S\_SS-Clines\_N-c  
1231219



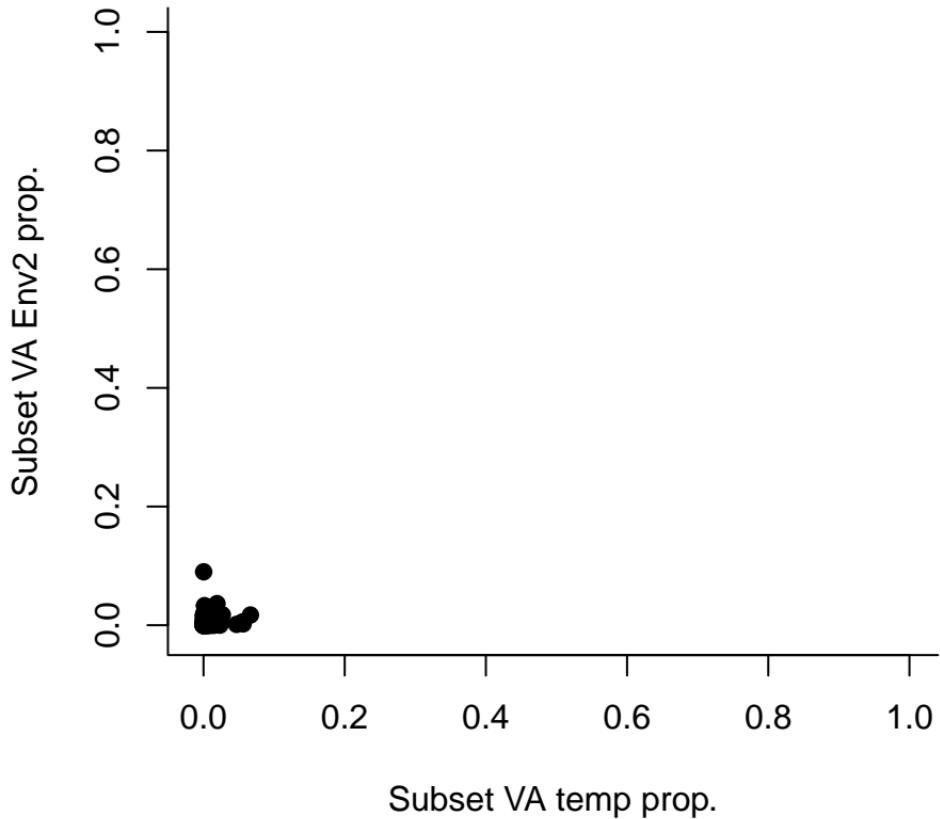
trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to-  
1231219



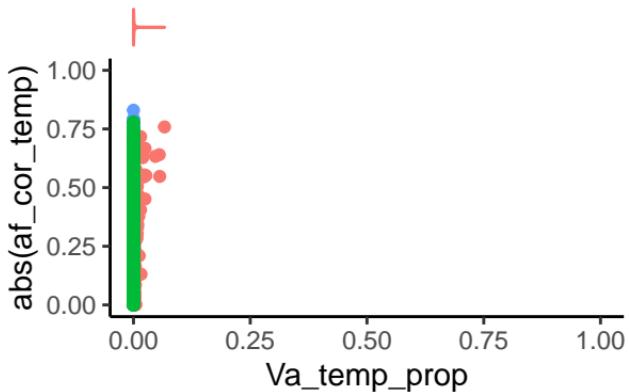
trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to  
1231219



trait-pleiotropy-equal-S\_SS-Clines\_N-cline-center-to  
1231219

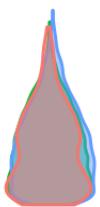


polygenic\_2-trait-pleiotropy-ec  
1231219

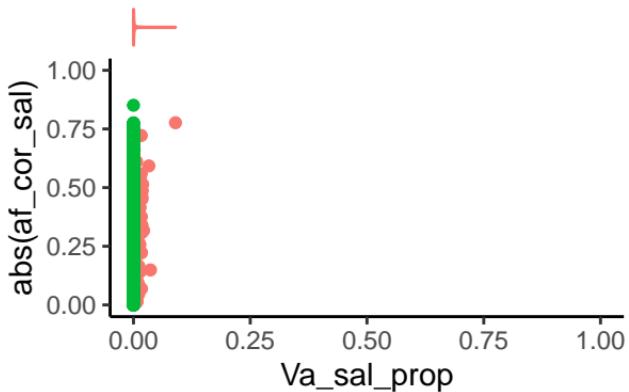


causal\_temp

- causal
- neutral
- neutral-linked



polygenic\_2-trait-pleiotropy-ec  
1231219



causal\_sal

- causal
- neutral
- neutral-linked

