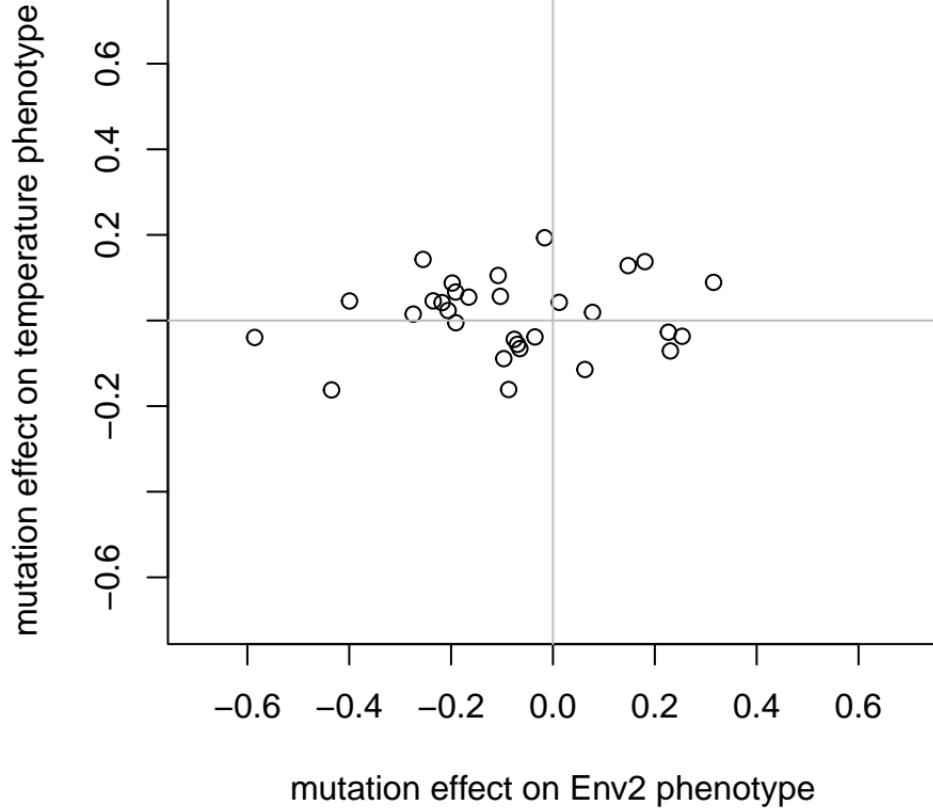
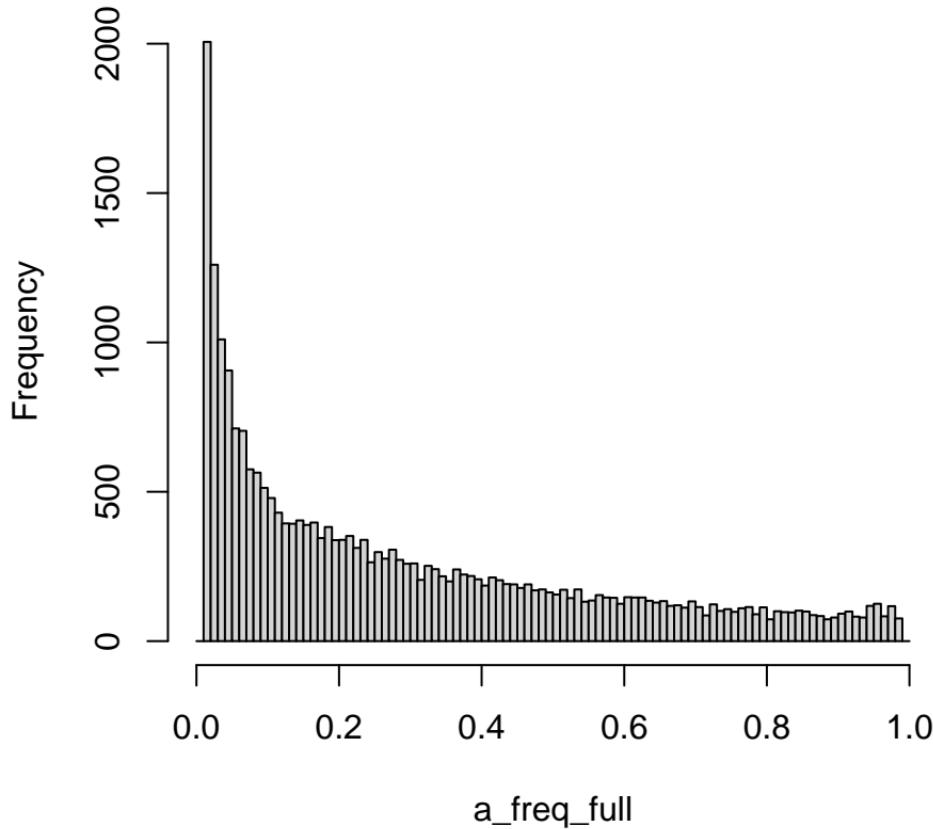


mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1232114



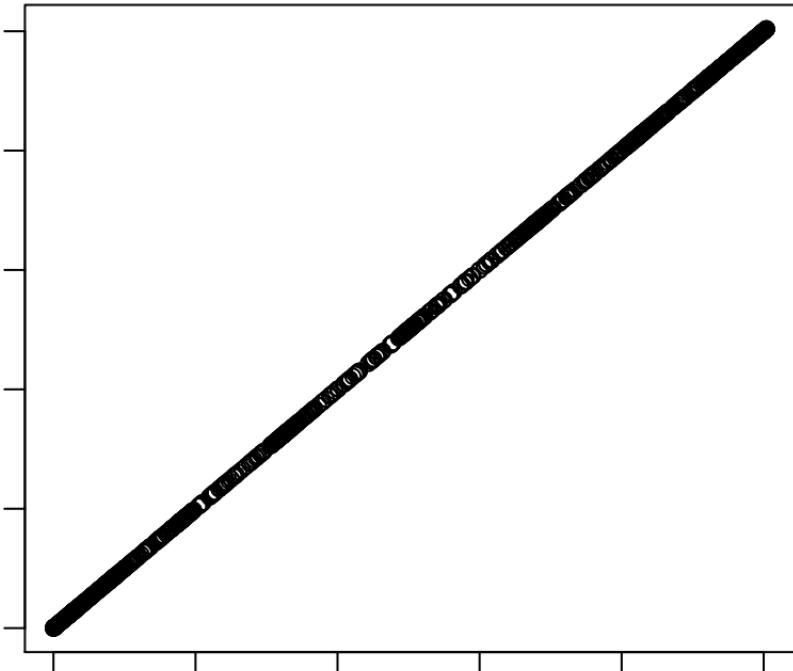
# Histogram of a\_freq\_full

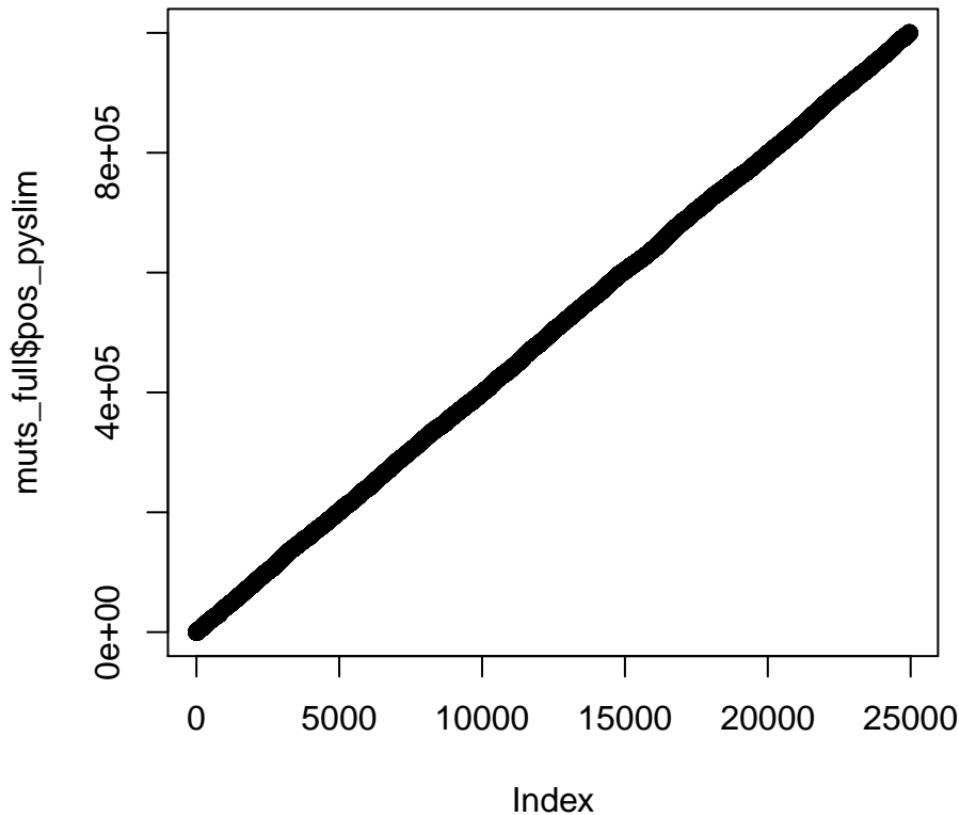


indPhen\_df\$indID[which(indPhen\_df\$subset)]

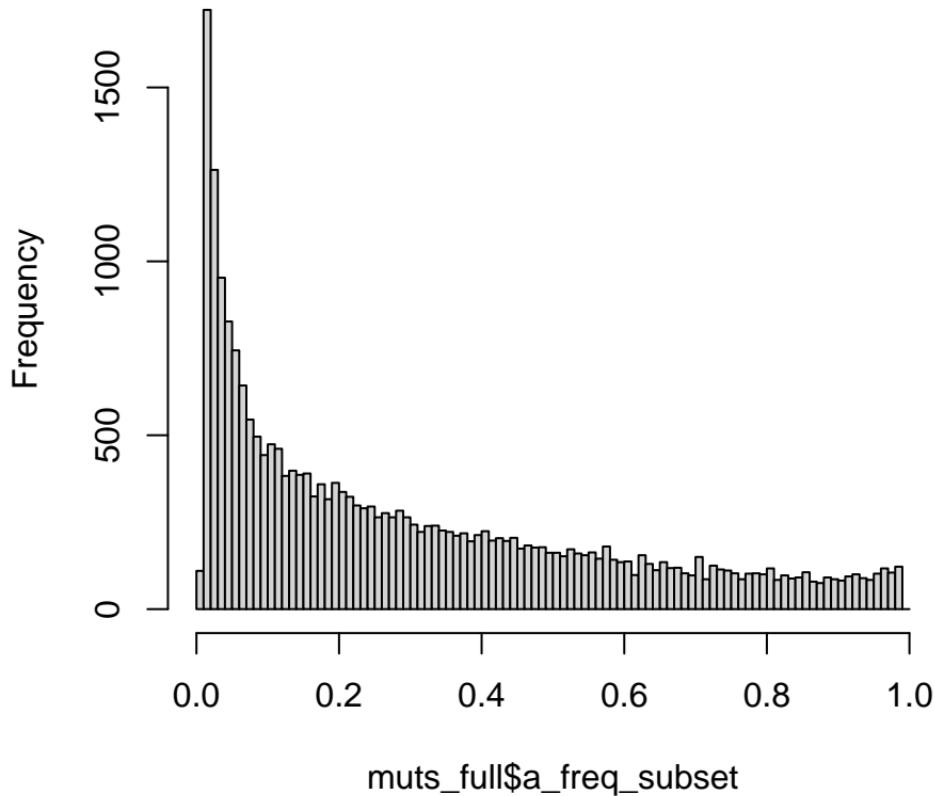
0 2000 4000 6000 8000 10000

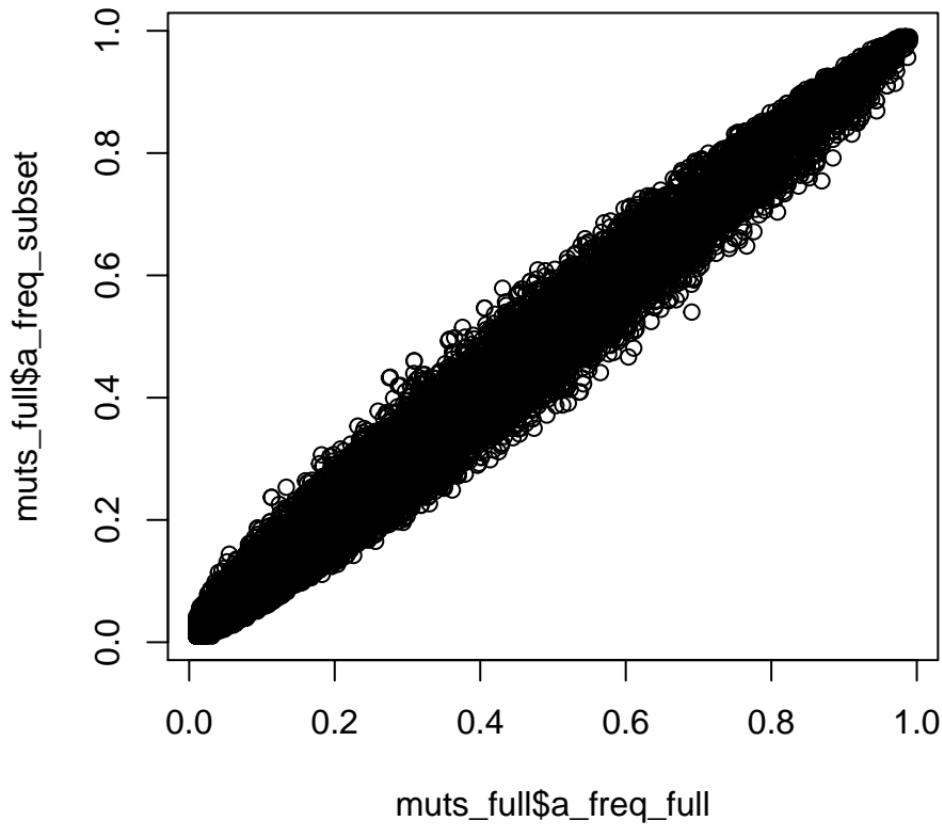
colnames(G\_full\_subset)



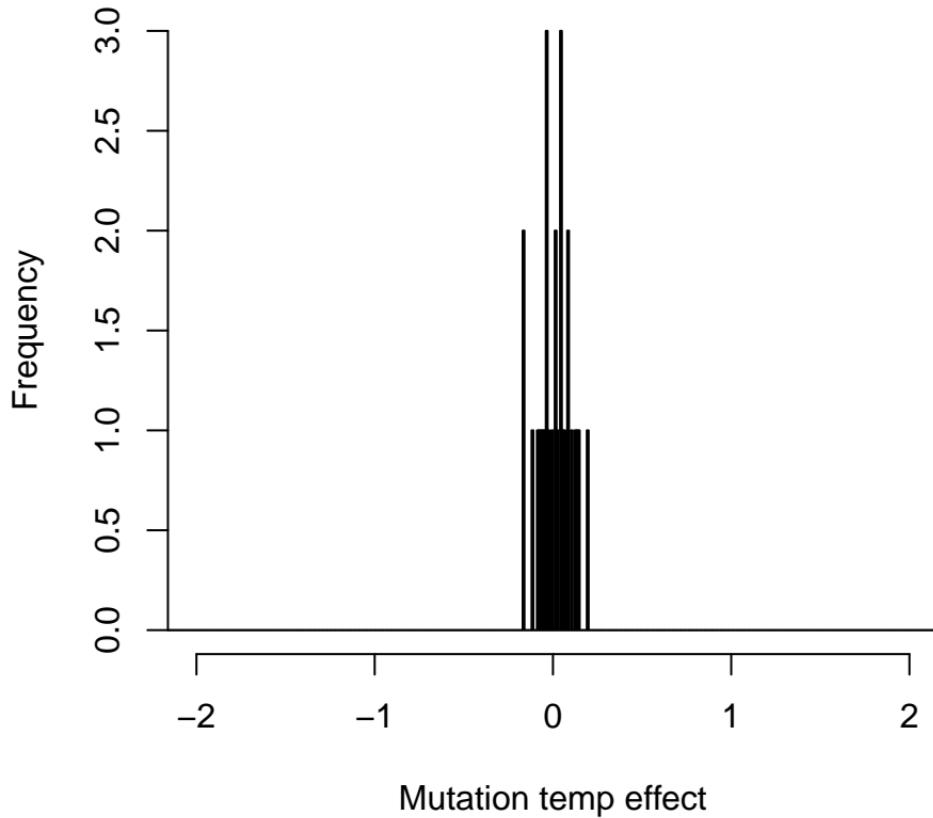


## Histogram of muts\_full\$a\_freq\_subset

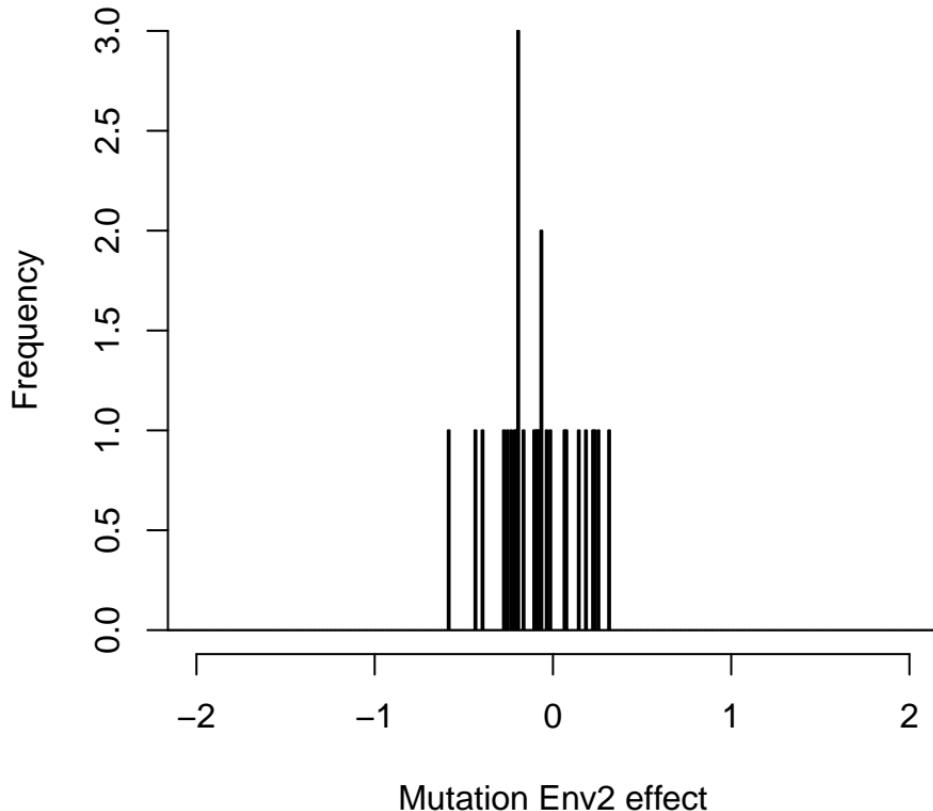




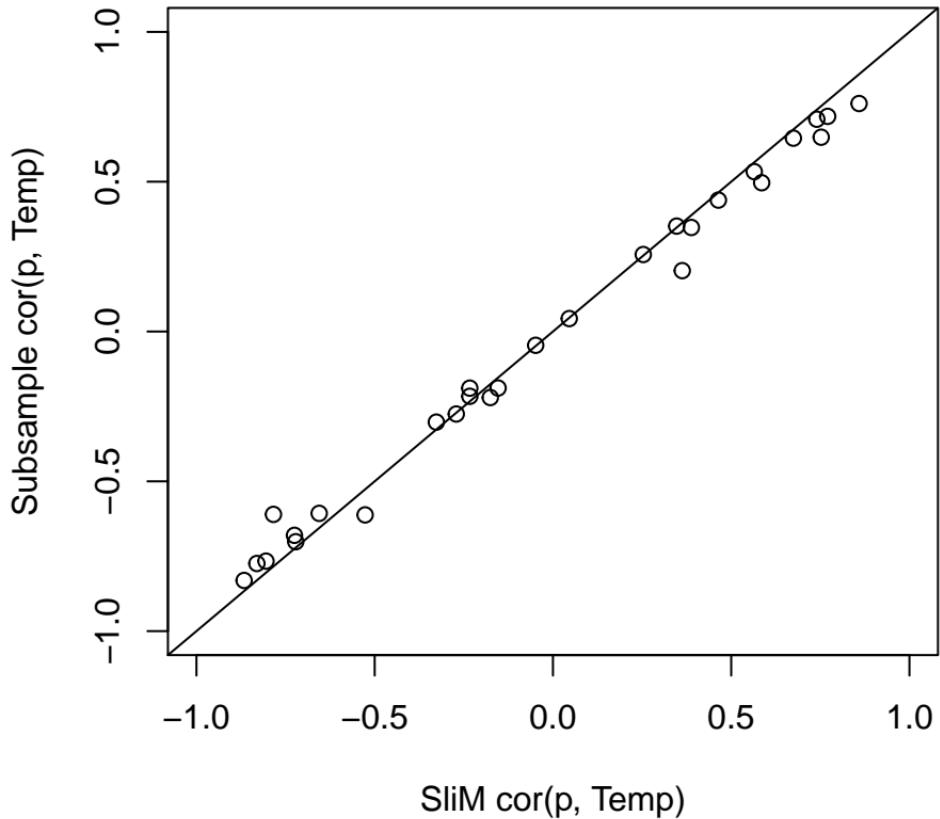
mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1232114



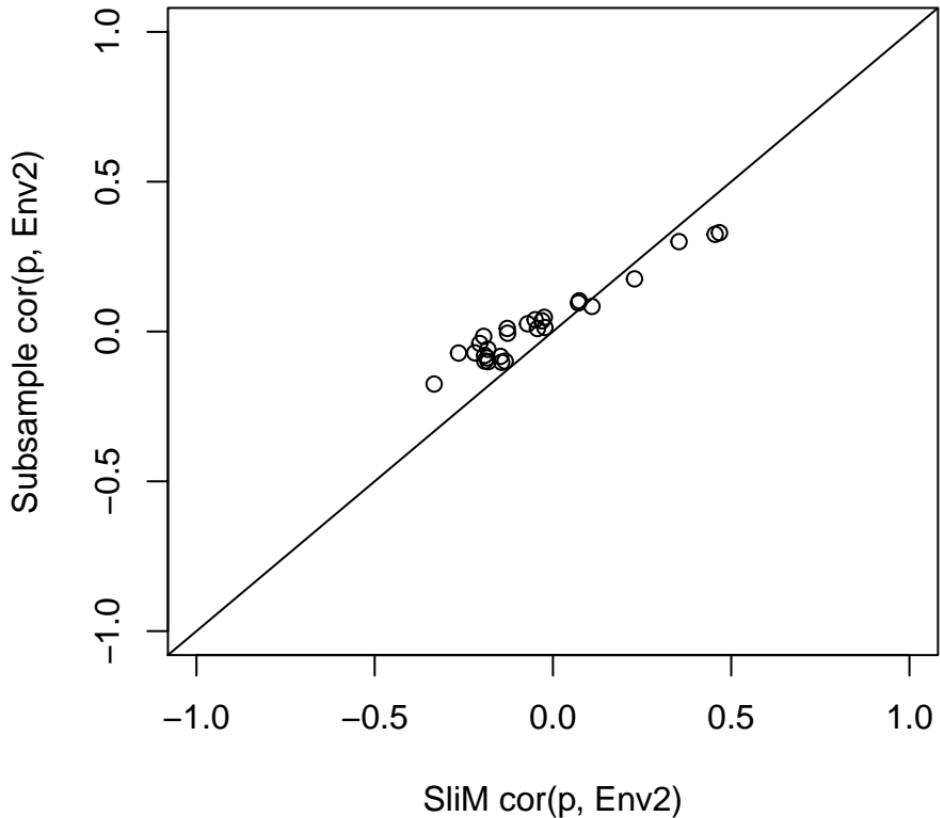
mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1232114



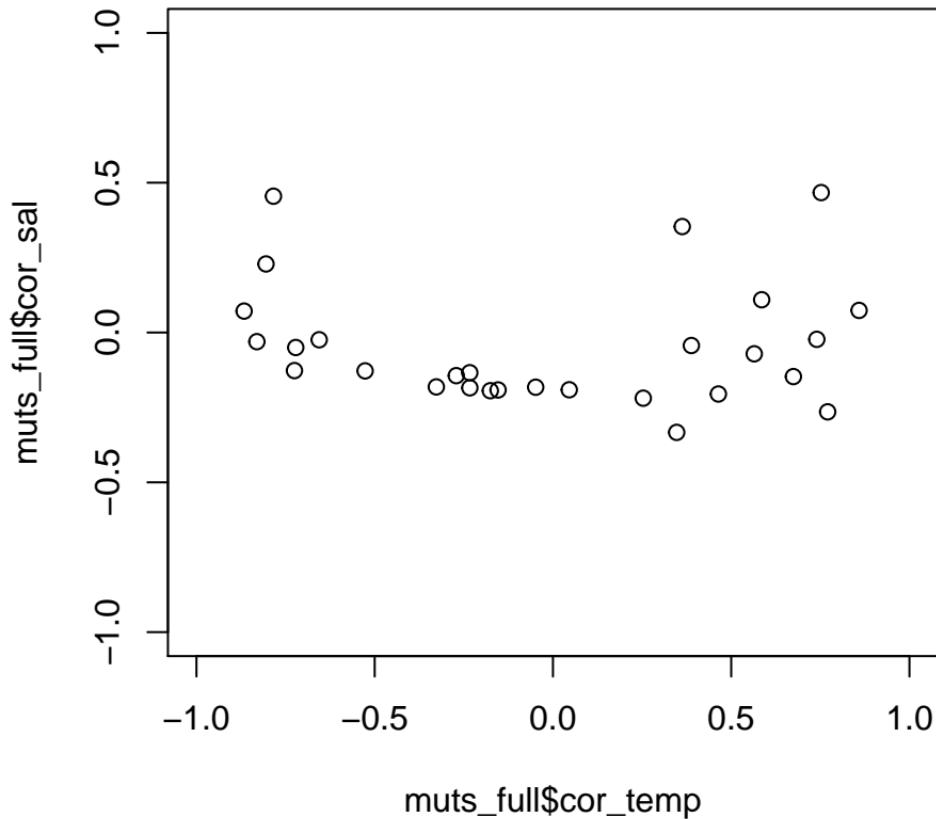
-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center  
1232114



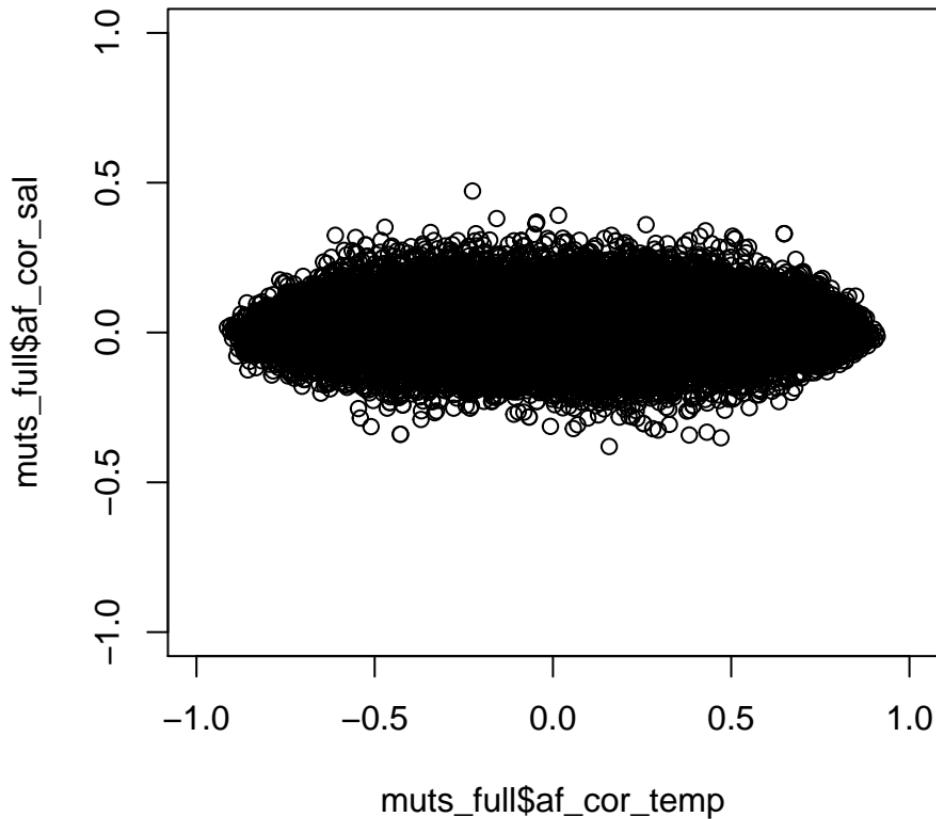
-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center  
1232114



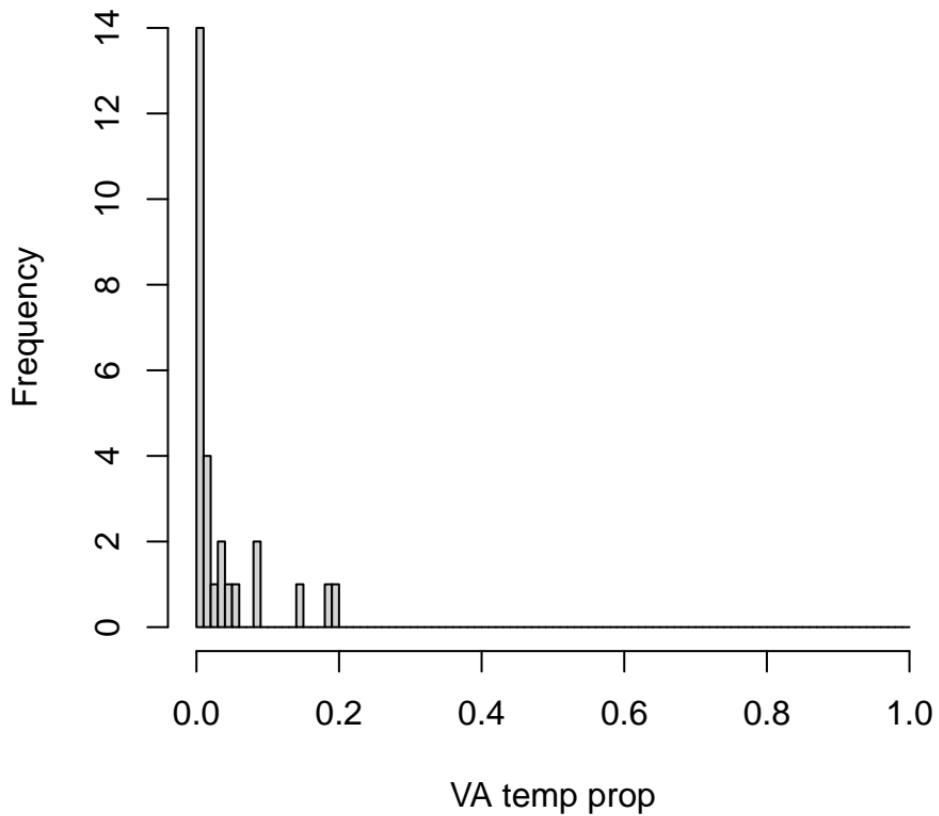
## SLiM Correlation (AF, ENV) all samples



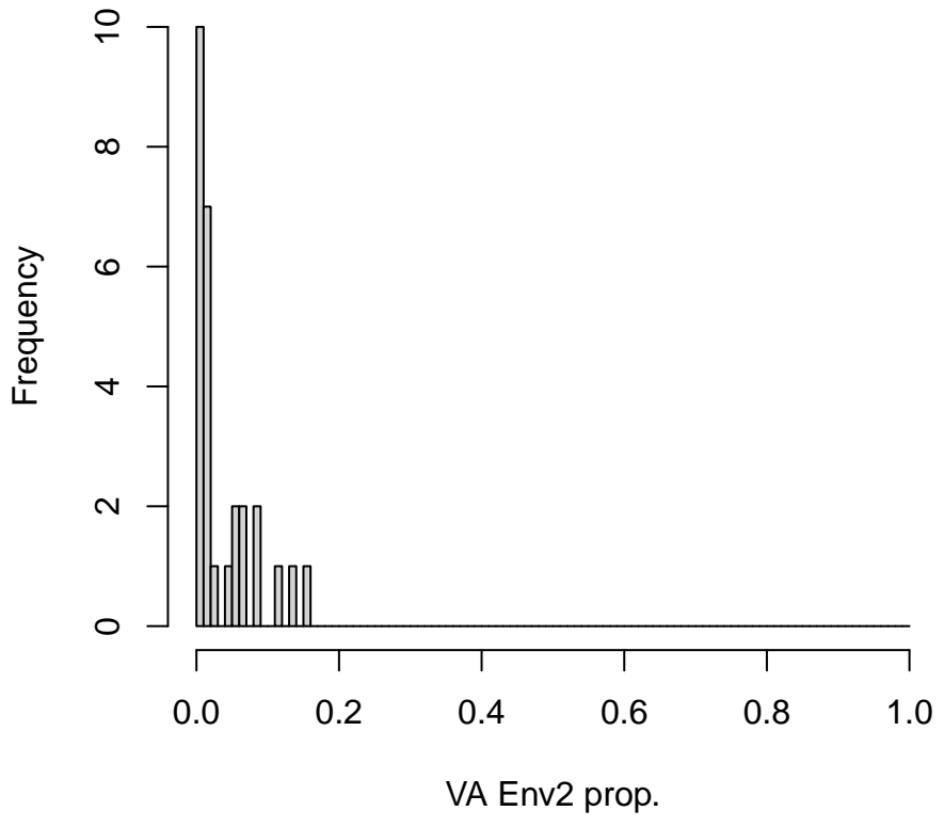
## Subsample Correlation (AF, ENV) all samples

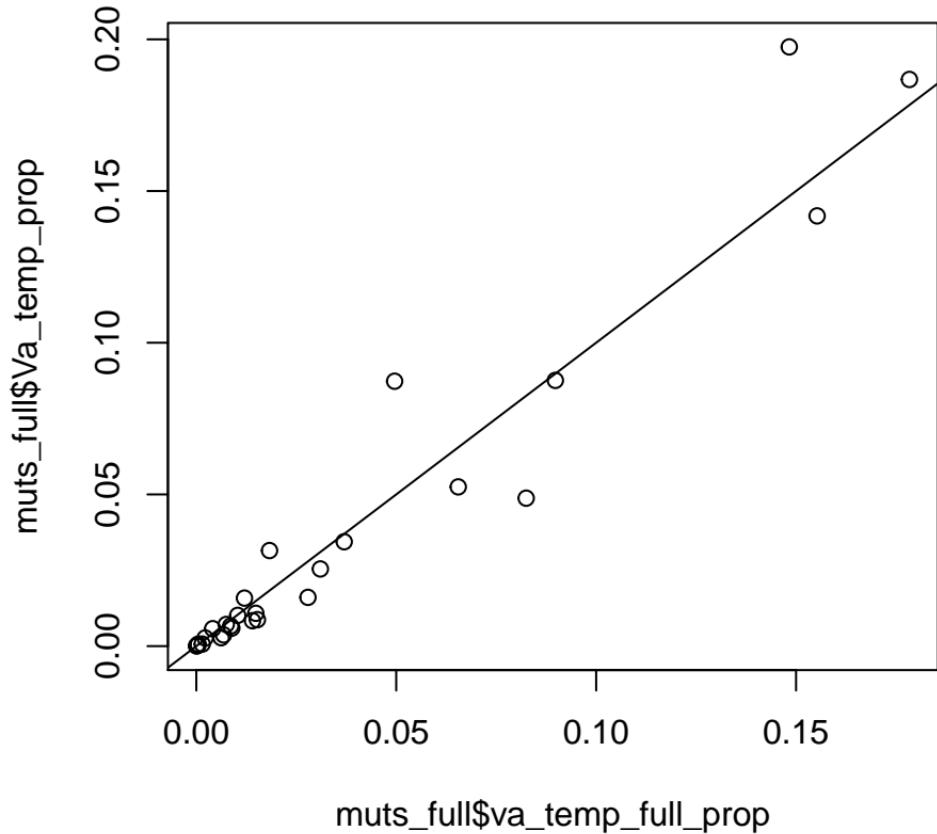


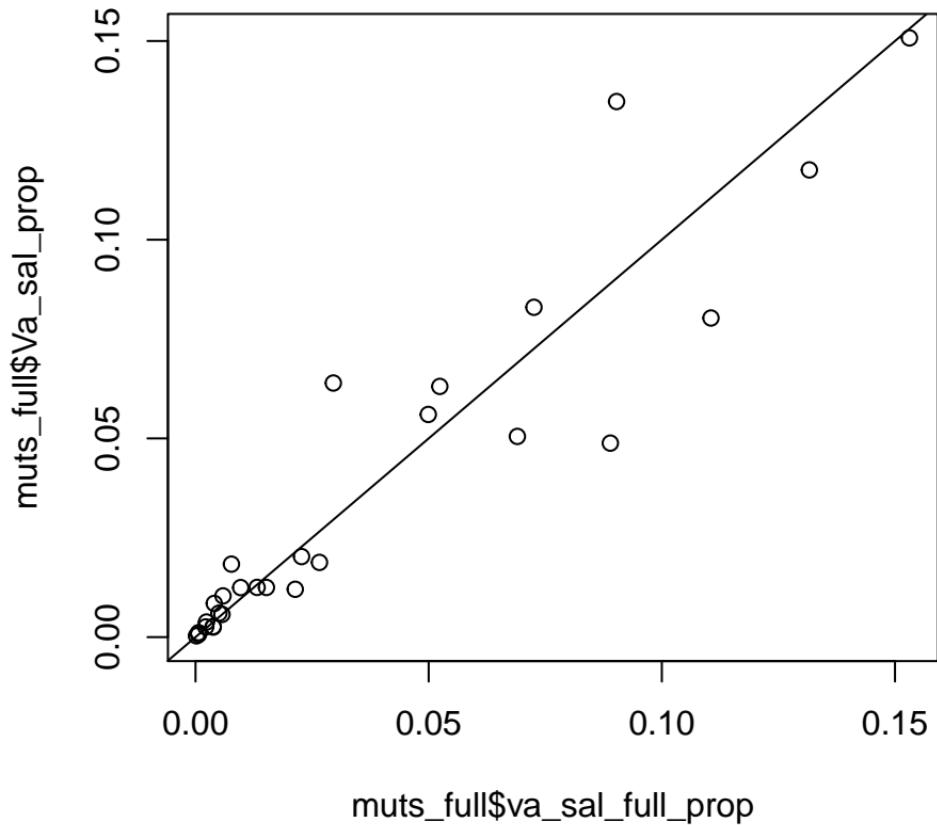
-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center  
1232114



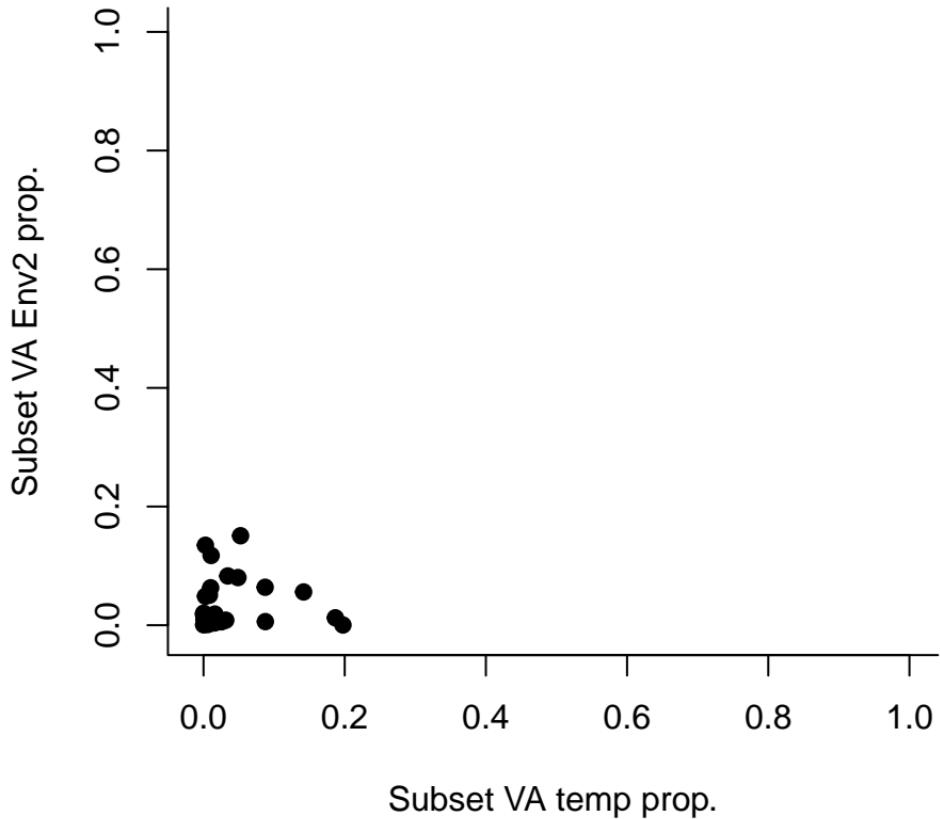
-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center  
1232114



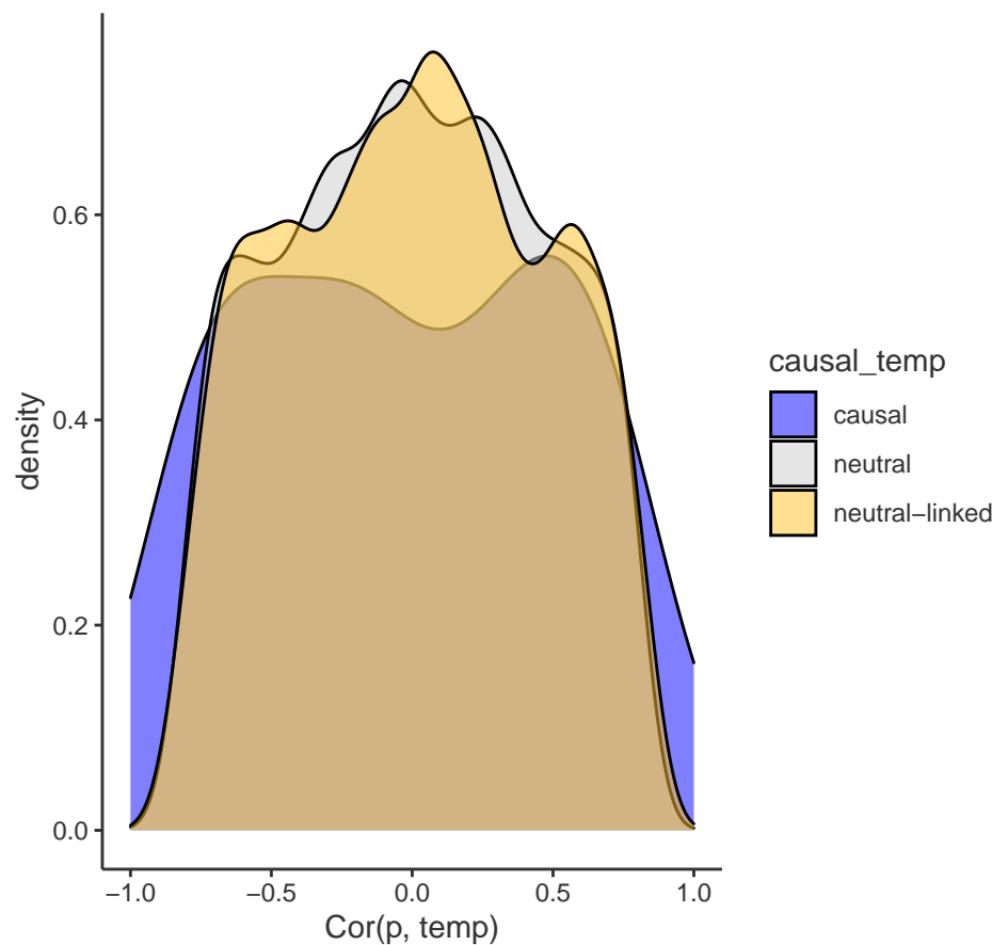




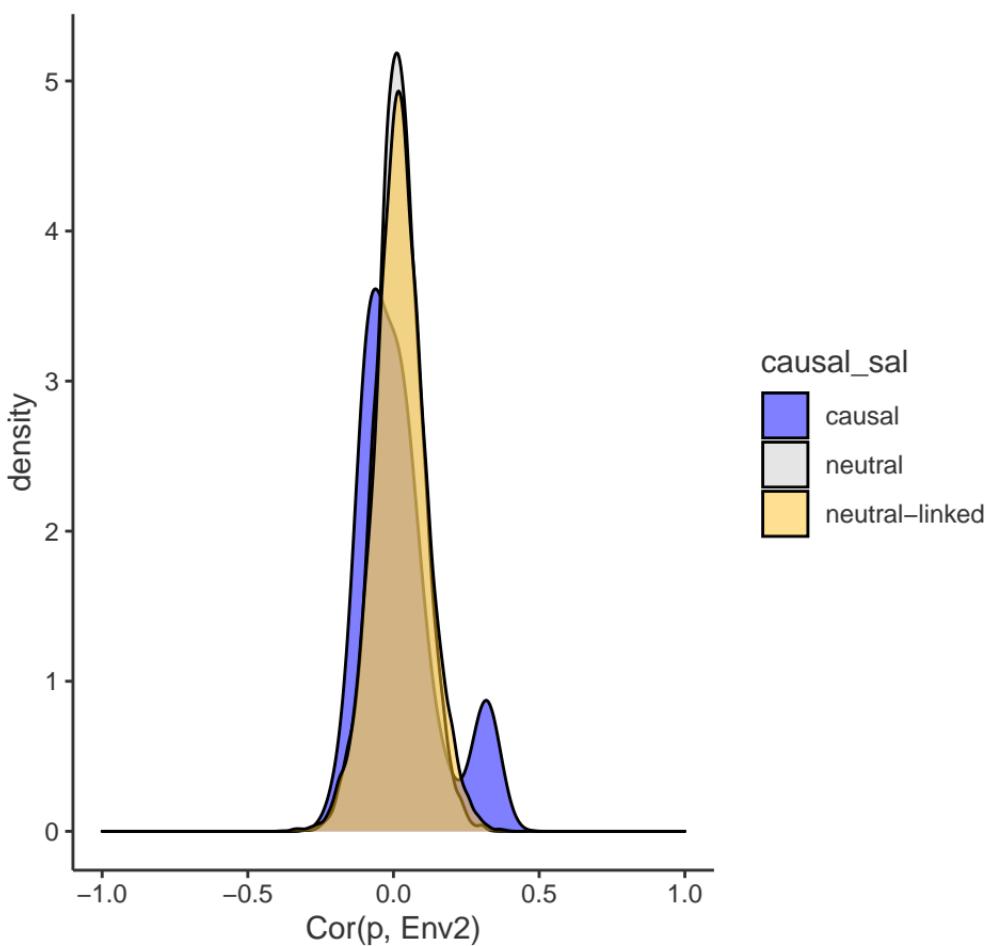
-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center  
1232114



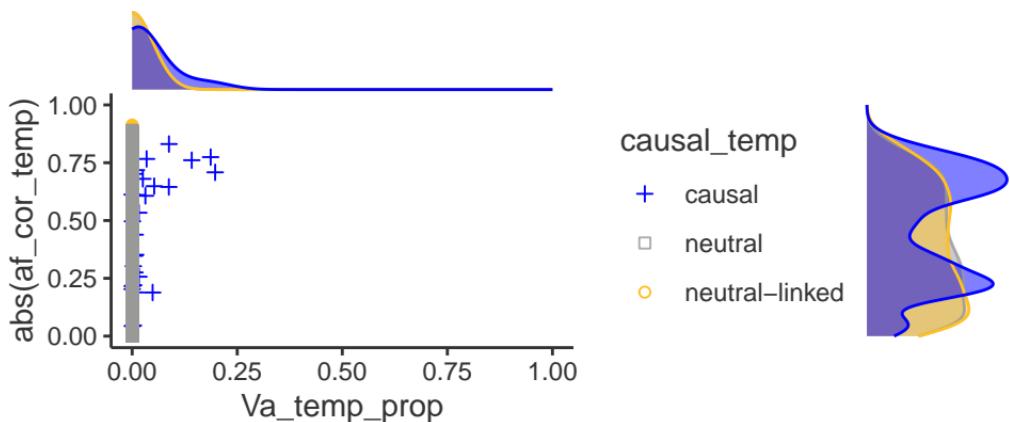
mod-polygenic\_2-trait-pleiotropy-equal-S\_\_Est-Clines\_N-cline-center-to-edge\_m-cons  
1232114



mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-cline-center-to-edge\_m-consta  
1232114



mod-polygenic\_2-trait-pleiotropy-equal-S\_\_  
1232114



mod-polygenic\_2-trait-pleiotropy-equal-S\_\_  
1232114

