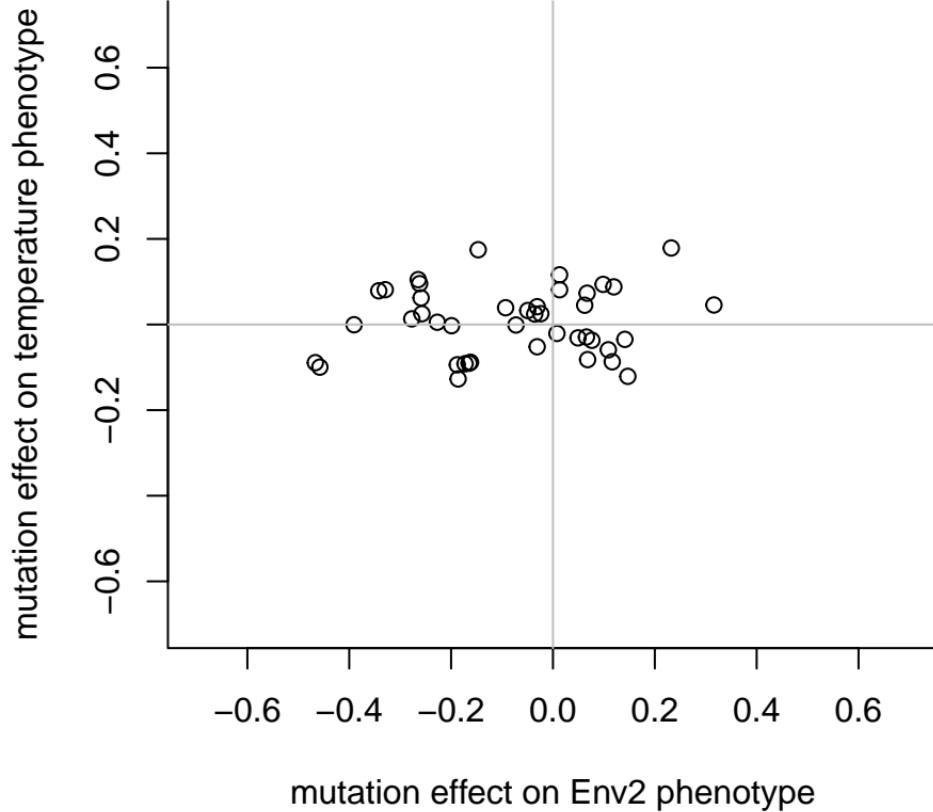
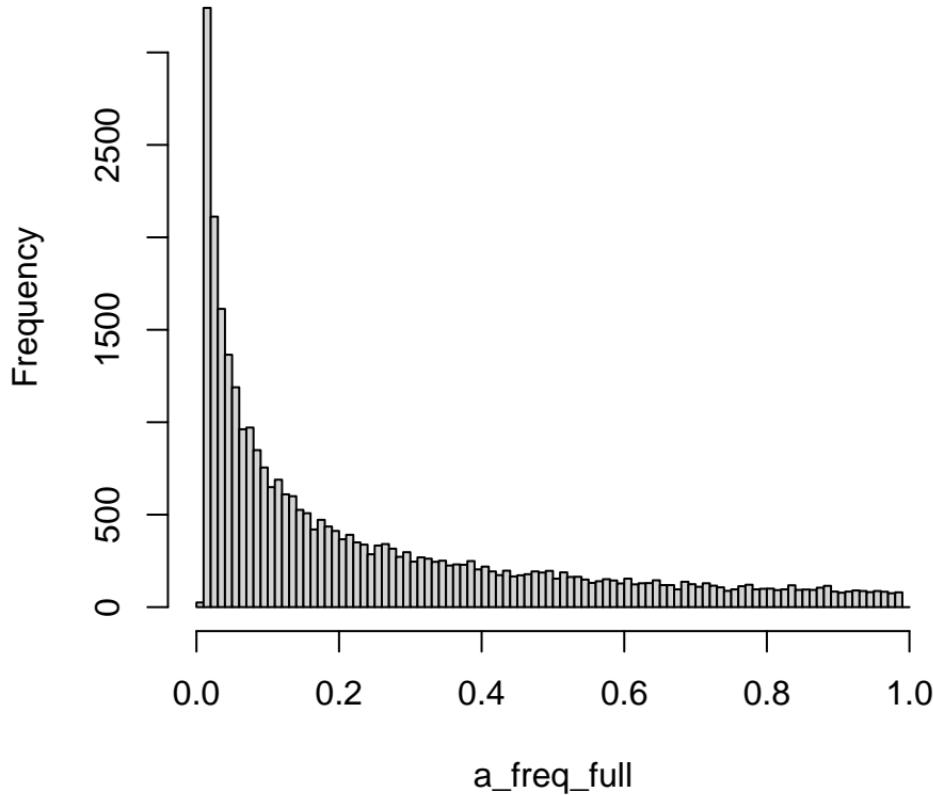


mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equal\_m-constant  
1232117



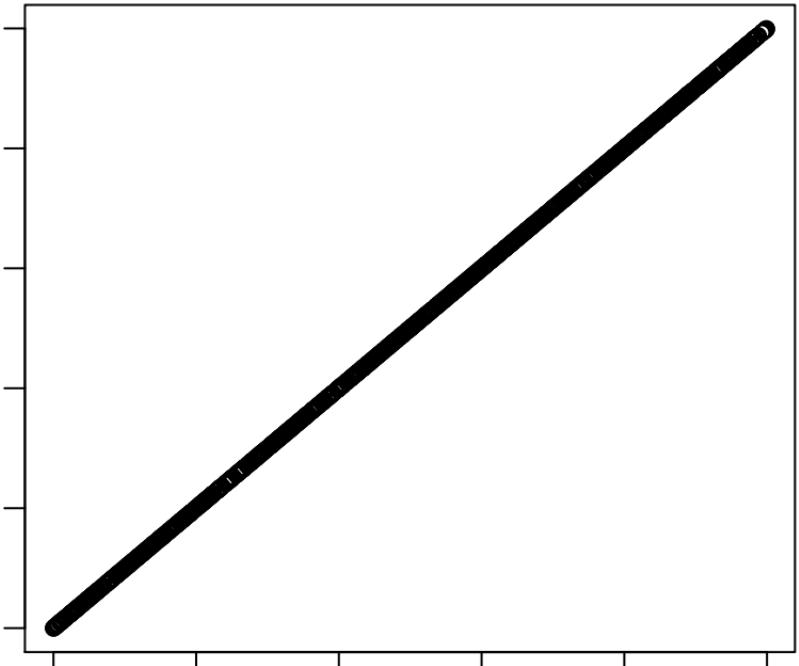
## Histogram of a\_freq\_full

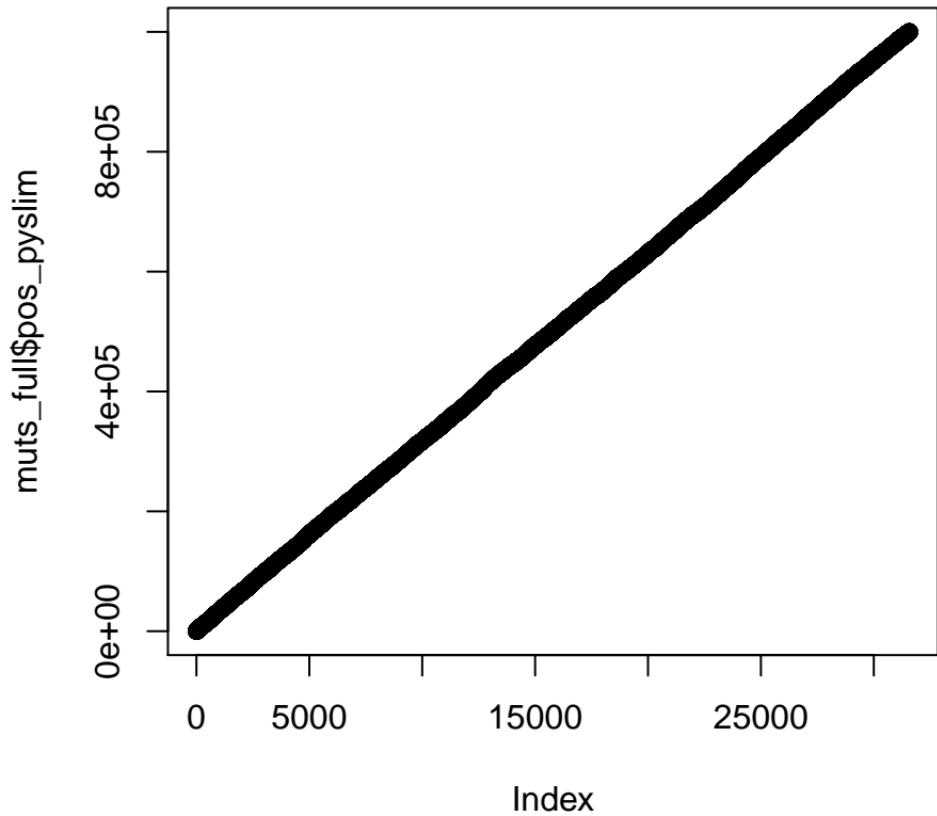


indPhen\_df\$indID[which(indPhen\_df\$subset)]

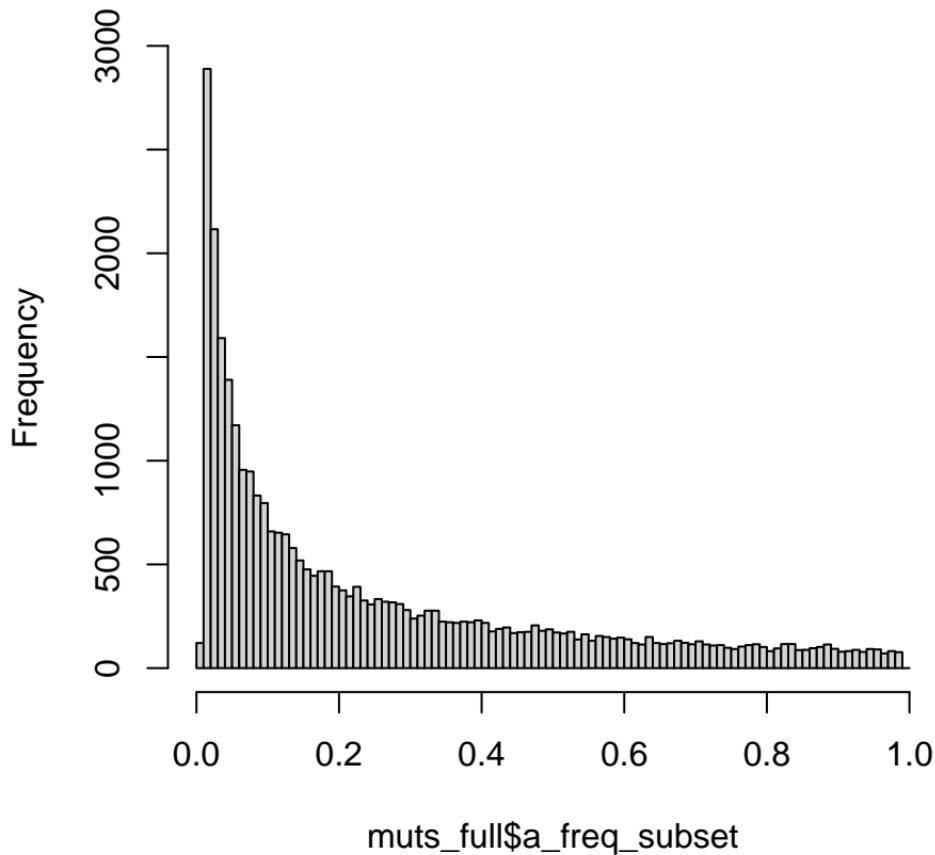
0 2000 4000 6000 8000 10000

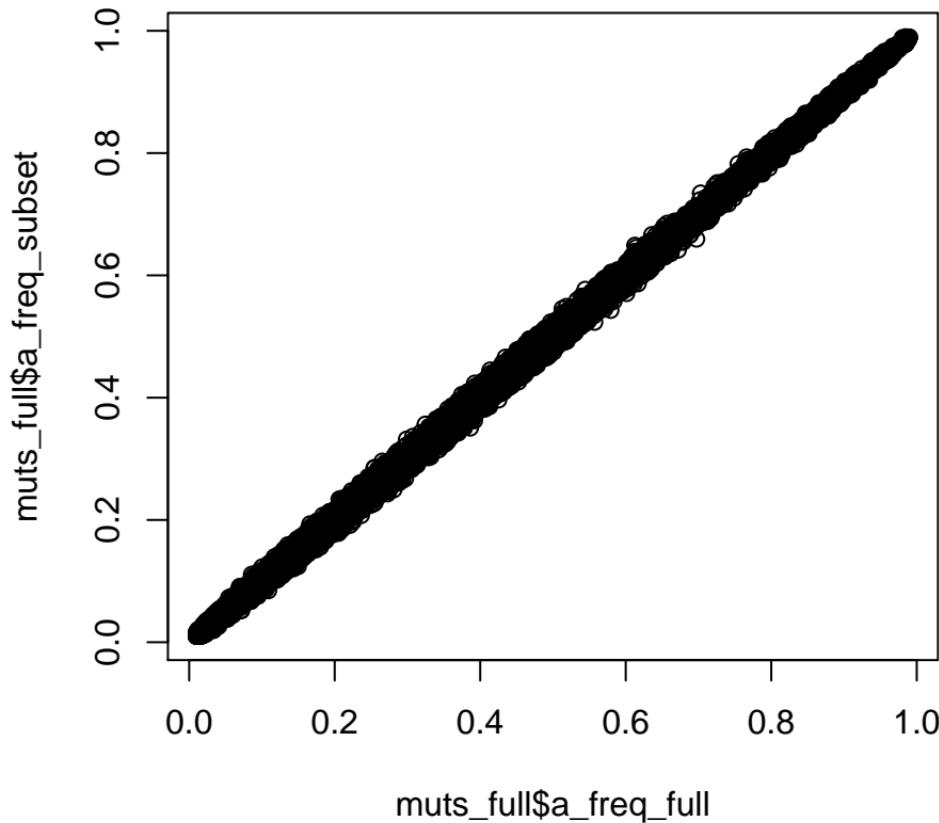
colnames(G\_full\_subset)



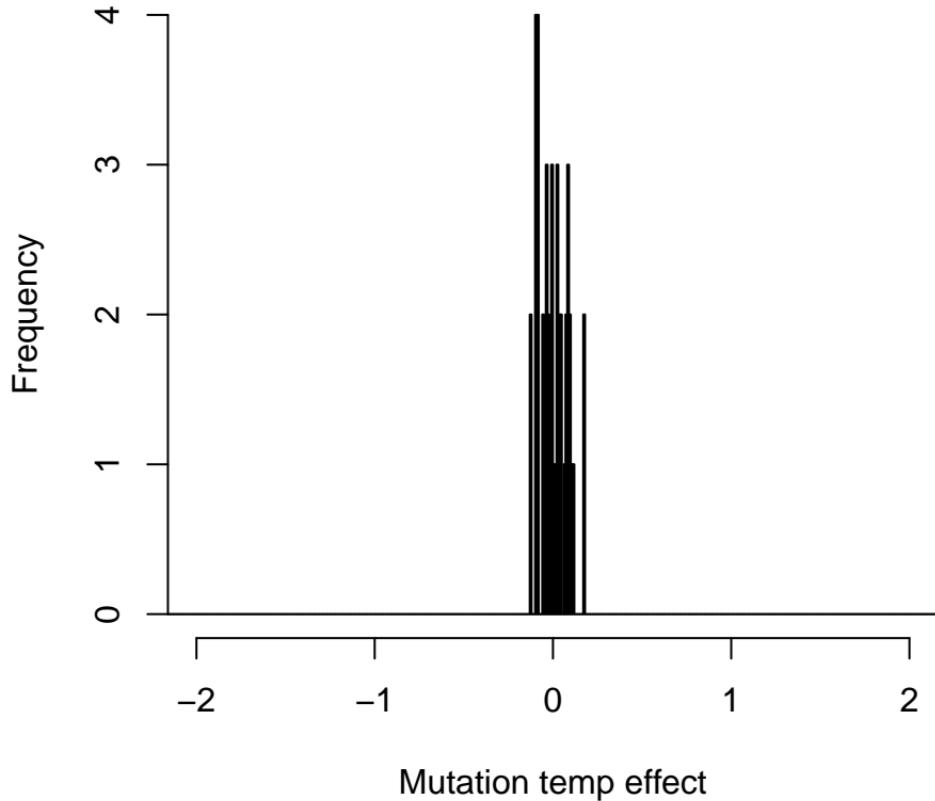


## Histogram of muts\_full\$a\_freq\_subset

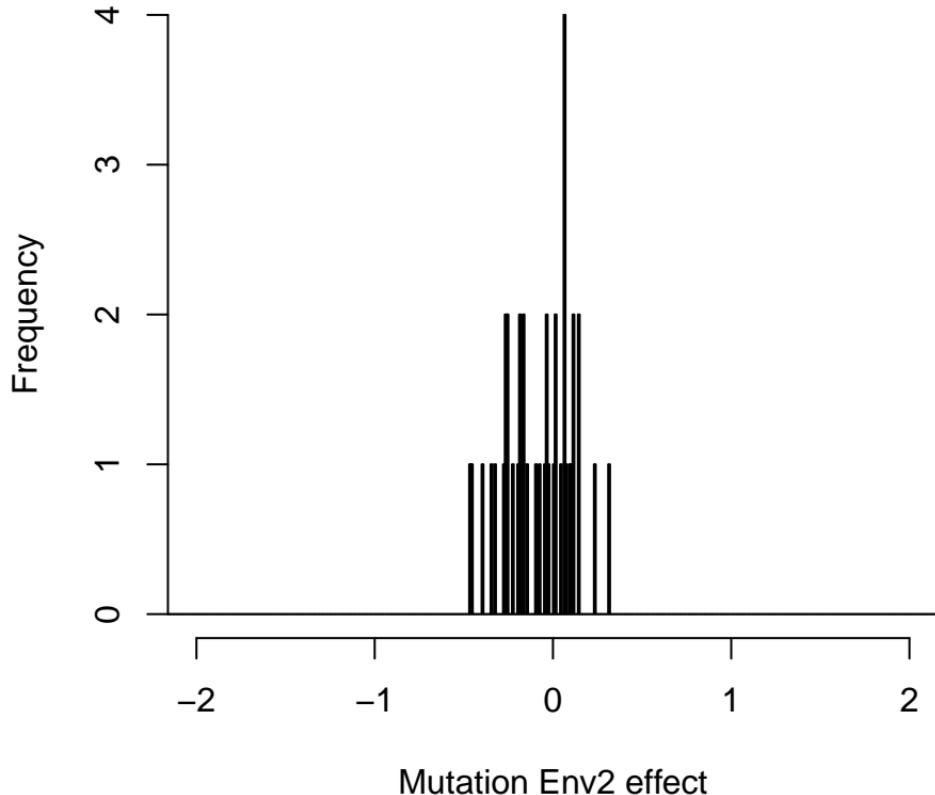




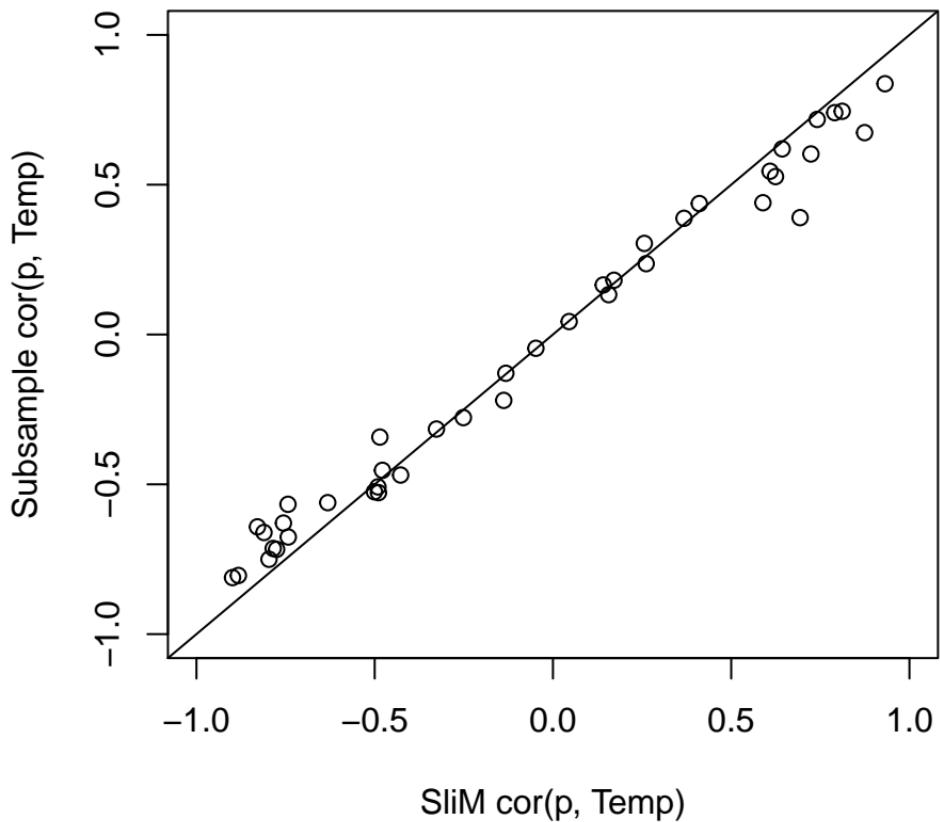
mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equal\_m-constant  
1232117



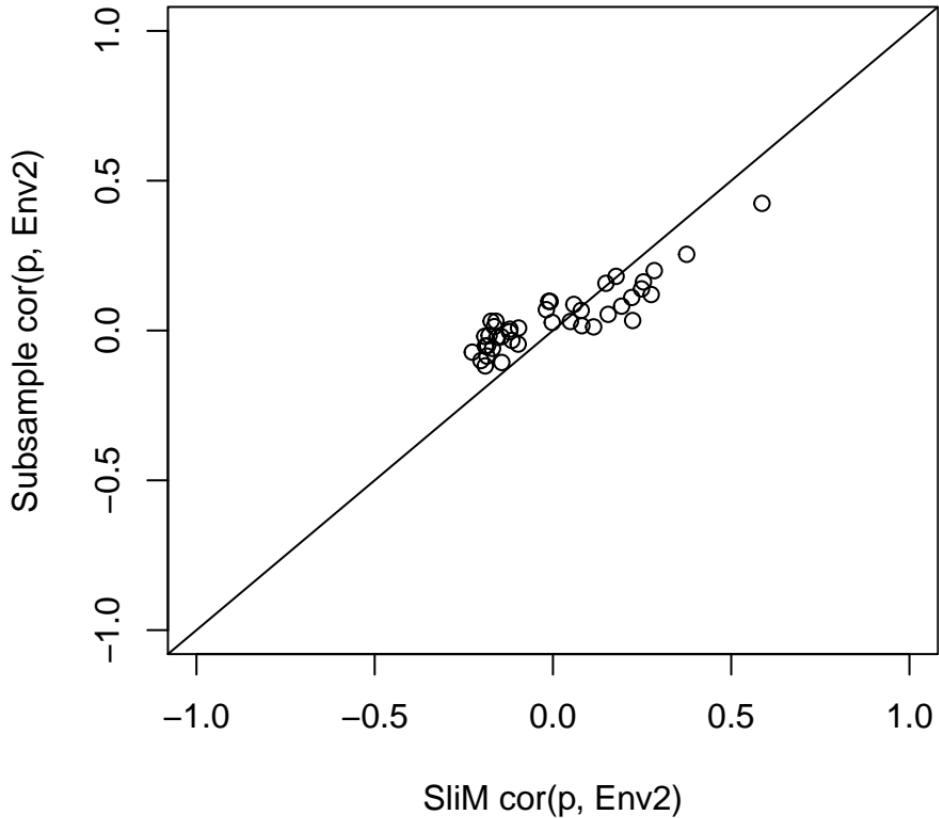
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1232117



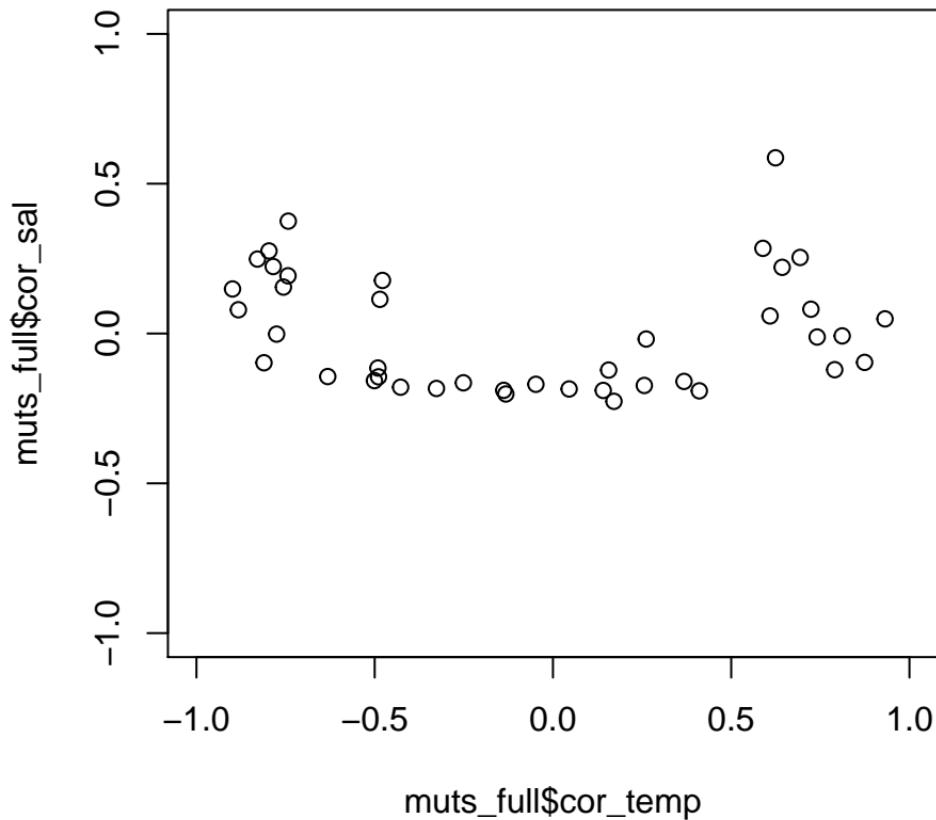
genic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equ  
1232117



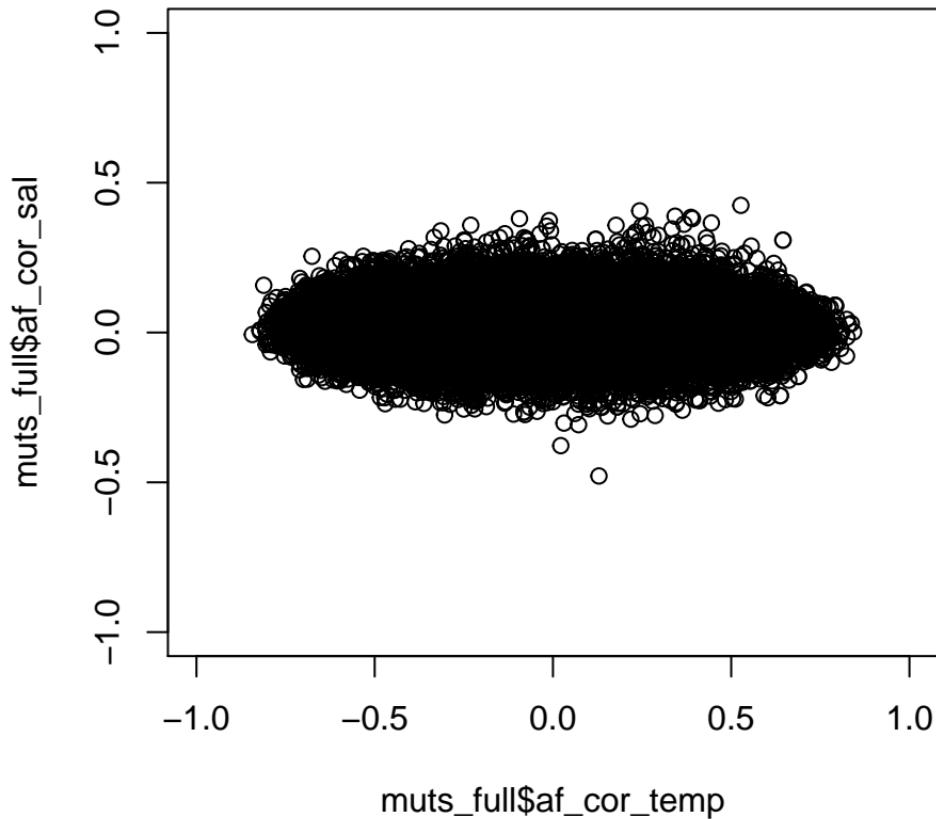
genic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equa  
1232117



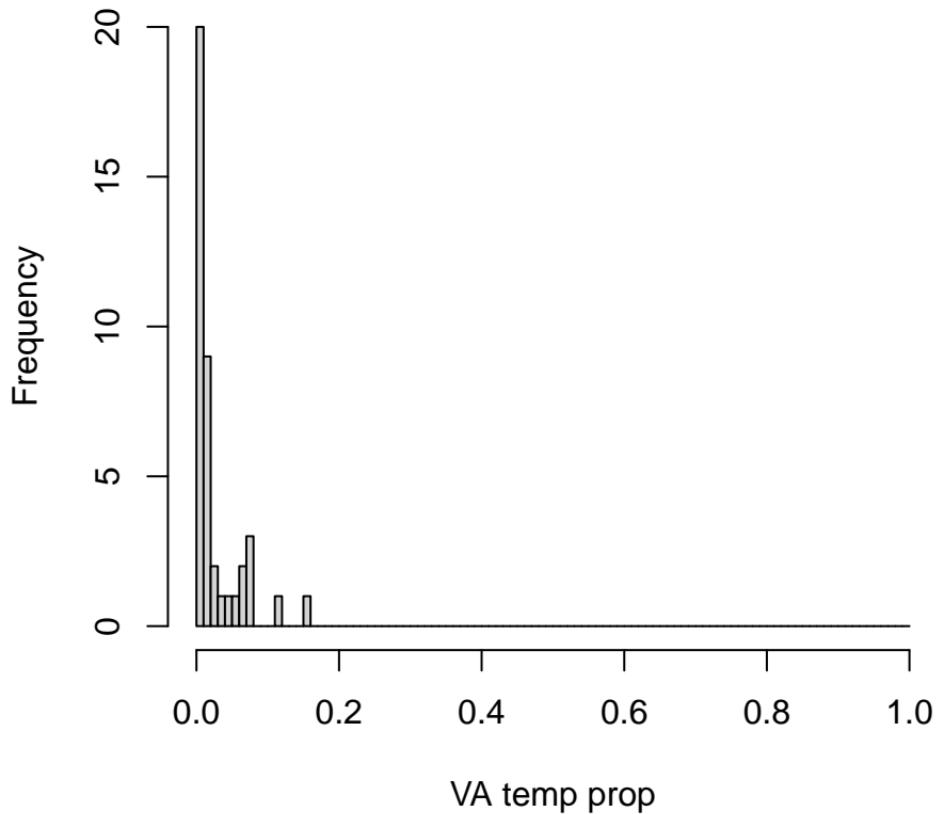
## SiM Correlation (AF, ENV) all samples



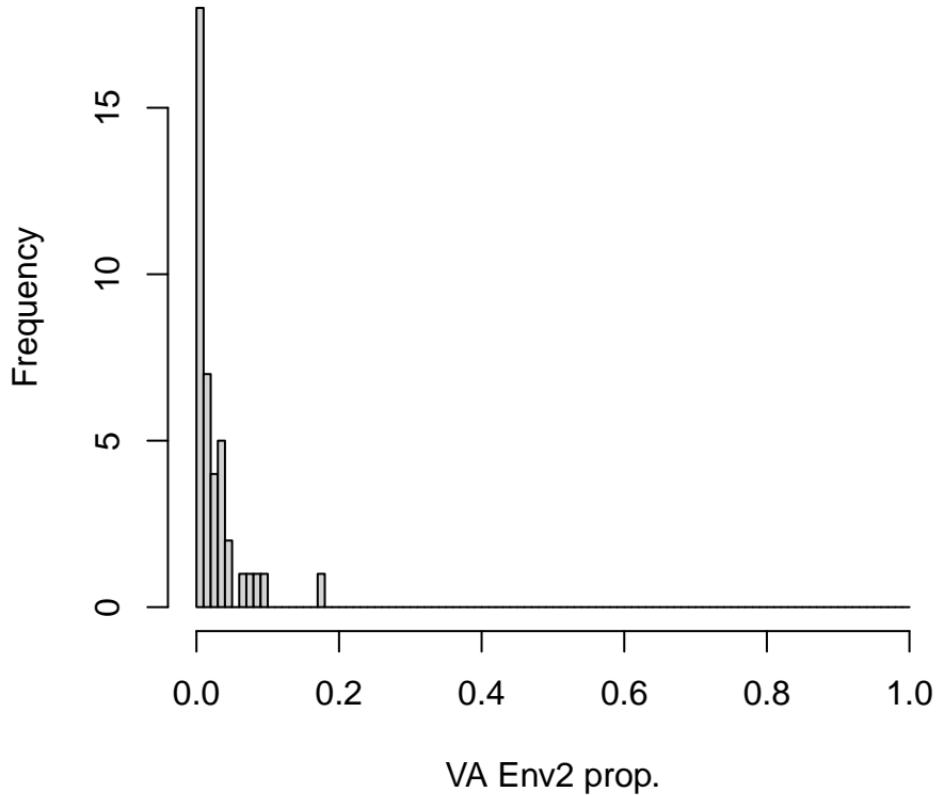
## Subsample Correlation (AF, ENV) all samples

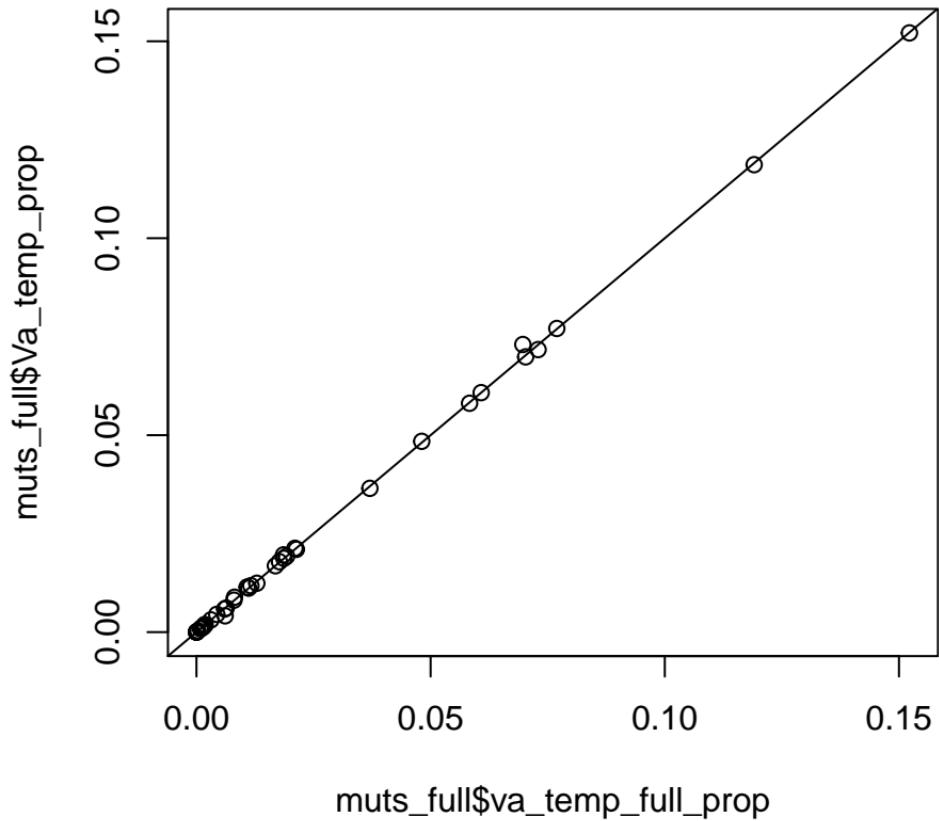


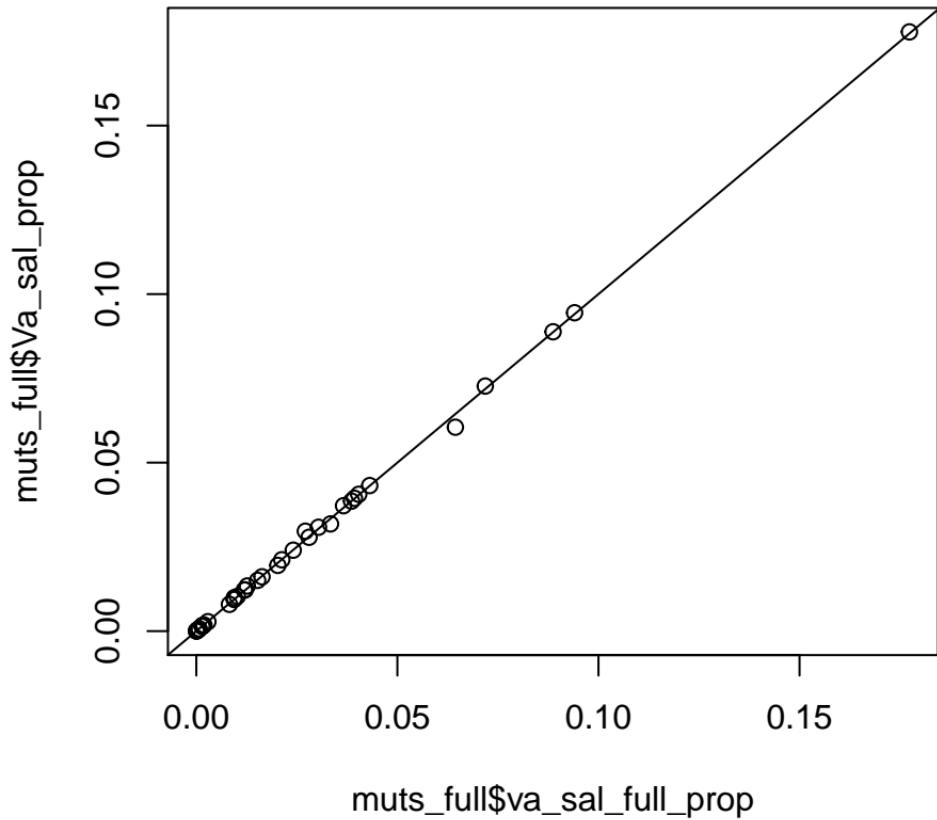
genic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equa  
1232117



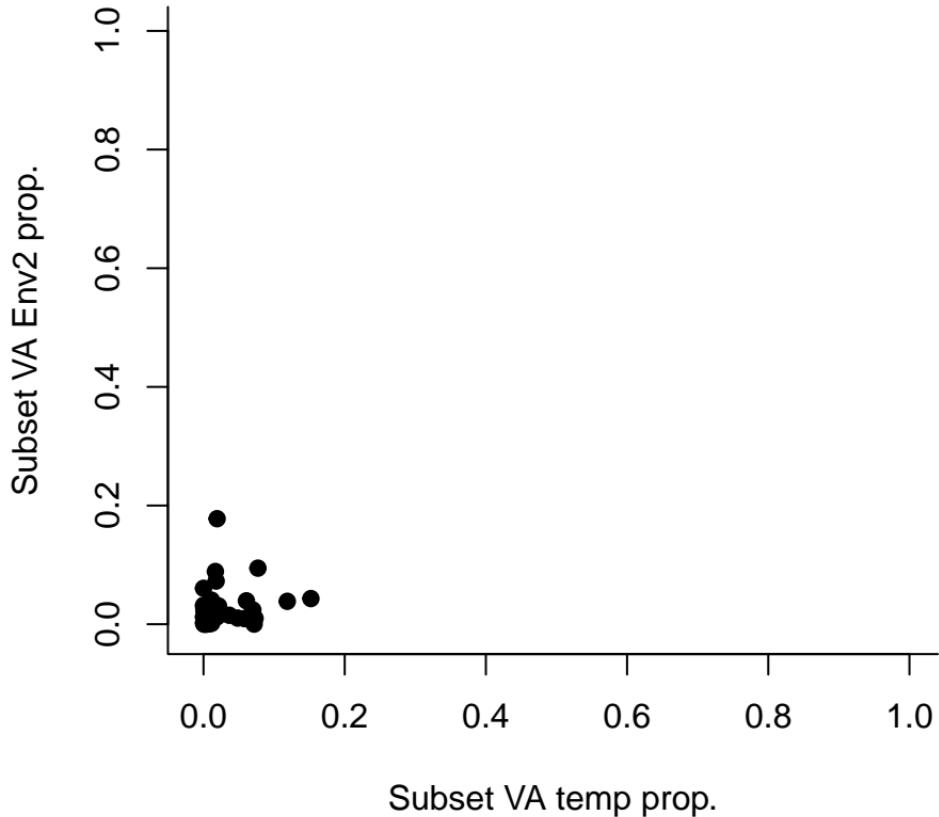
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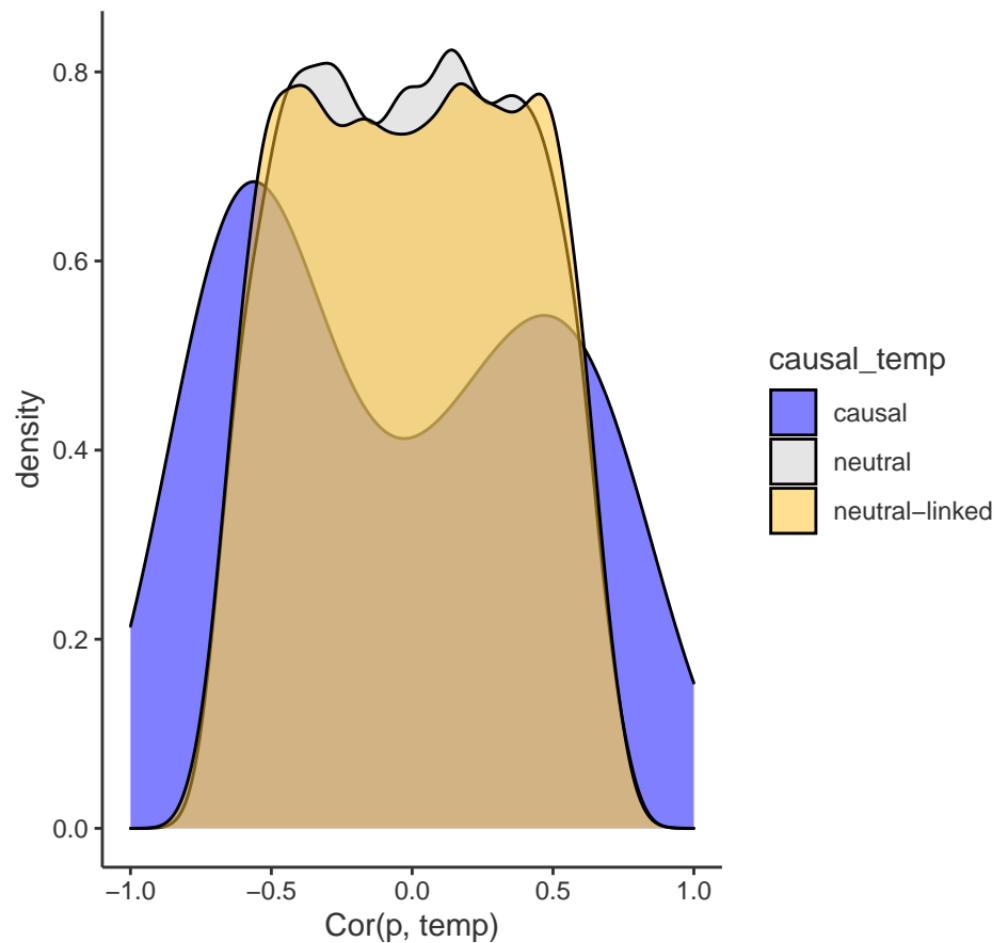




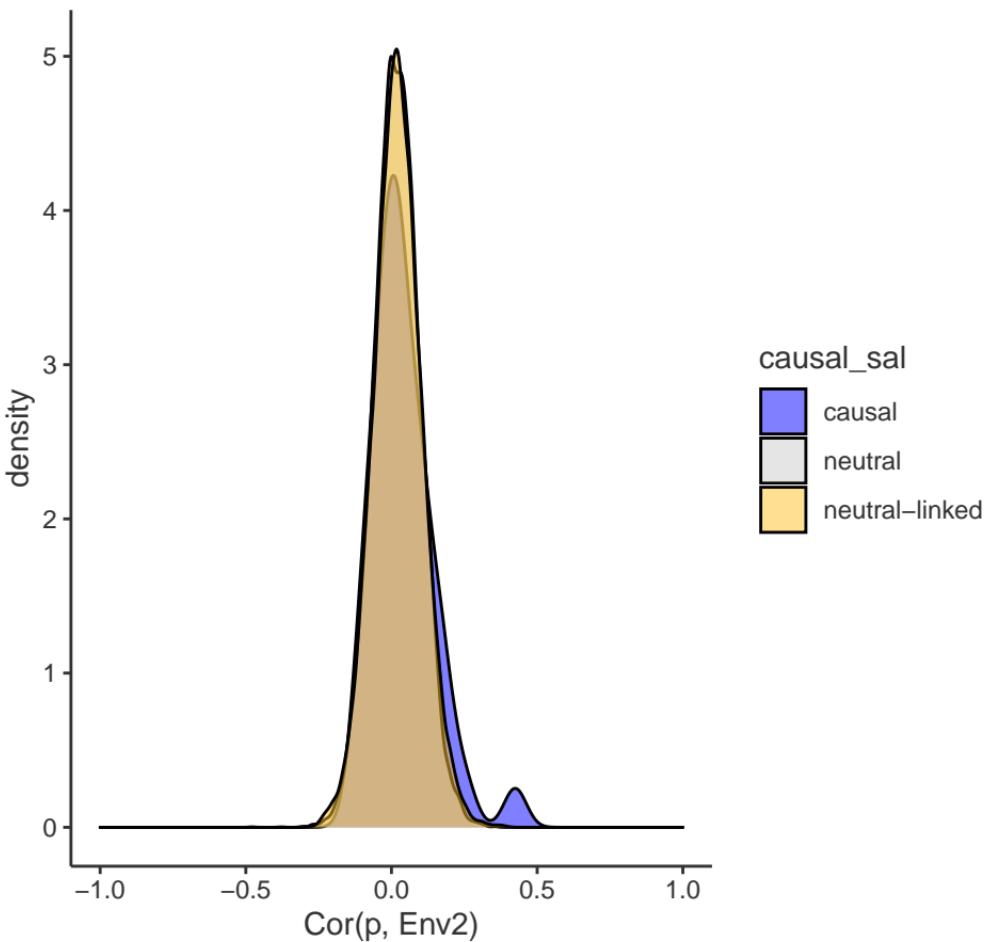
genic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equ  
1232117



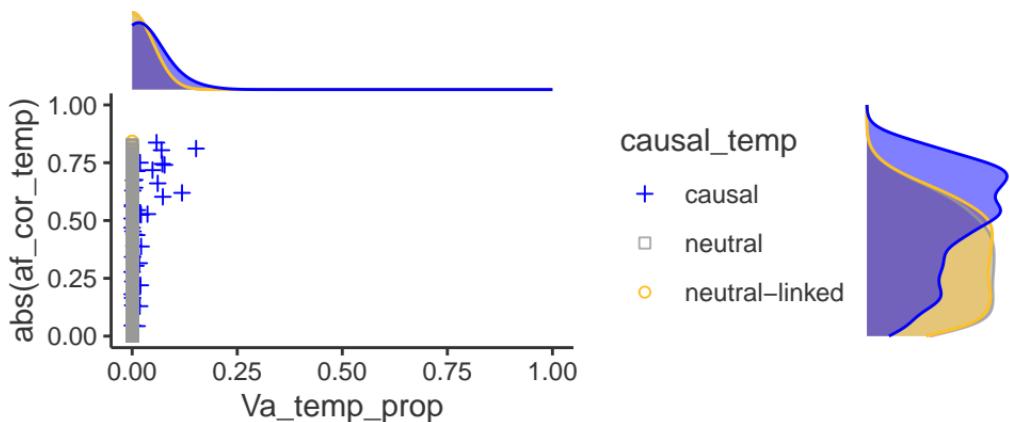
mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equal\_m-constant  
1232117



mod-polygenic\_2-trait-pleiotropy-equal-S\_Est-Clines\_N-equal\_m-constant  
1232117



mod-polygenic\_2-trait-pleiotropy-equal-S\_\_  
1232117



mod-polygenic\_2-trait-pleiotropy-equal-S\_\_  
1232117

