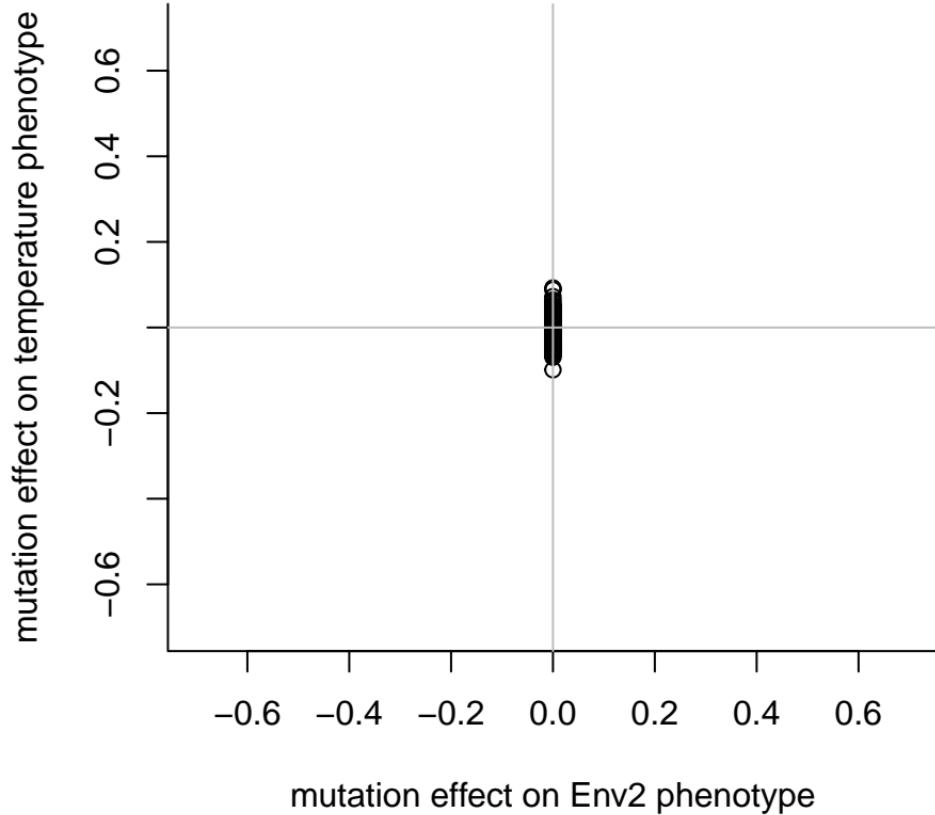
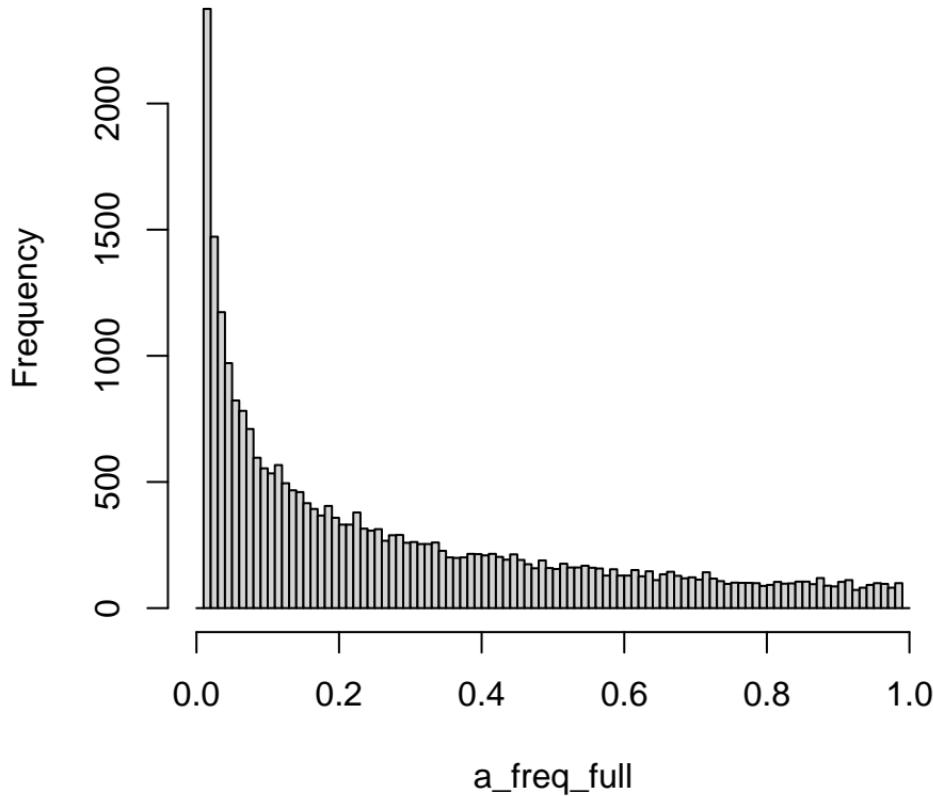


highly-polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1231094



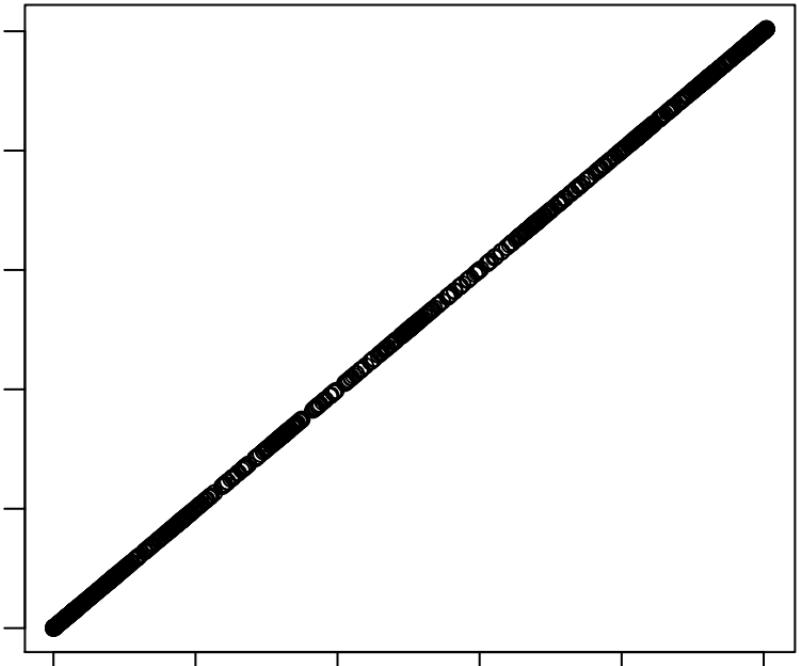
# Histogram of a\_freq\_full

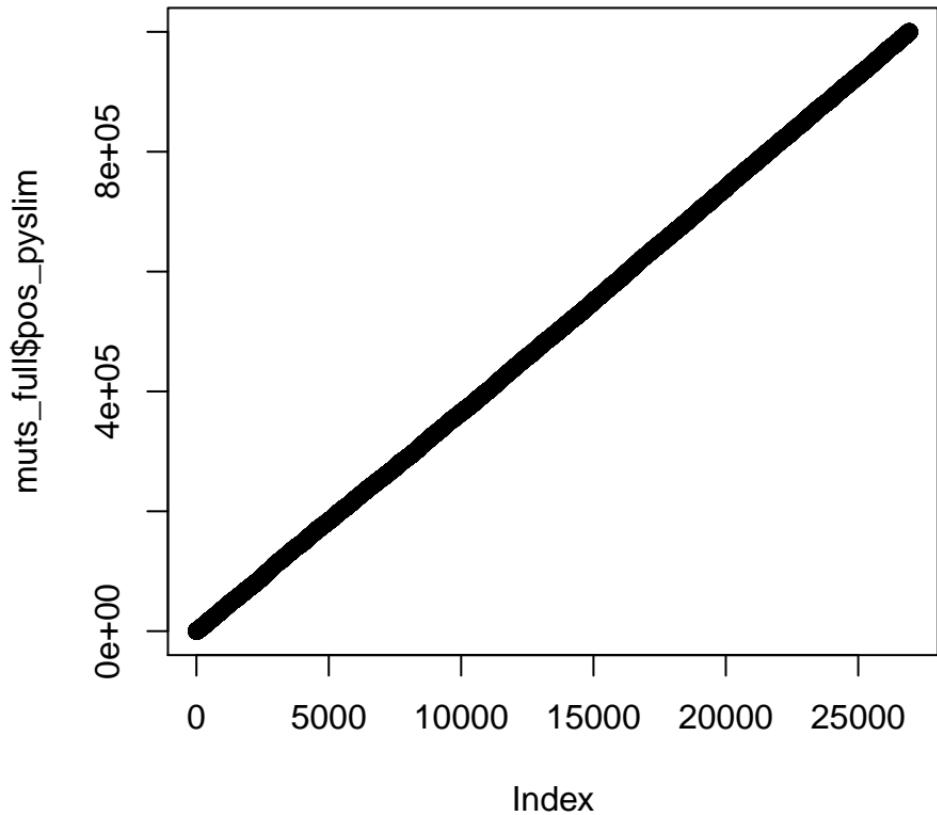


indPhen\_df\$indID[which(indPhen\_df\$subset)]

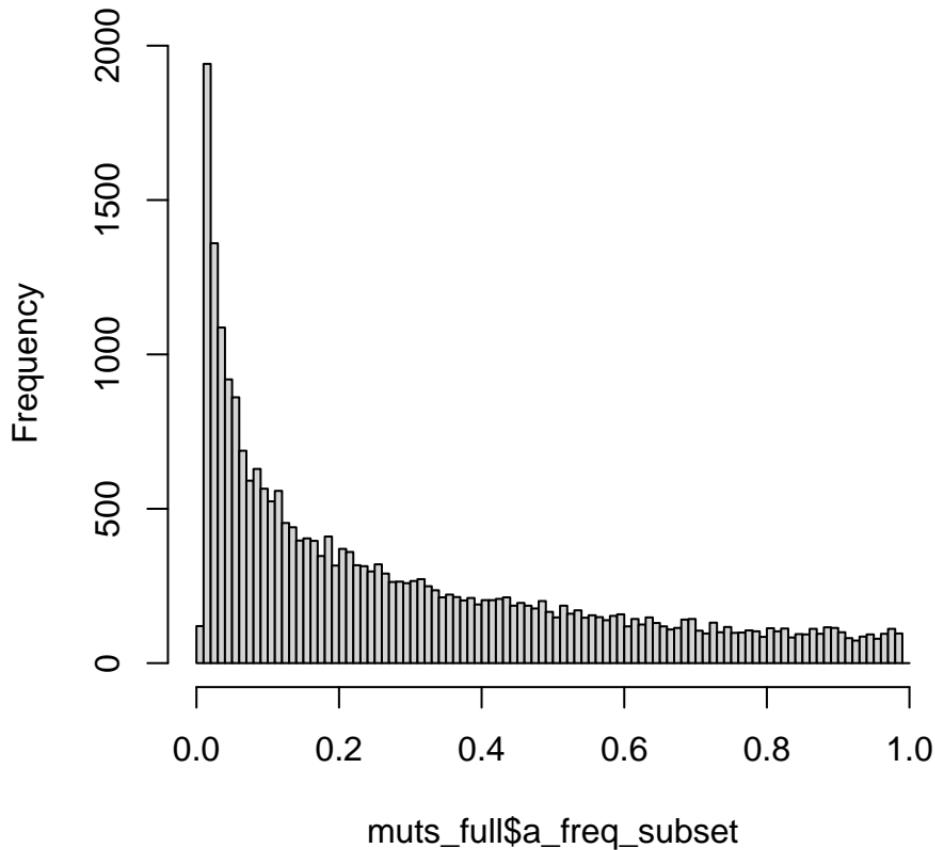
0 2000 4000 6000 8000 10000

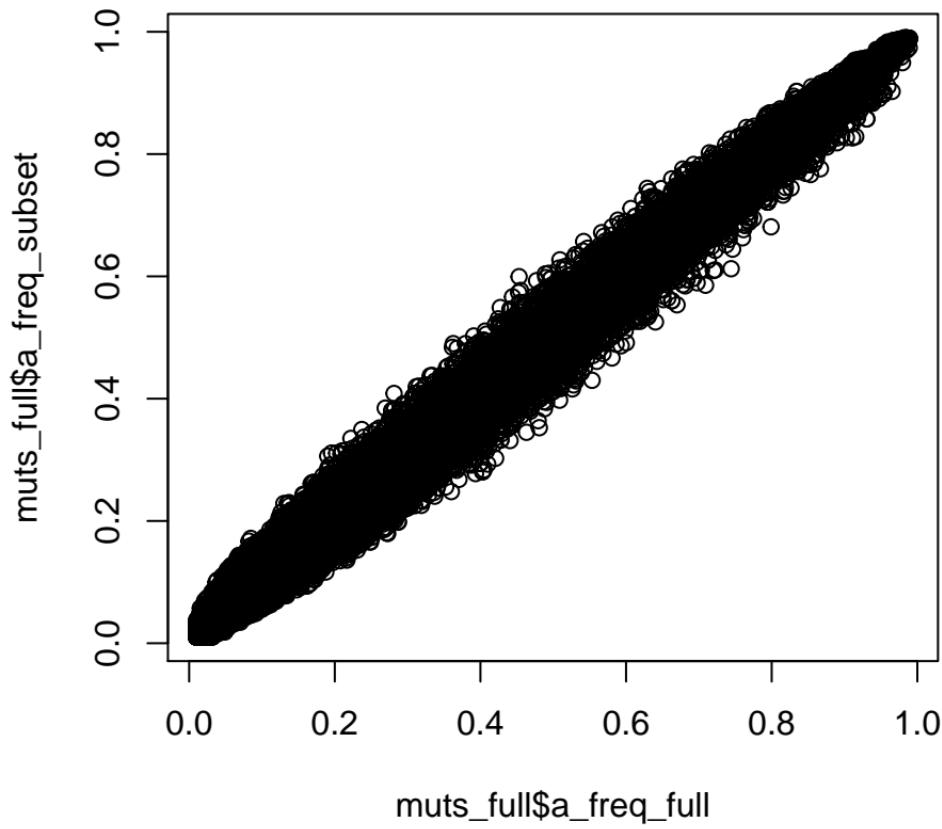
colnames(G\_full\_subset)



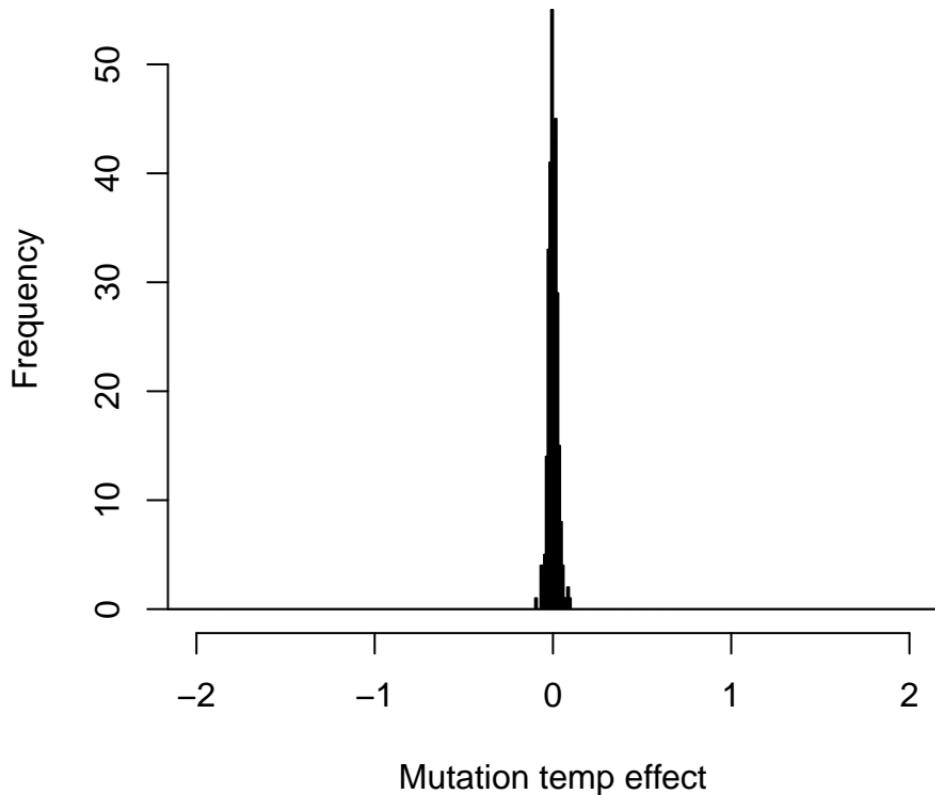


## Histogram of muts\_full\$a\_freq\_subset

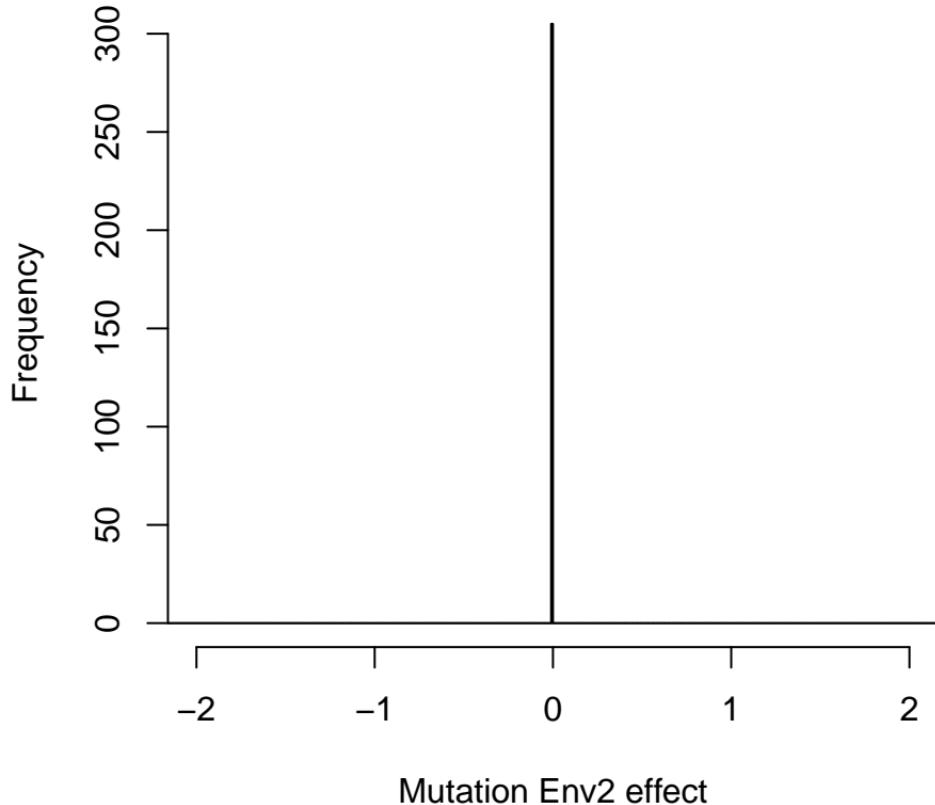




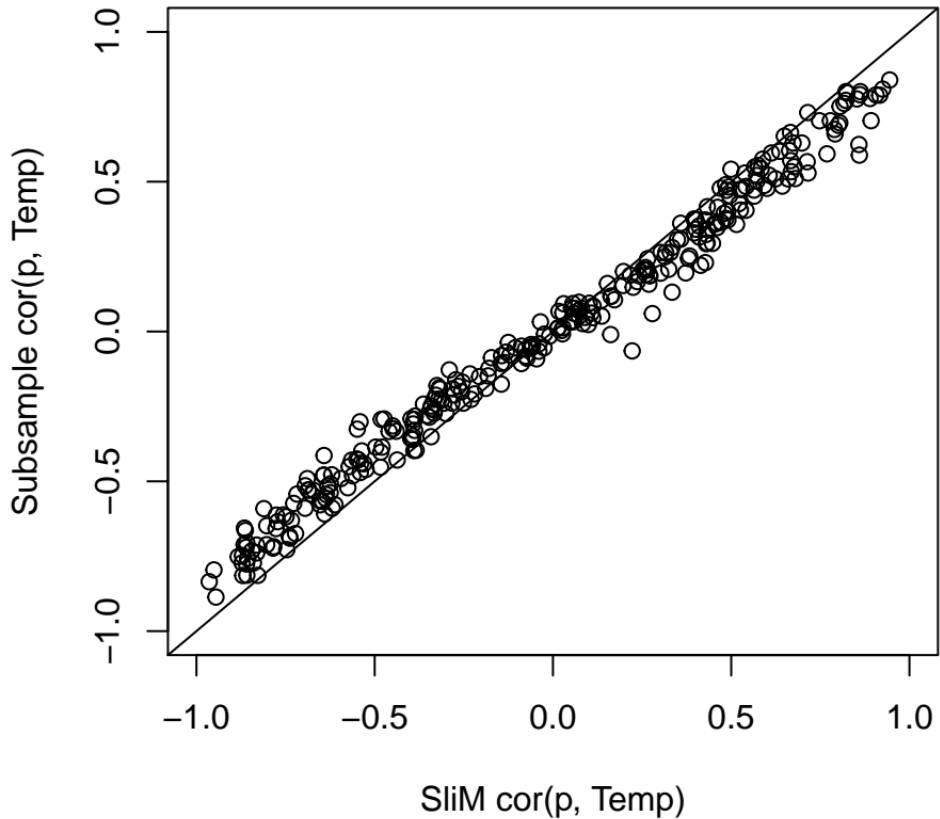
highly-polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1231094



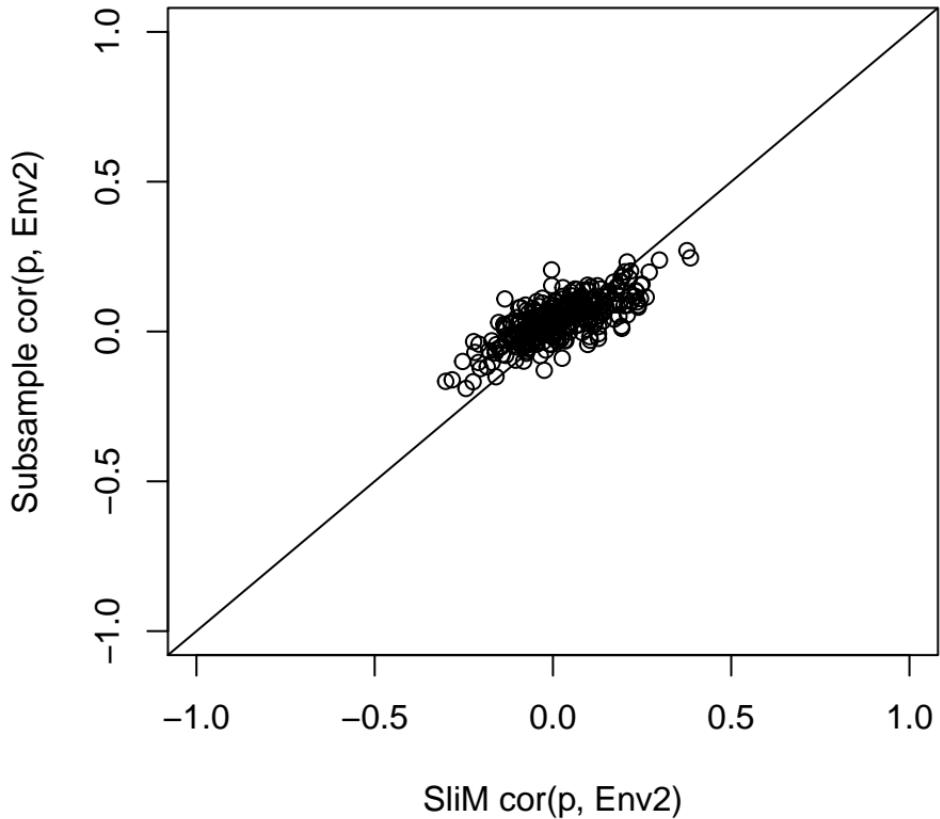
highly-polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1231094



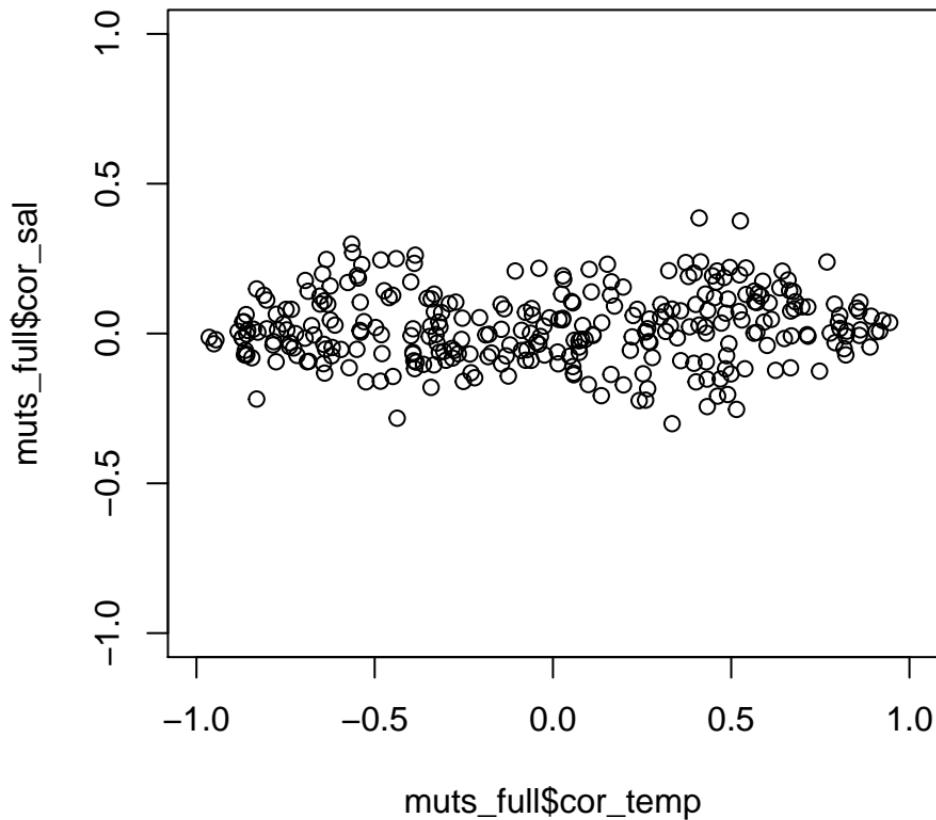
polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge  
1231094



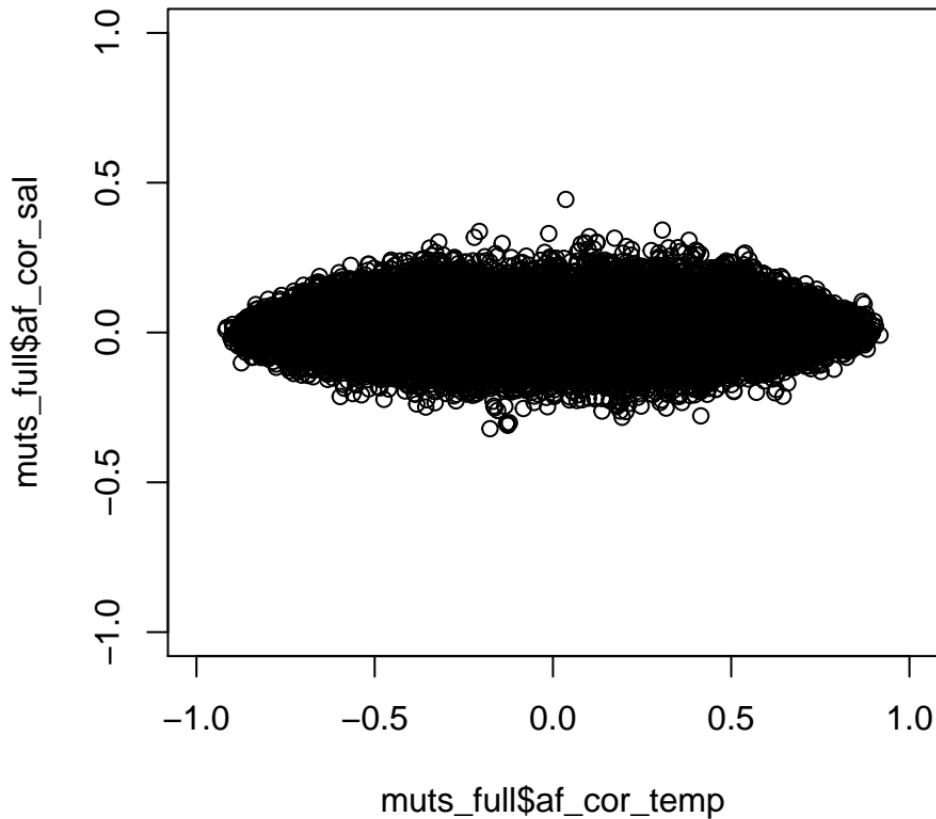
polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge  
1231094



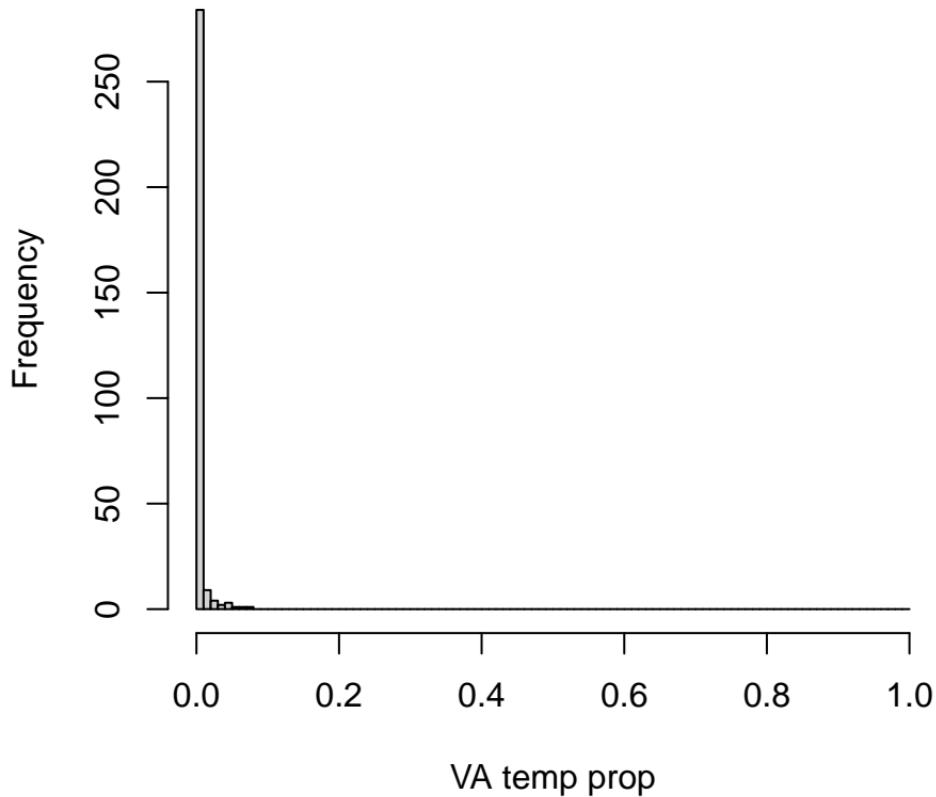
## SiM Correlation (AF, ENV) all samples



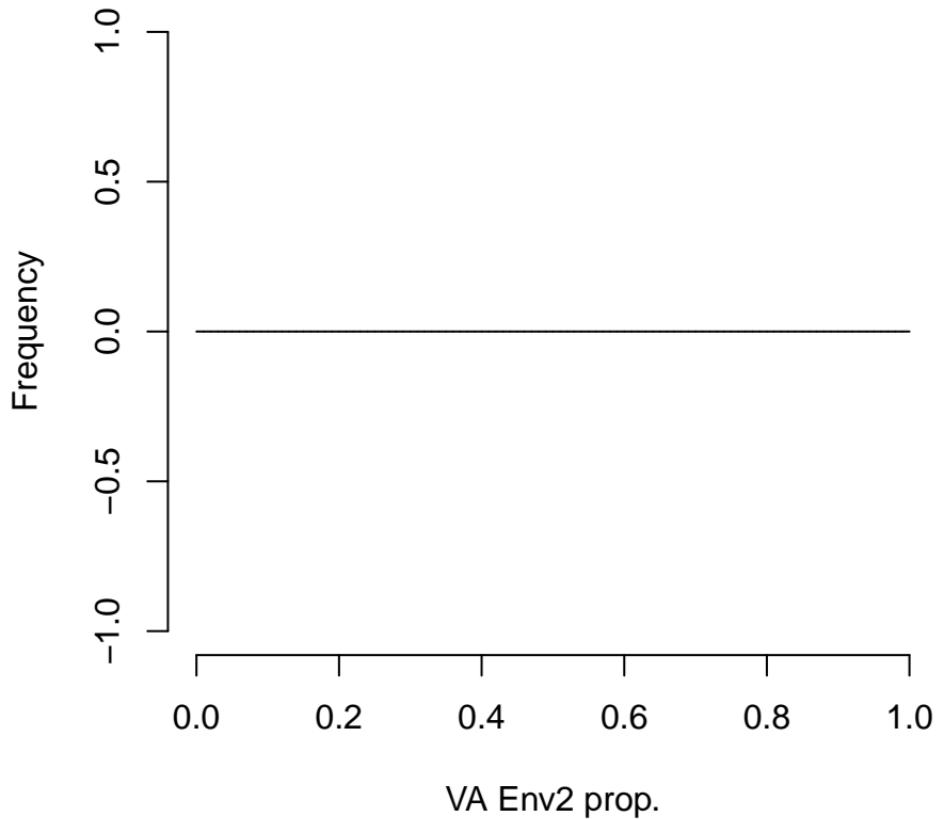
## Subsample Correlation (AF, ENV) all samples

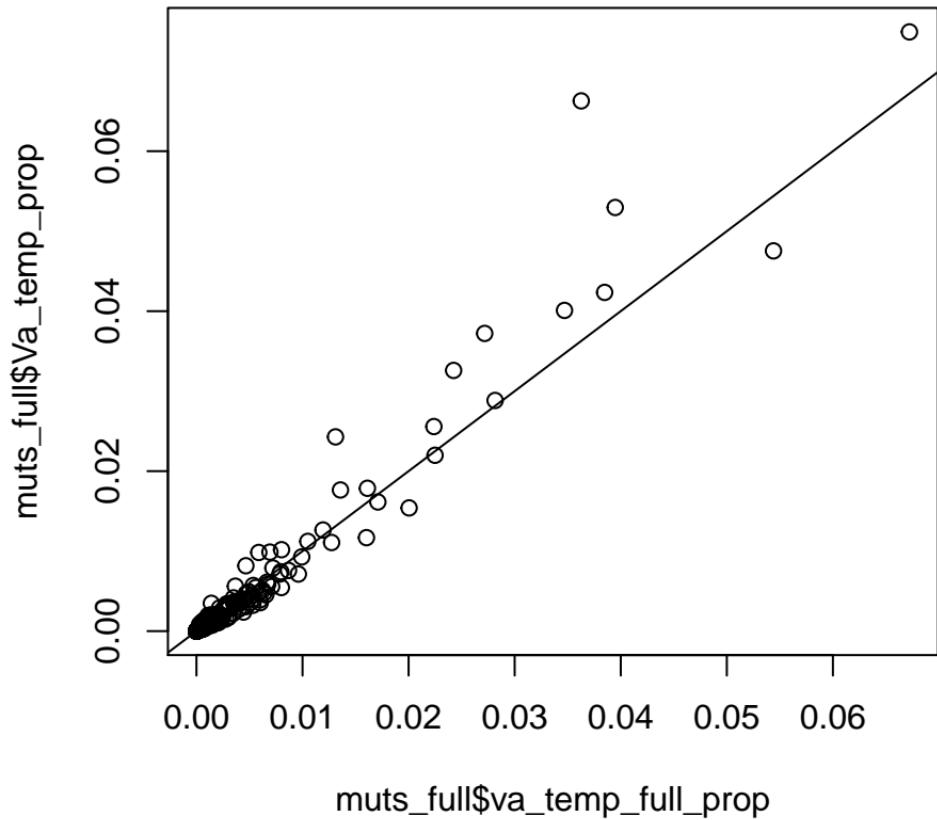


polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge  
1231094

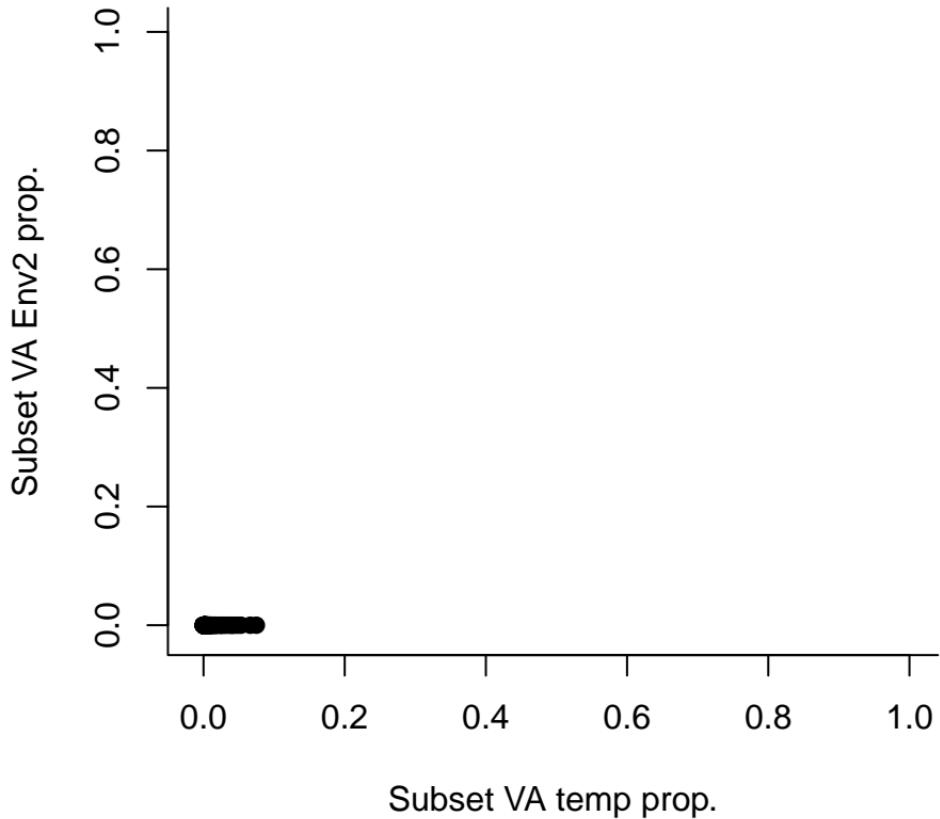


**polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge**  
**1231094**

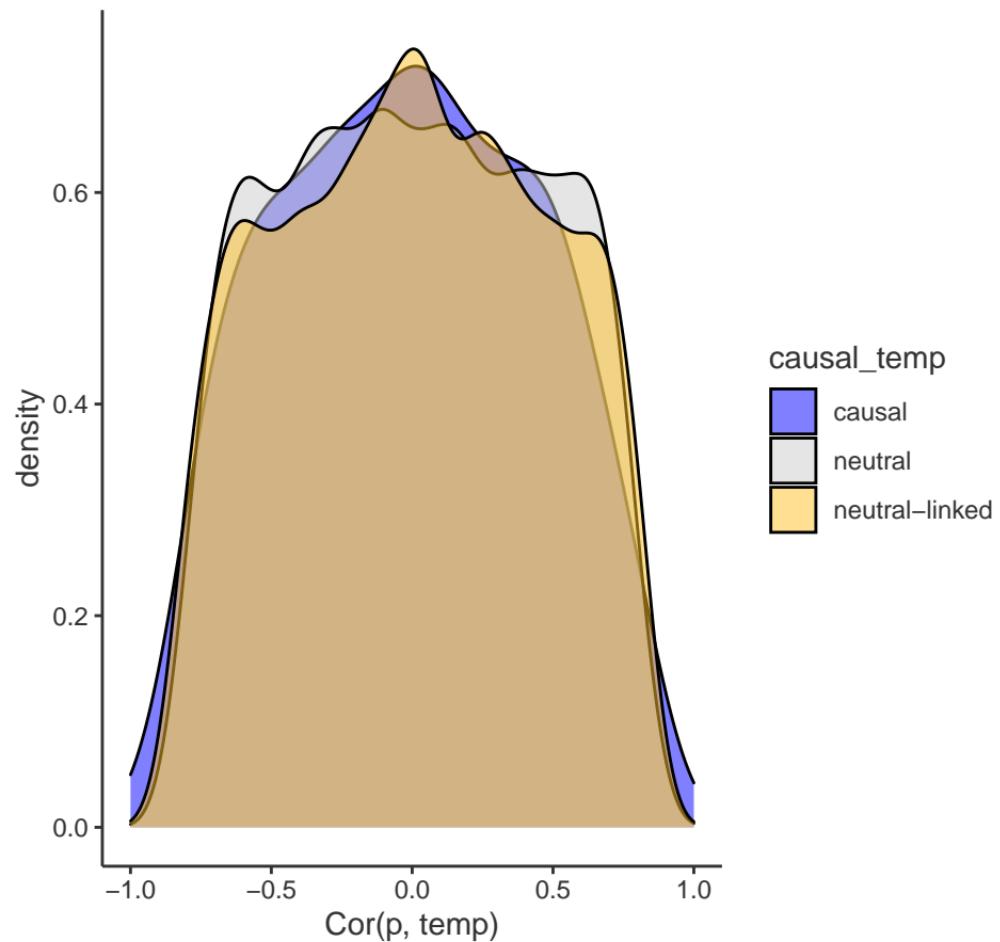




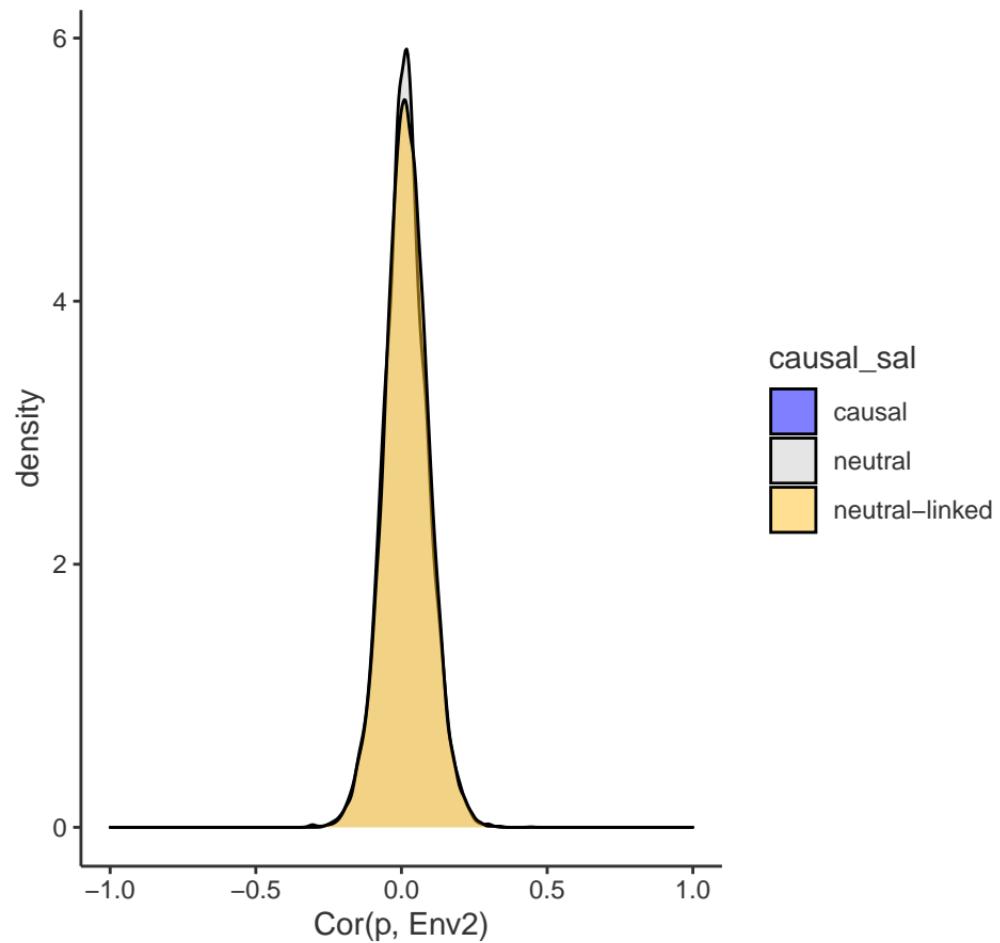
**polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge**  
**1231094**



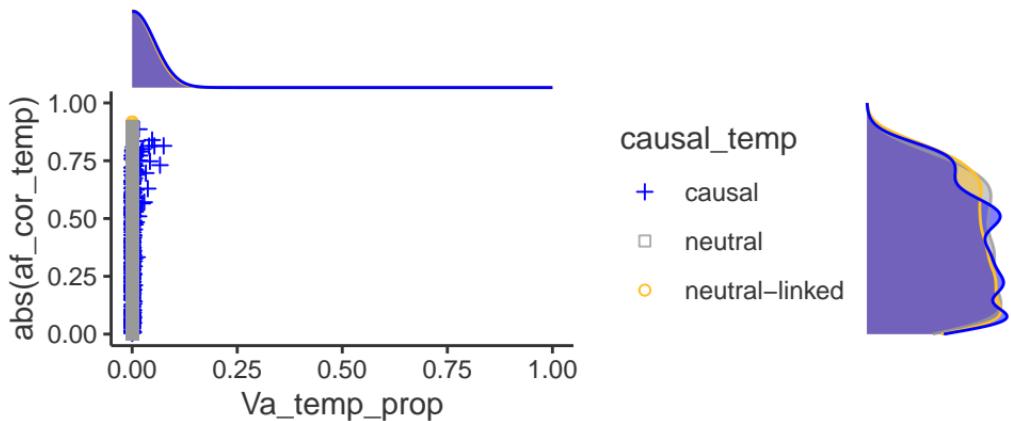
highly-polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1231094



highly-polygenic\_1-trait\_Est-Clines\_N-cline-center-to-edge\_m-constant  
1231094



highly-polygenic\_1-trait\_Est-Clines\_N-cline  
1231094



highly-polygenic\_1-trait\_Est-Clines\_N-cline  
1231094

