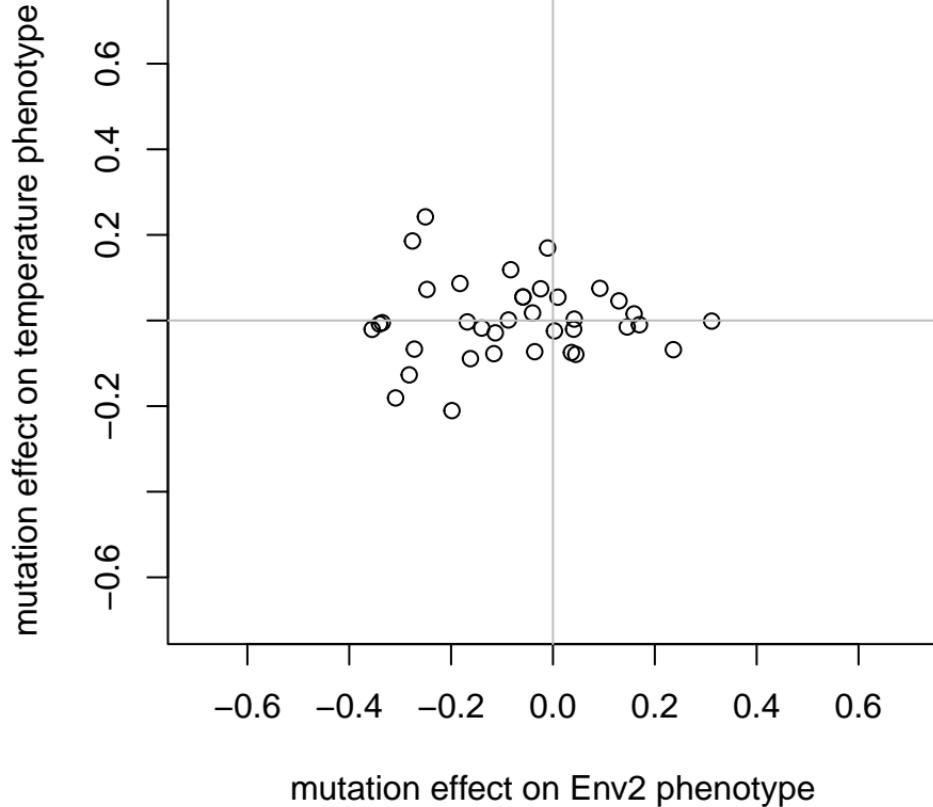
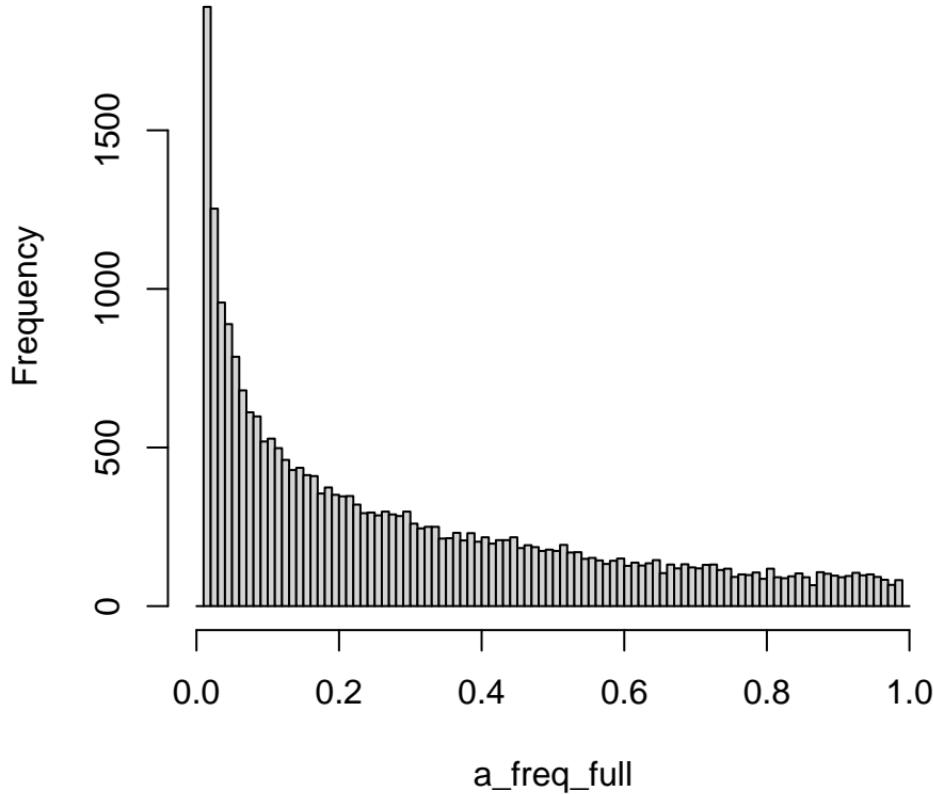


mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-cline-center-to-edge_m-constant
1231214



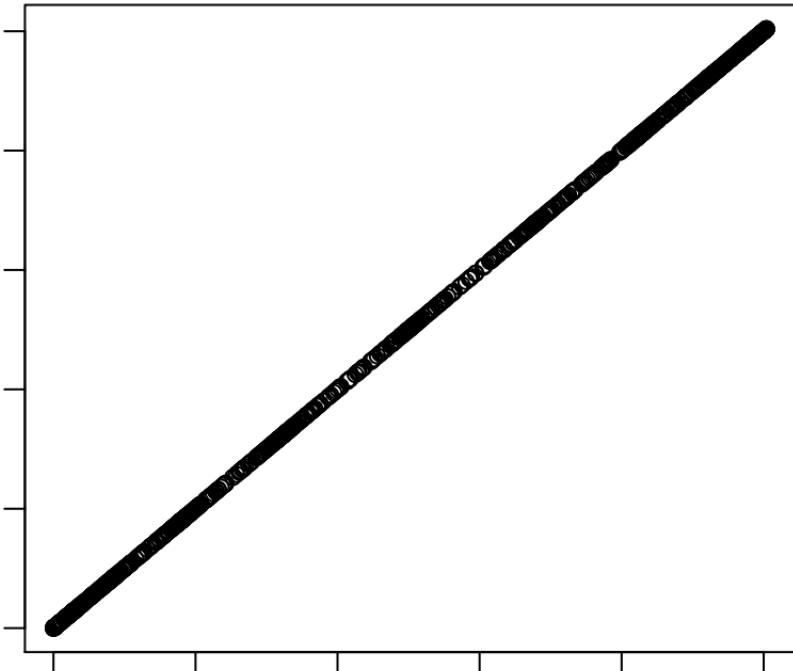
Histogram of a_freq_full

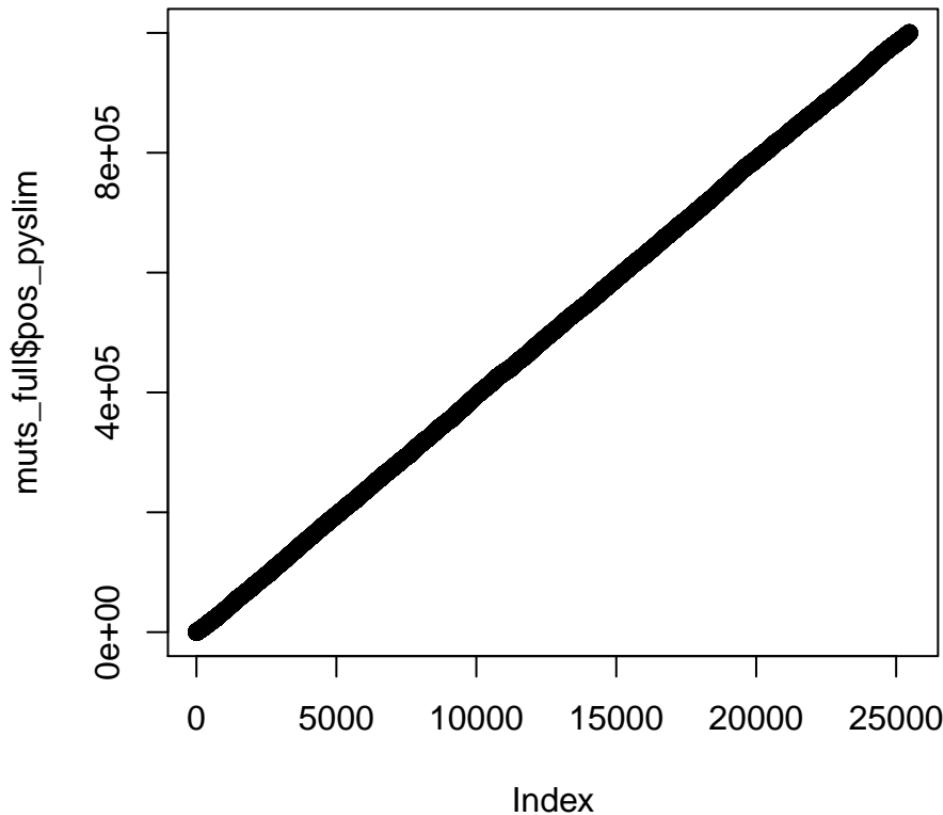


indPhen_df\$indID[which(indPhen_df\$subset)]

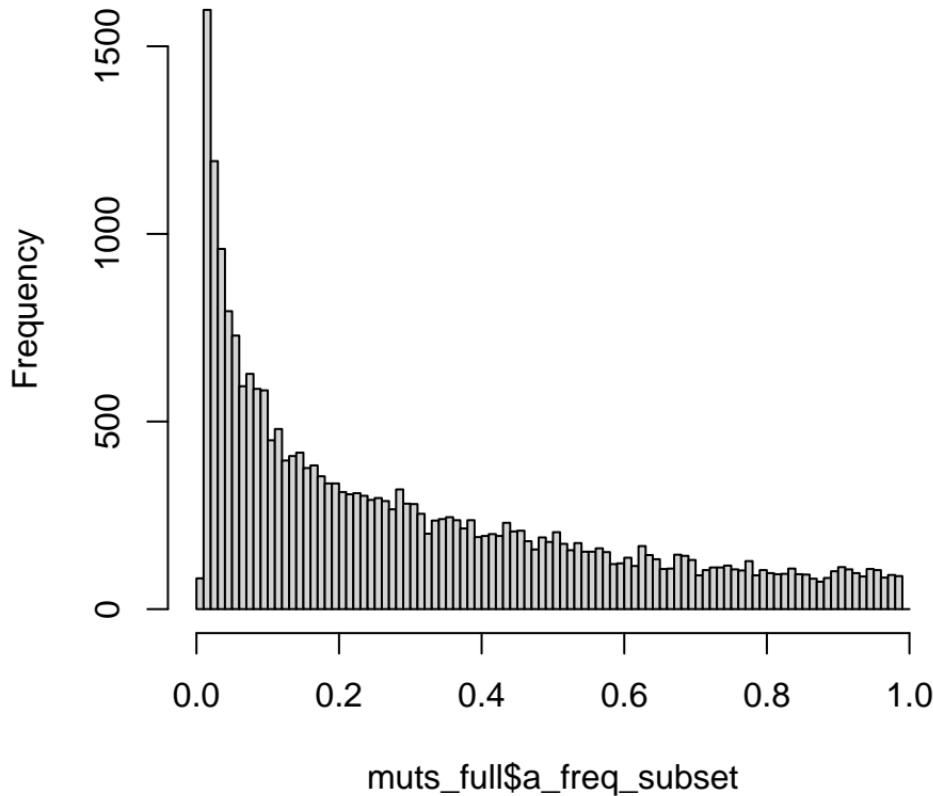
0 2000 4000 6000 8000 10000

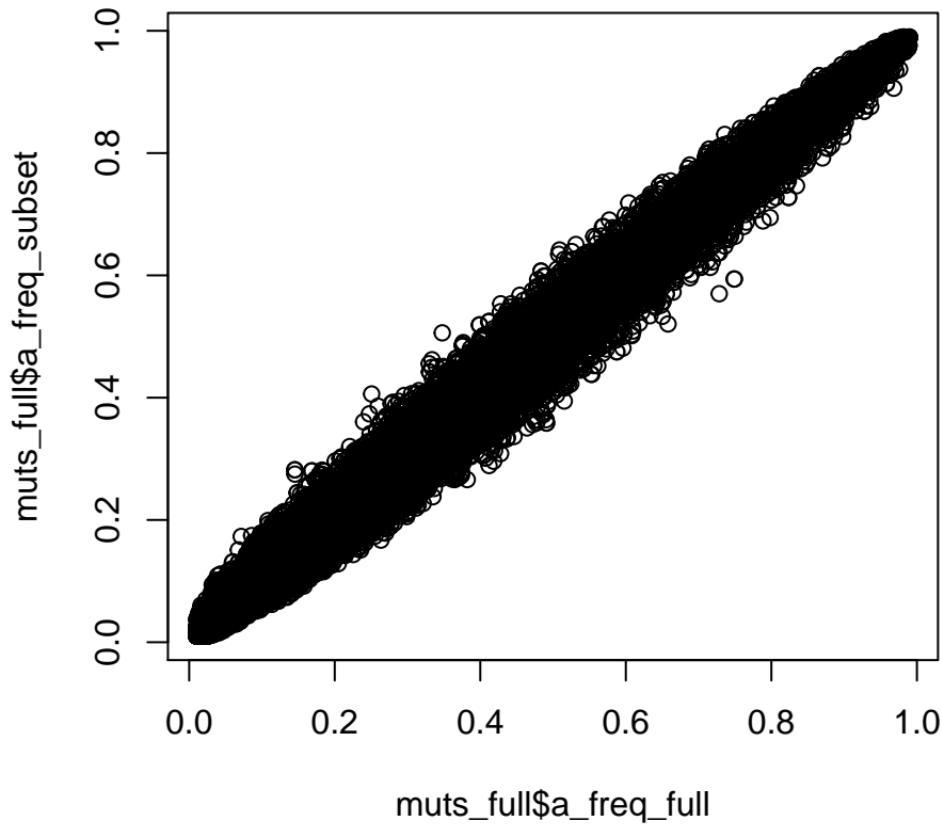
colnames(G_full_subset)



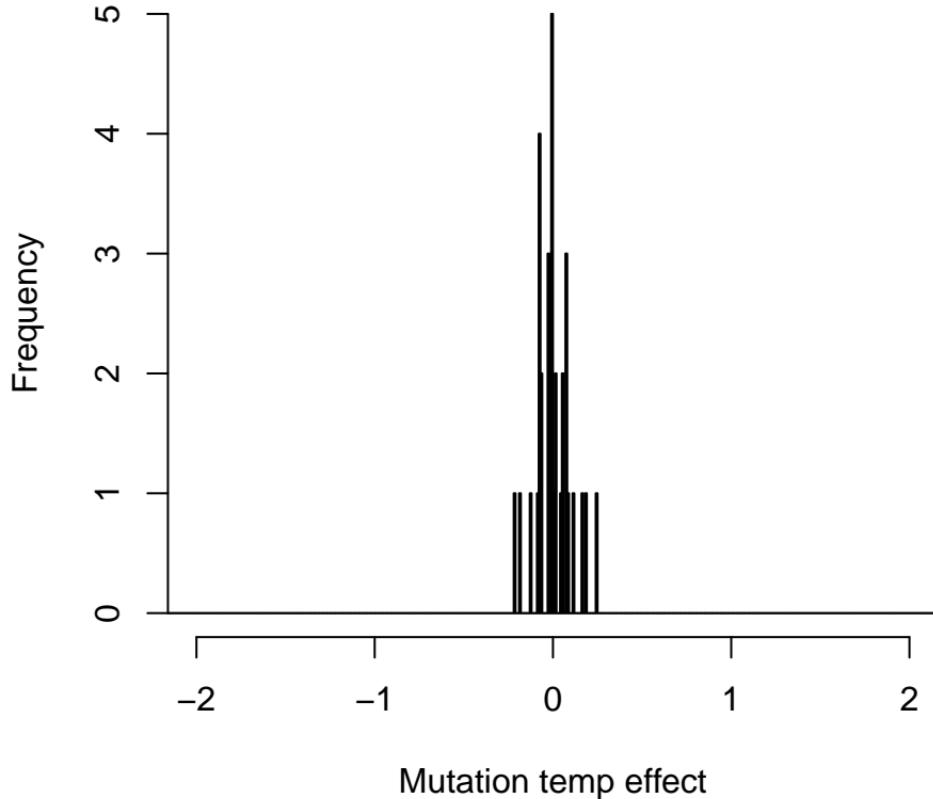


Histogram of muts_full\$a_freq_subset

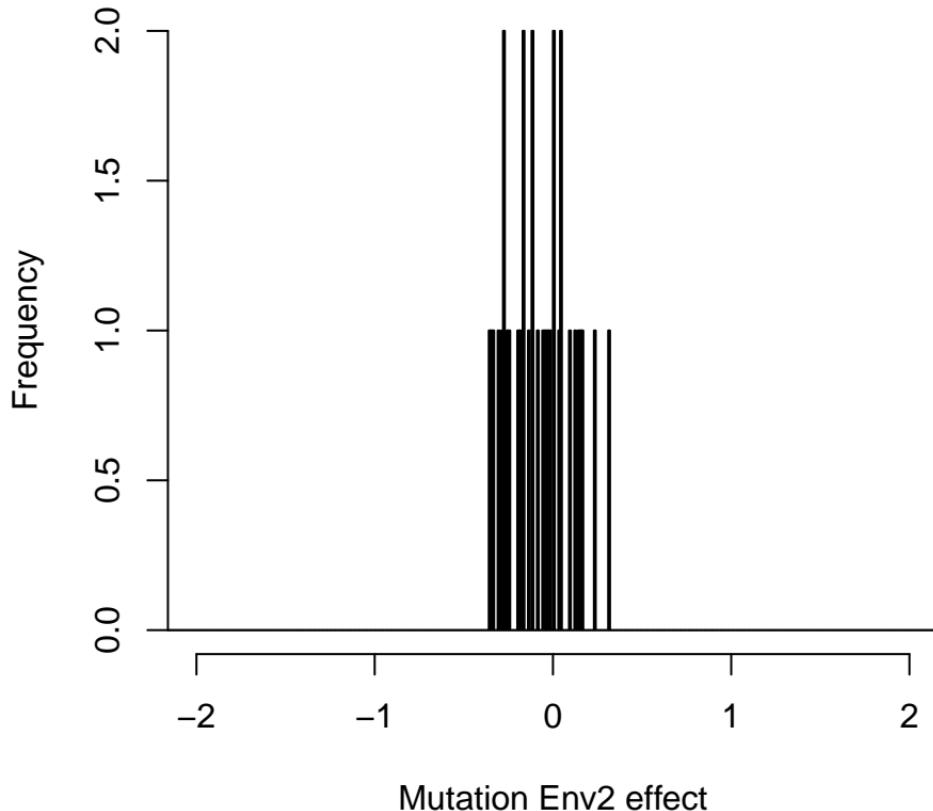




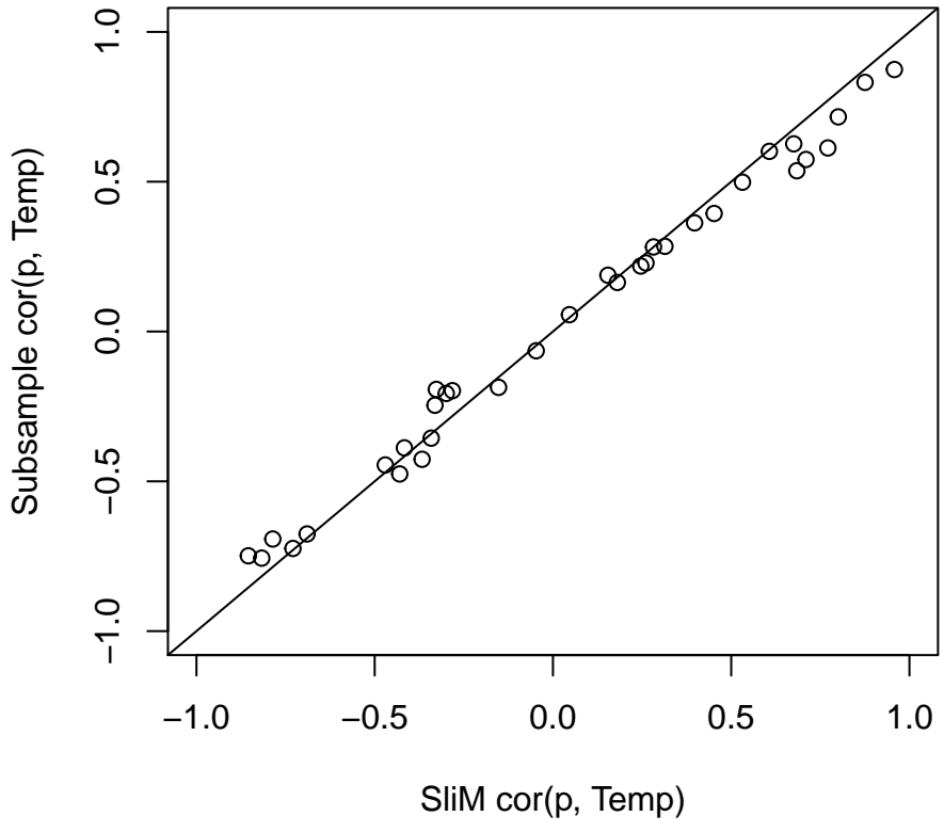
mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-cline-center-to-edge_m-constant
1231214



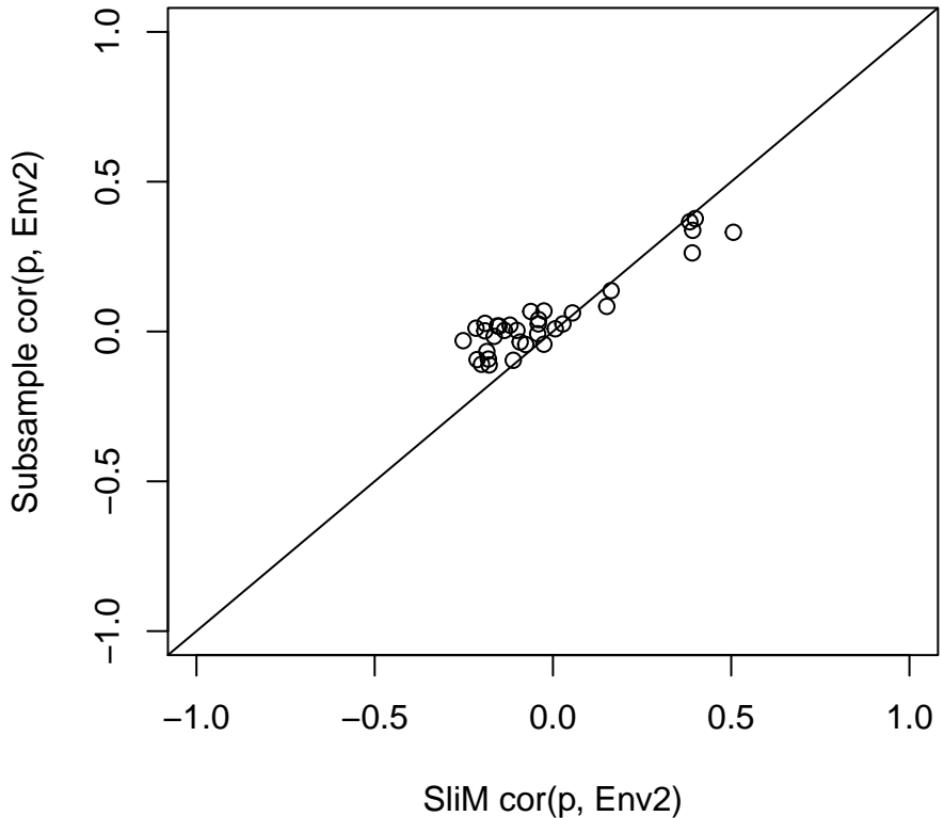
mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-cline-center-to-edge_m-constant
1231214



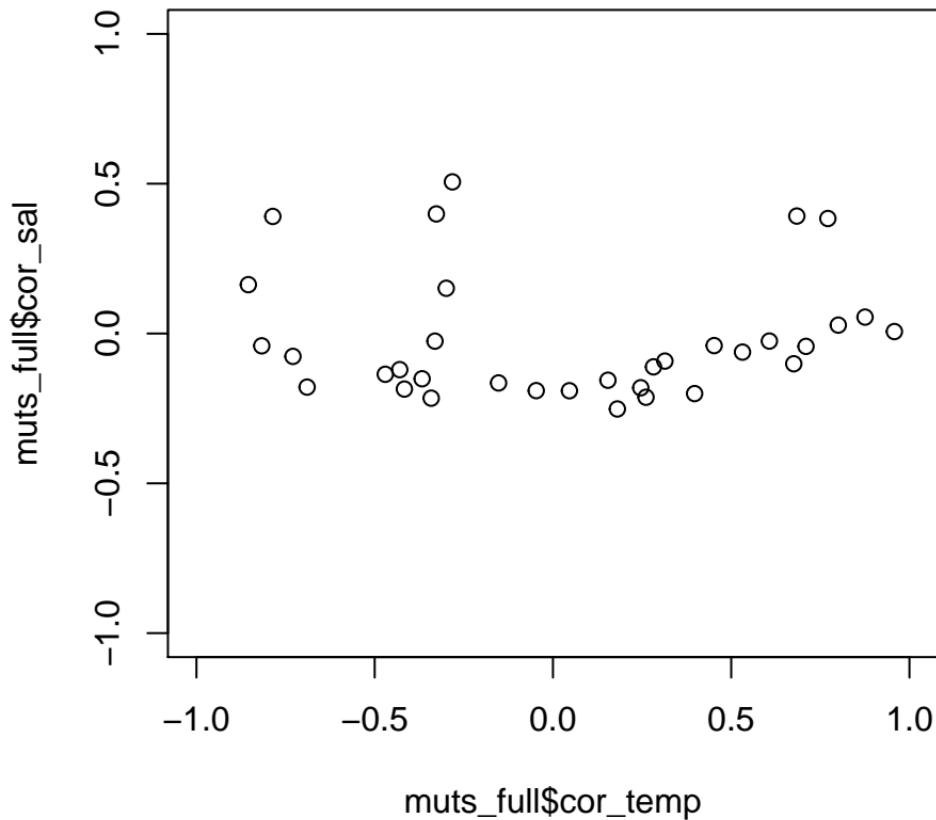
-trait-pleiotropy-equal-S_Est-Clines_N-cline-center
1231214



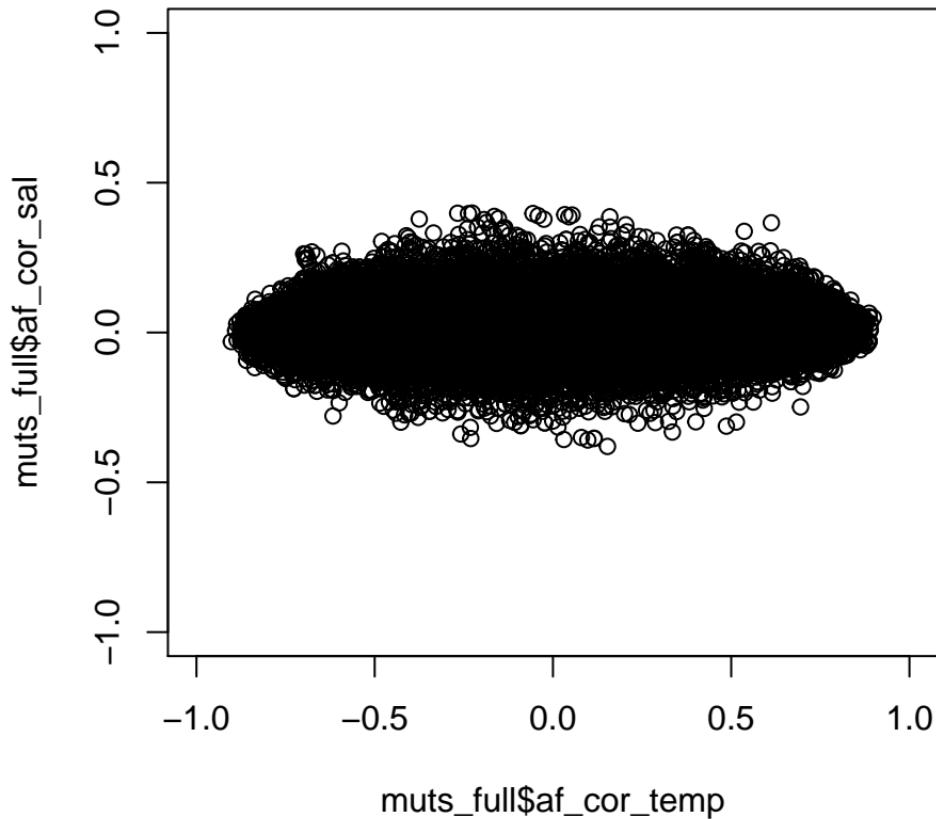
-trait-pleiotropy-equal-S_Est-Clines_N-cline-center
1231214



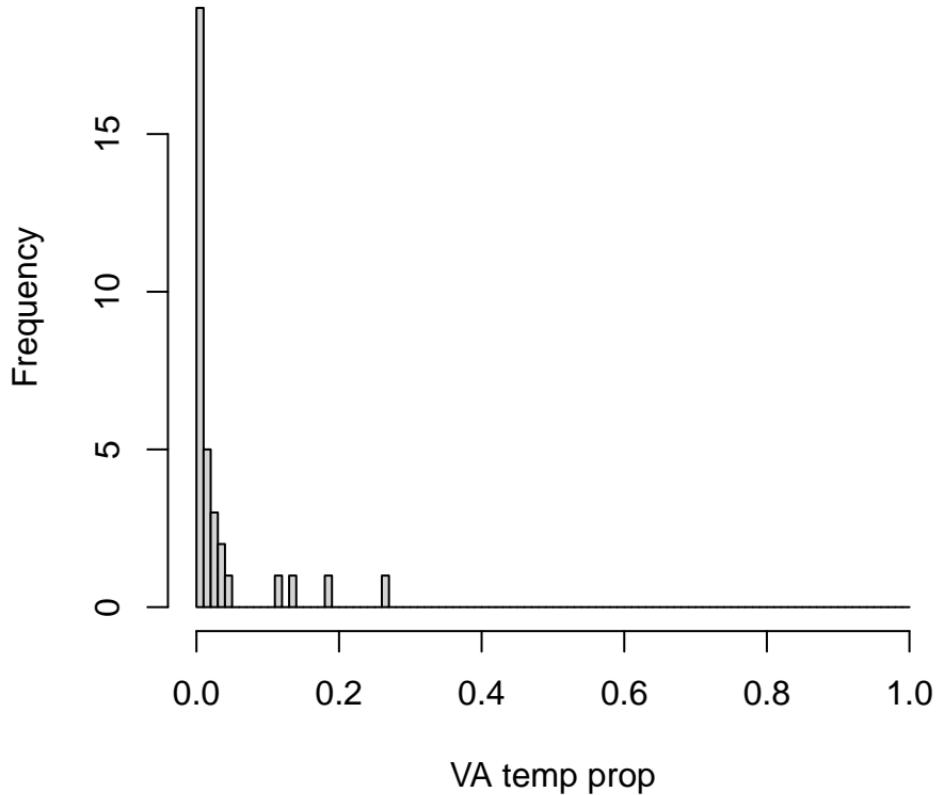
SiM Correlation (AF, ENV) all samples



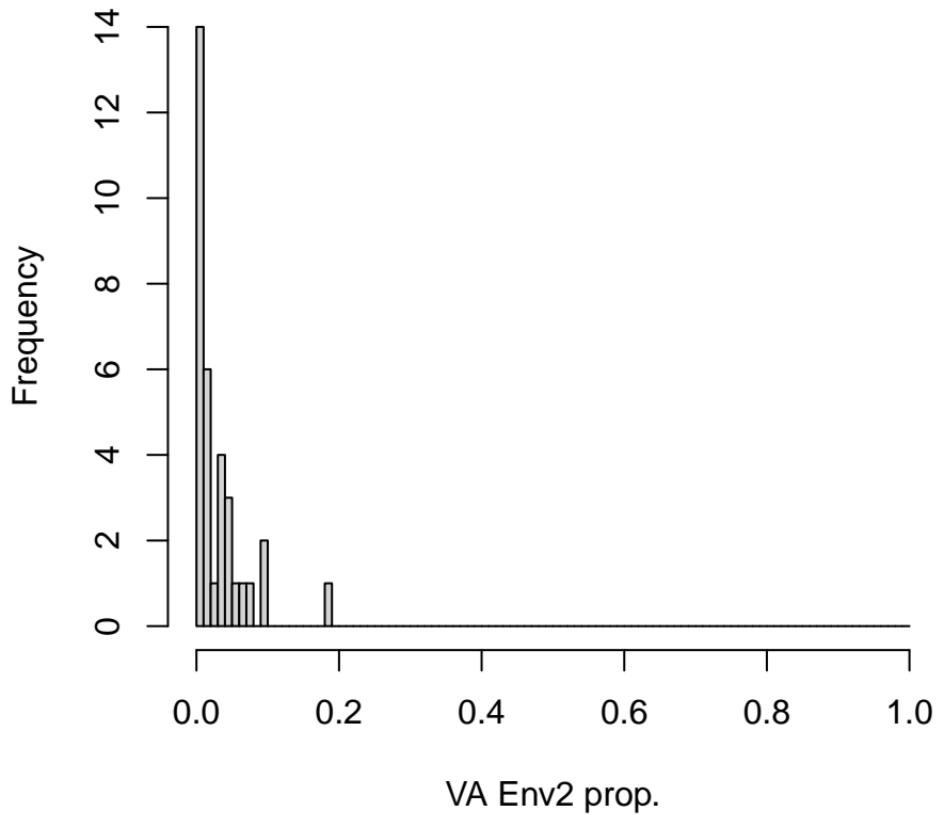
Subsample Correlation (AF, ENV) all samples

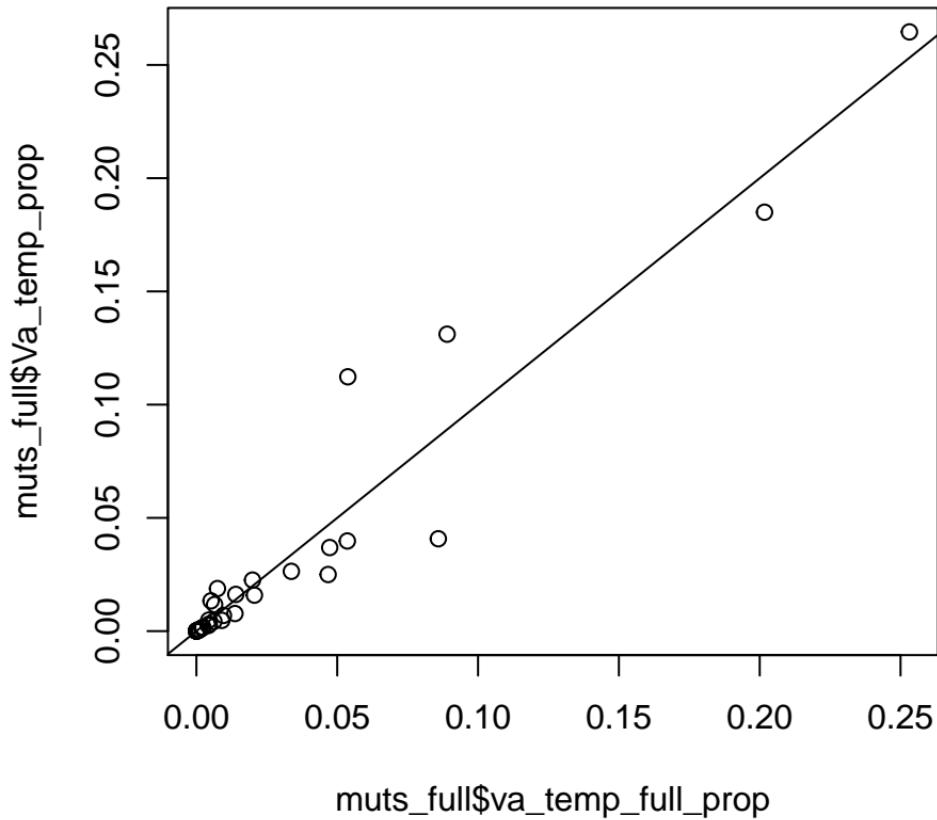


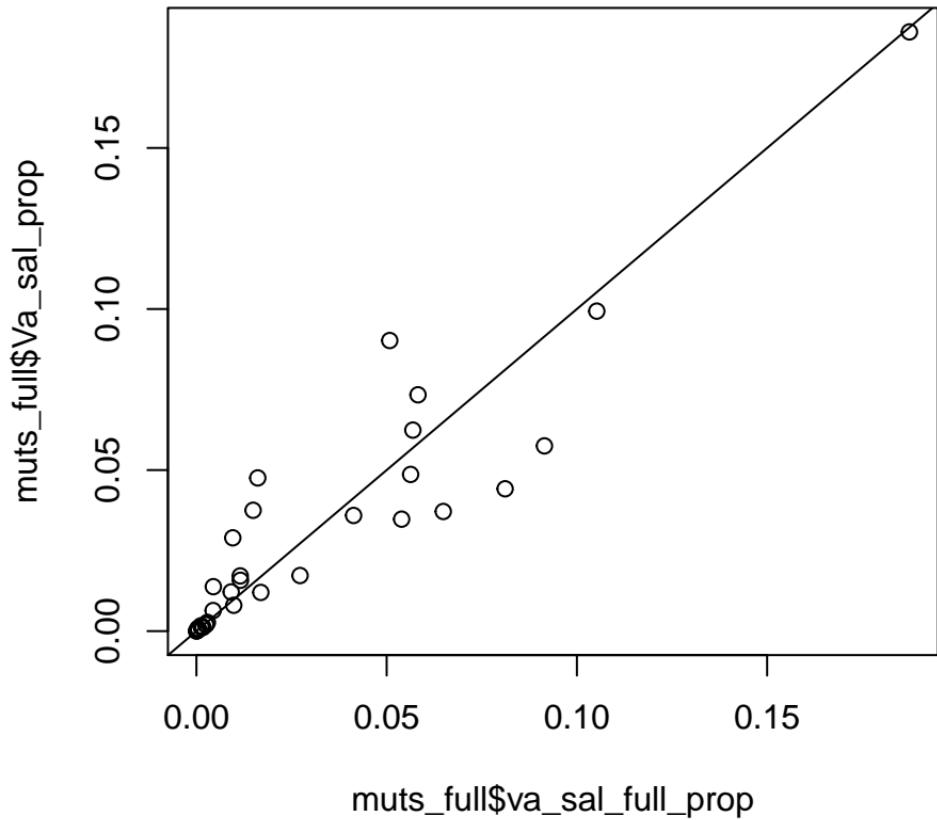
-trait-pleiotropy-equal-S_Est-Clines_N-cline-center
1231214



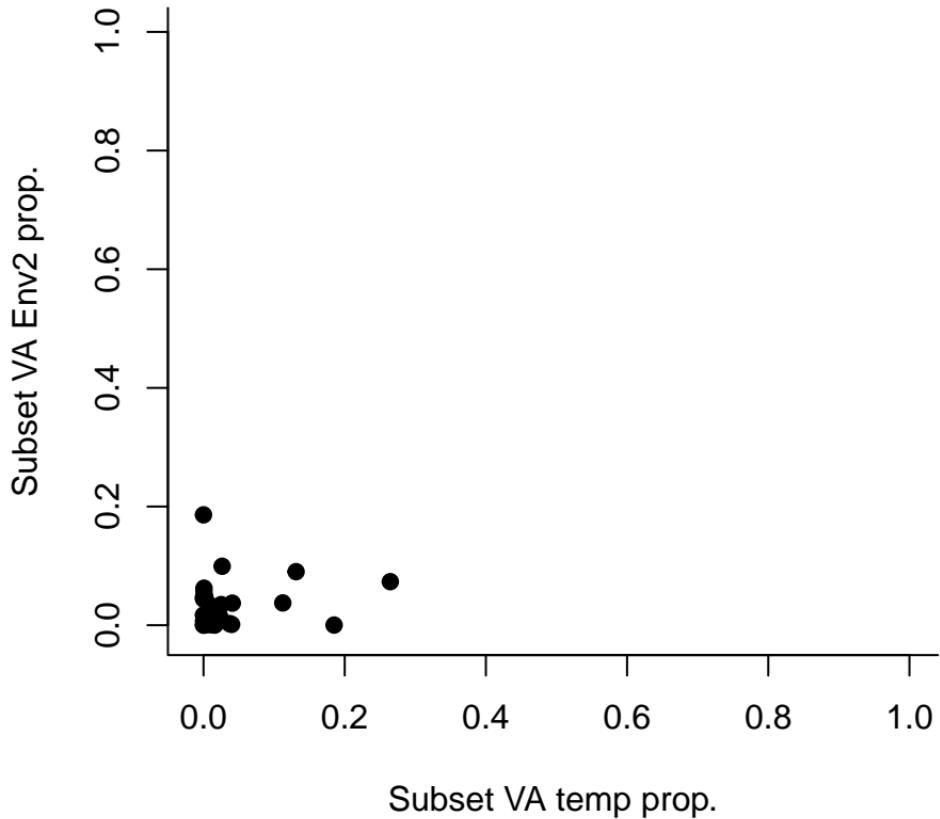
-trait-pleiotropy-equal-S_Est-Clines_N-cline-center
1231214



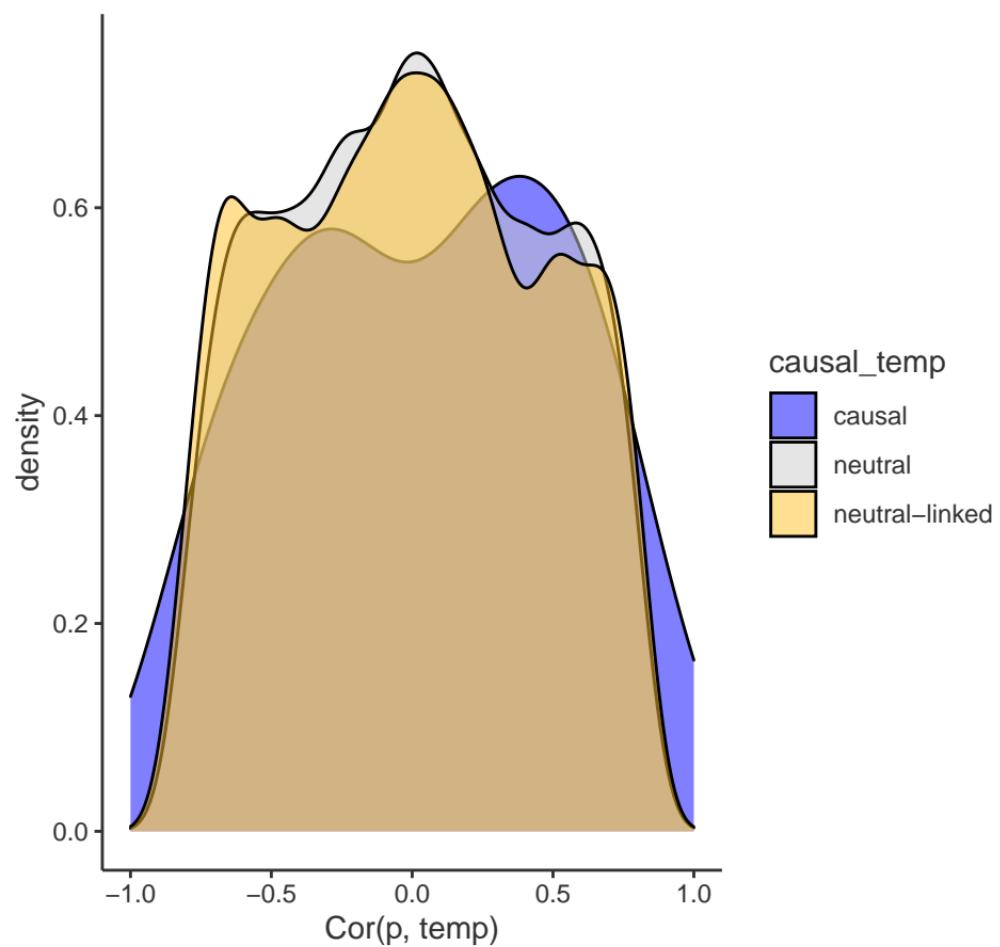




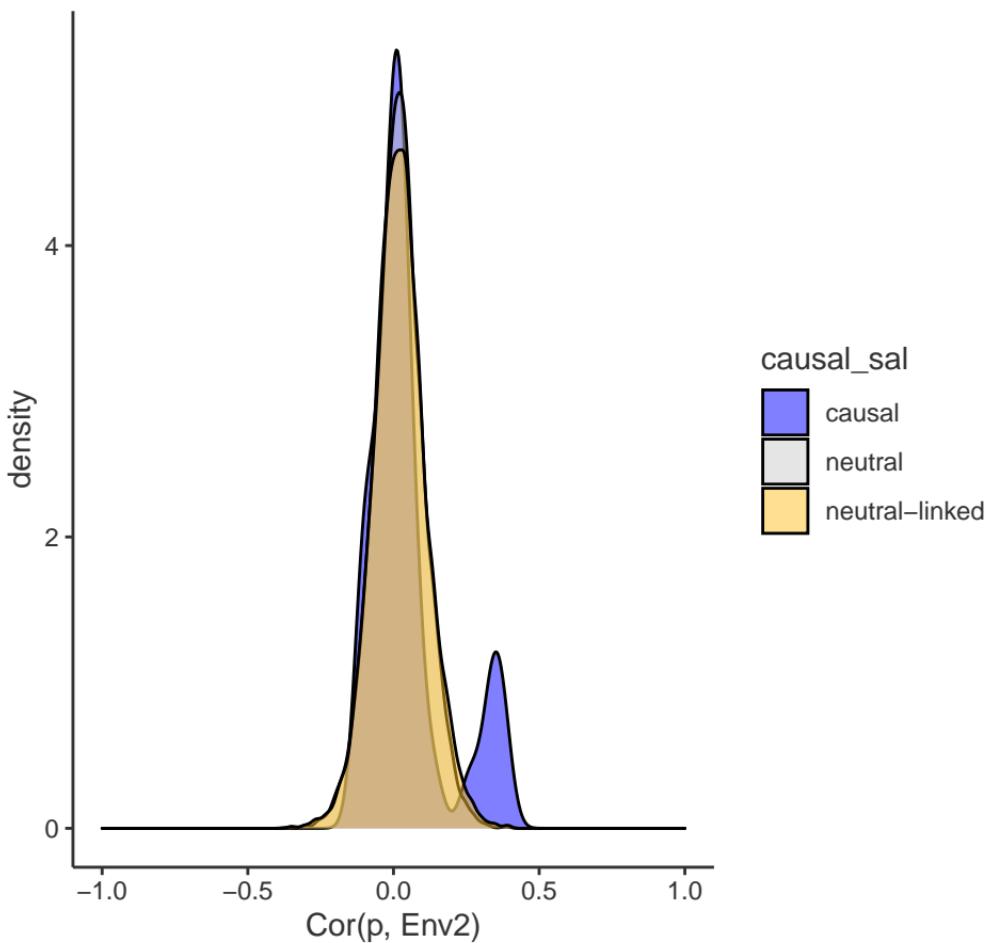
-trait-pleiotropy-equal-S_Est-Clines_N-cline-center
1231214



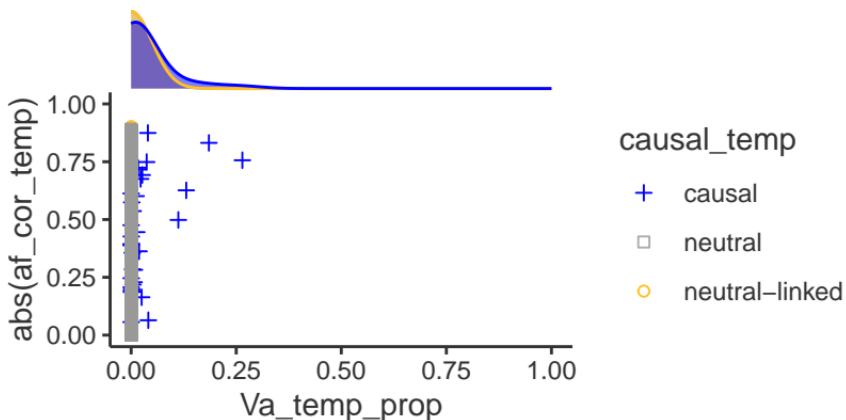
mod-polygenic_2-trait-pleiotropy-equal-S__Est-Clines_N-cline-center-to-edge_m-cons
1231214



mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-cline-center-to-edge_m-consta
1231214



mod-polygenic_2-trait-pleiotropy-equal-S__
1231214



mod-polygenic_2-trait-pleiotropy-equal-S__
1231214

