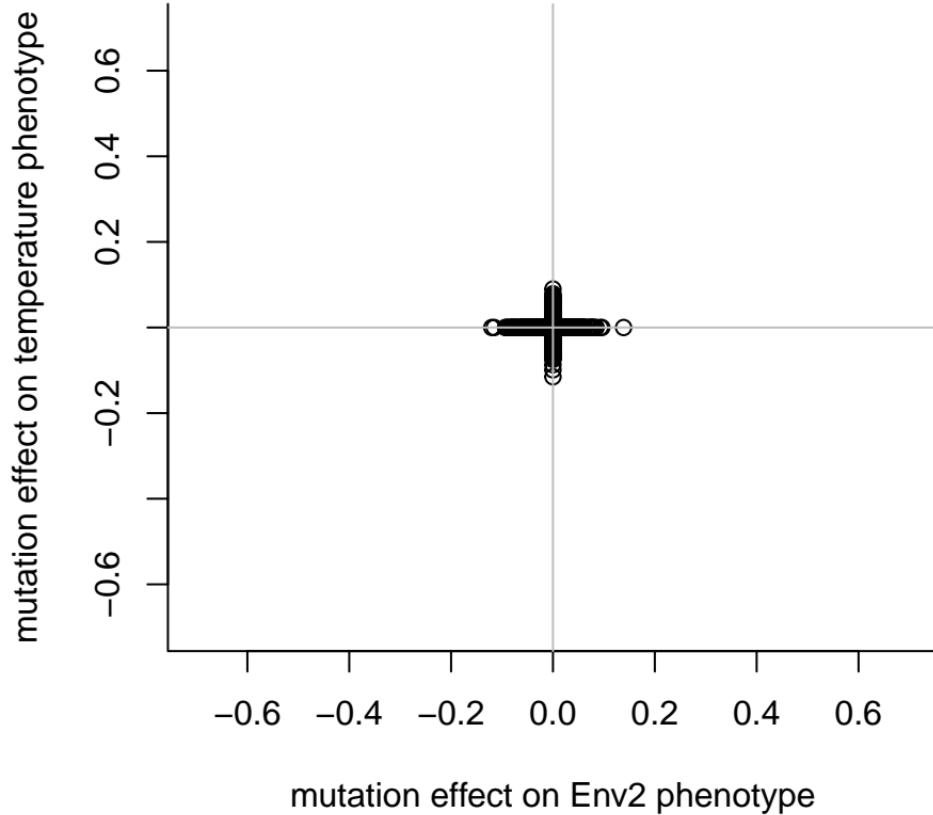
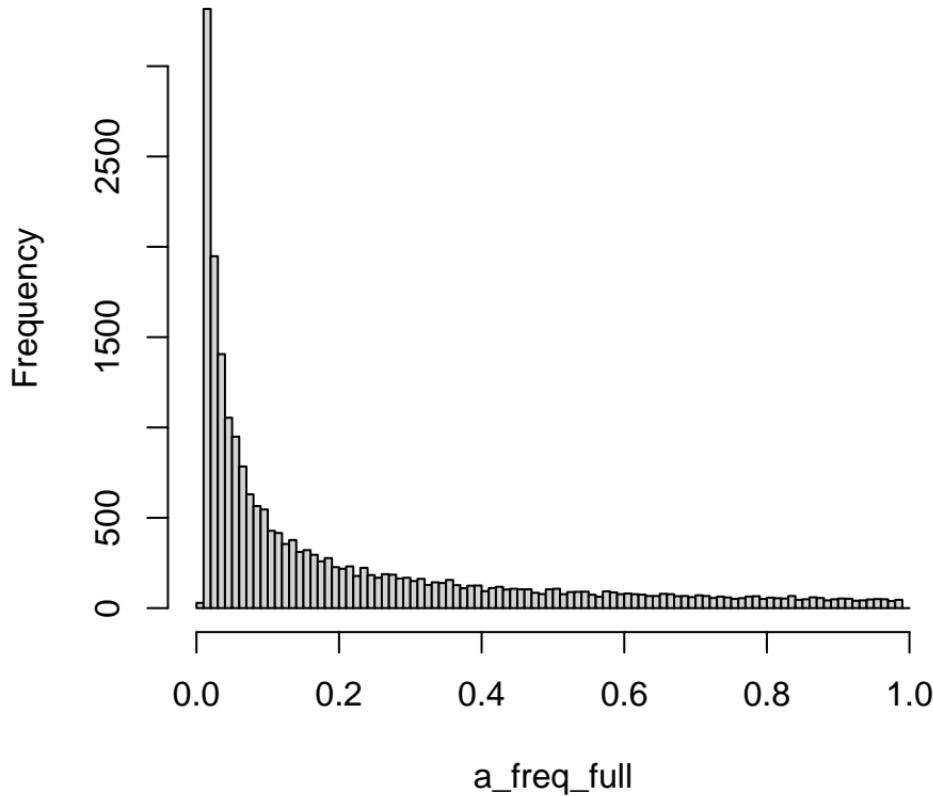


highly-polygenic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-equal_m-constant
1231122



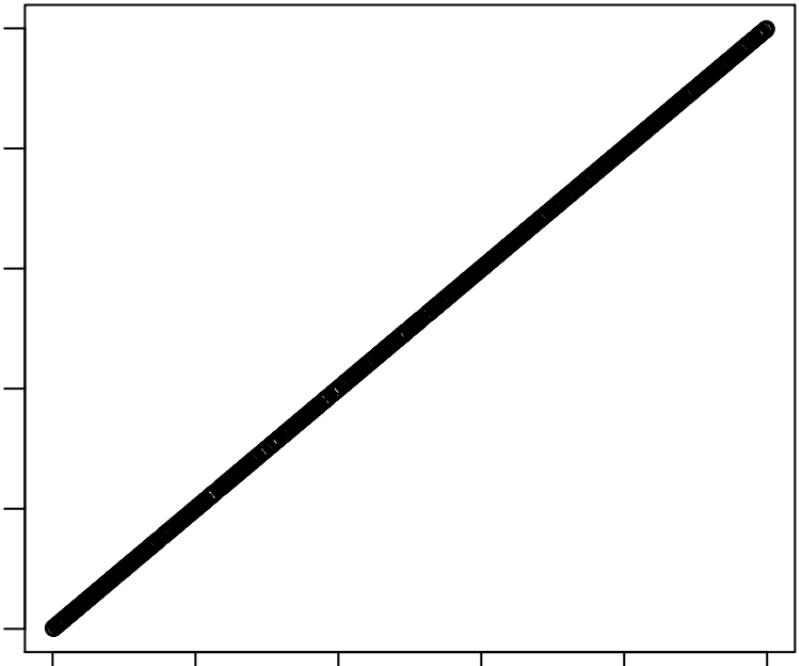
Histogram of a_freq_full

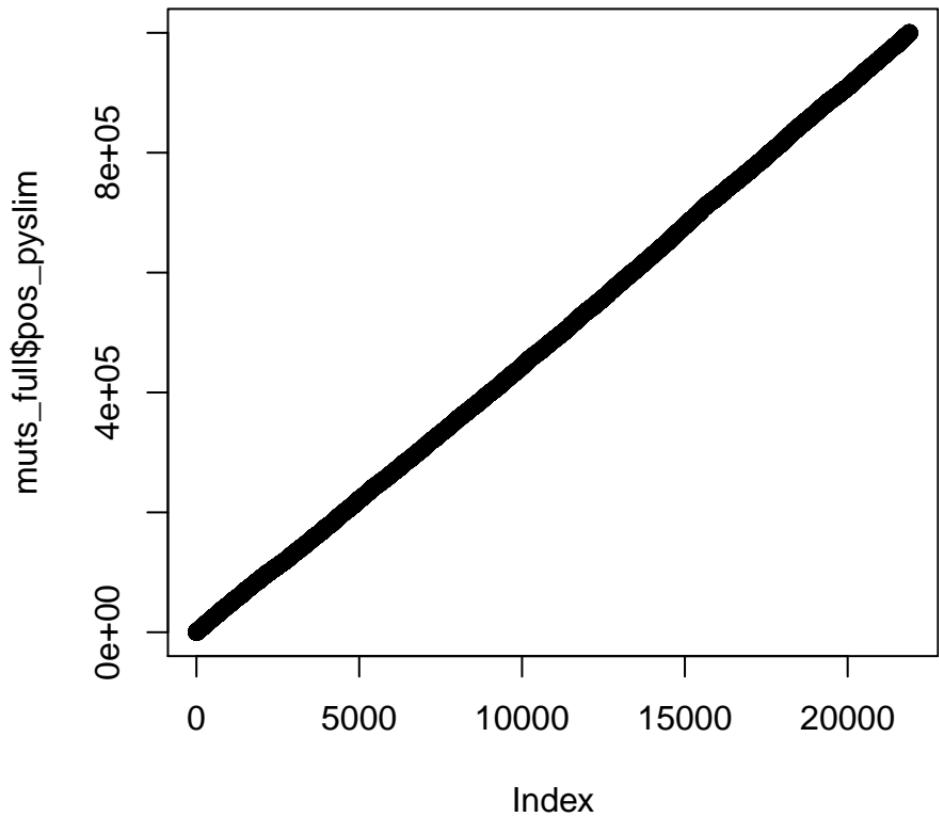


indPhen_df\$indID[which(indPhen_df\$subset)]

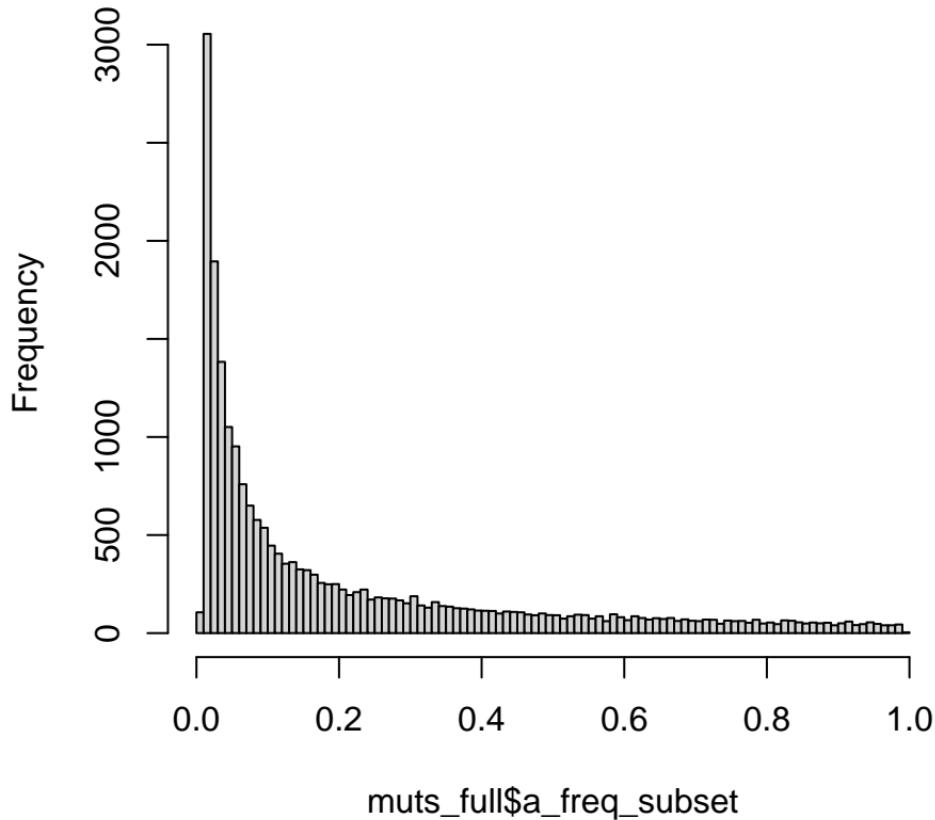
0 2000 4000 6000 8000 10000

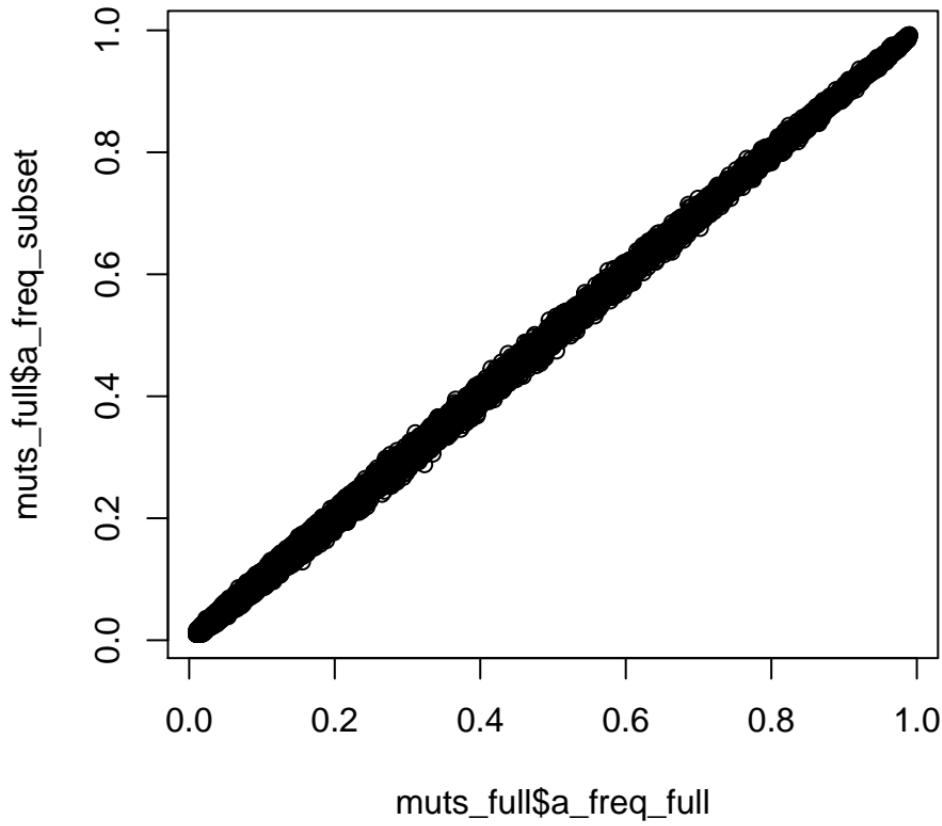
colnames(G_full_subset)



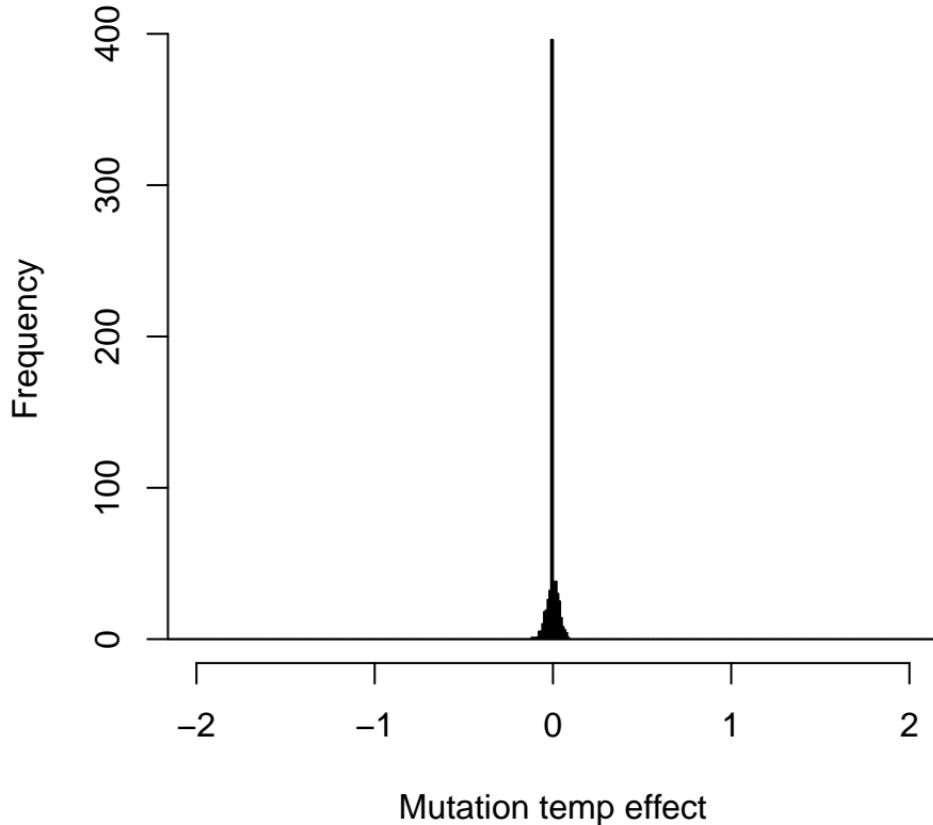


Histogram of muts_full\$a_freq_subset

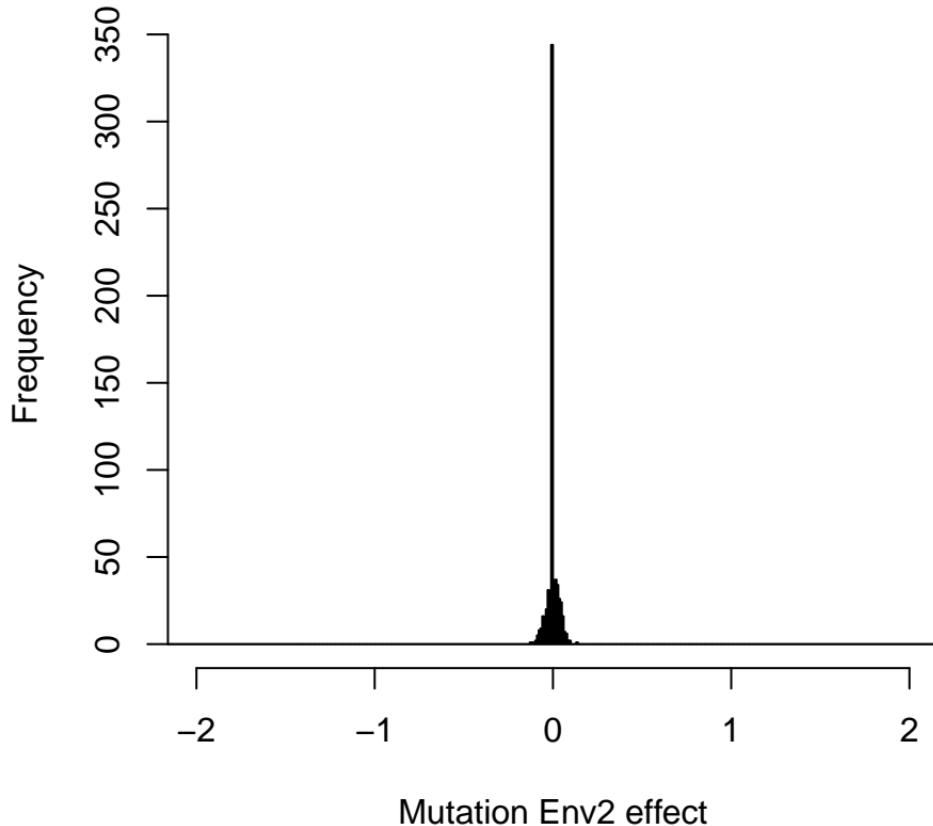




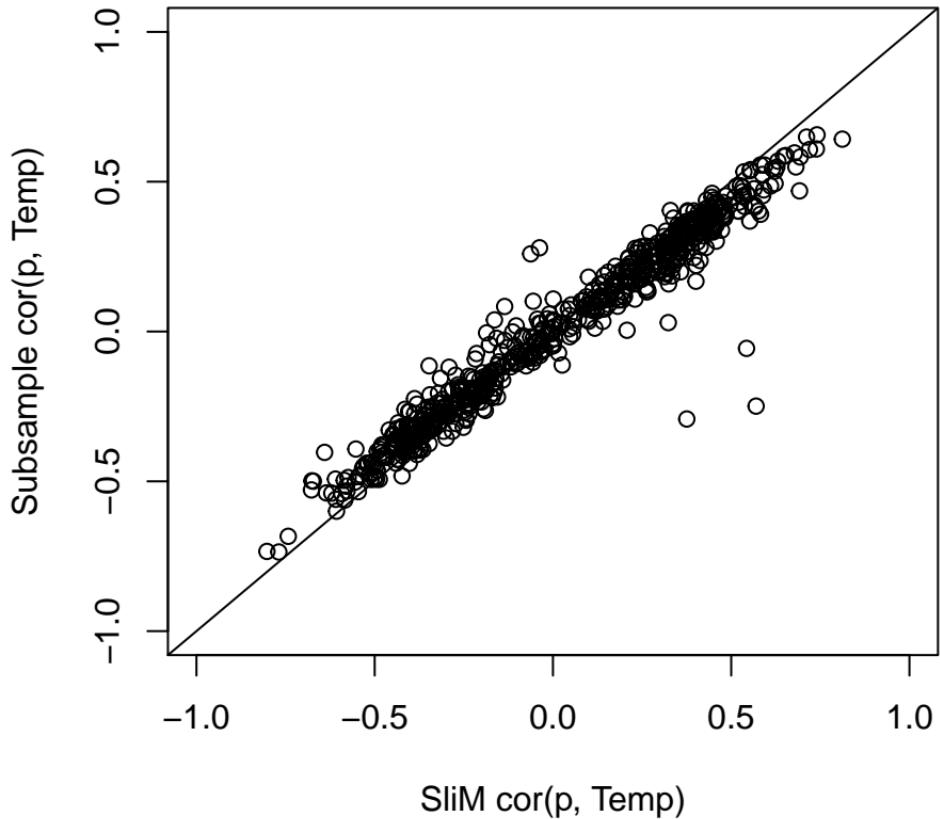
highly-polygenic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-equal_m-constant
1231122



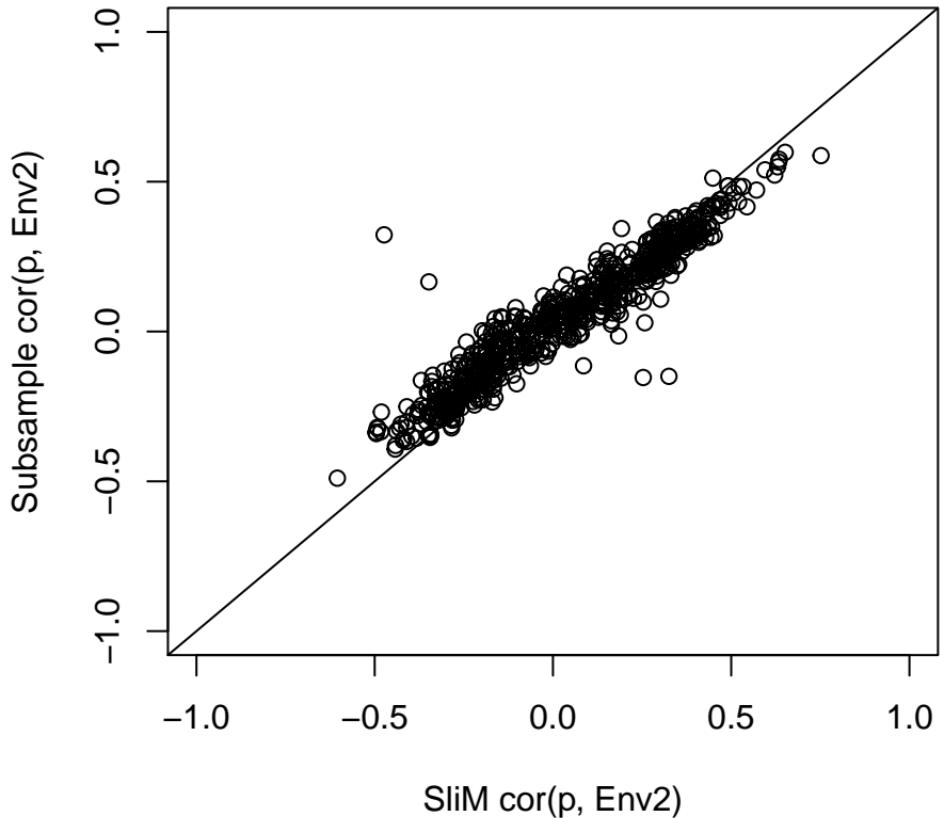
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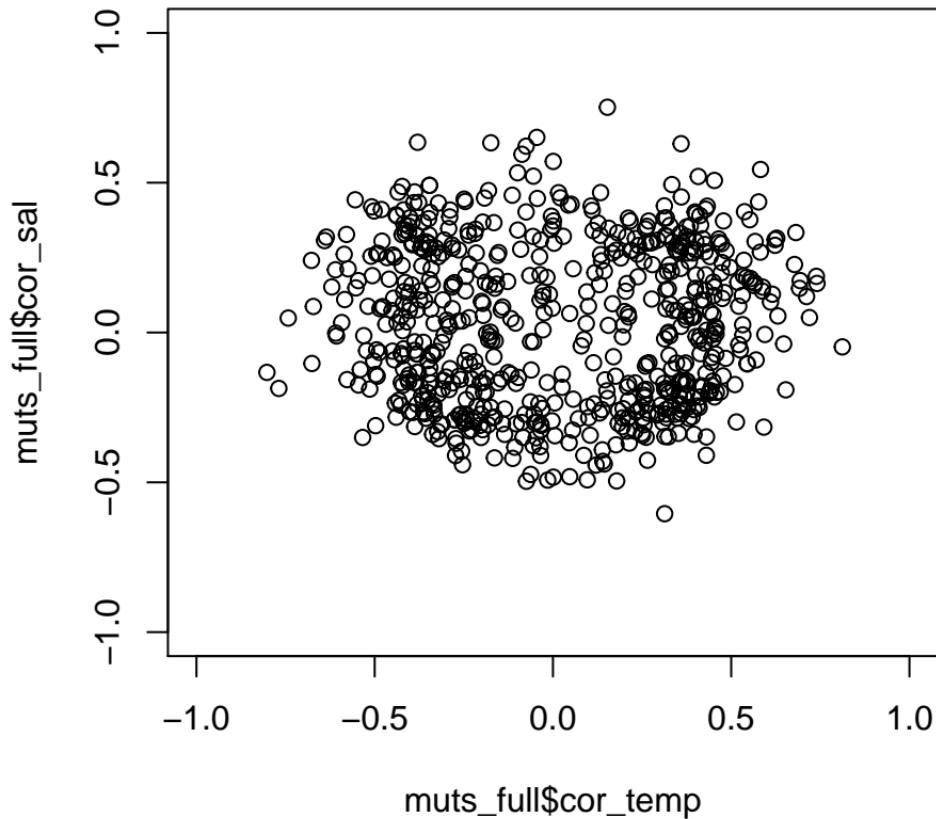
genic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-eq
1231122



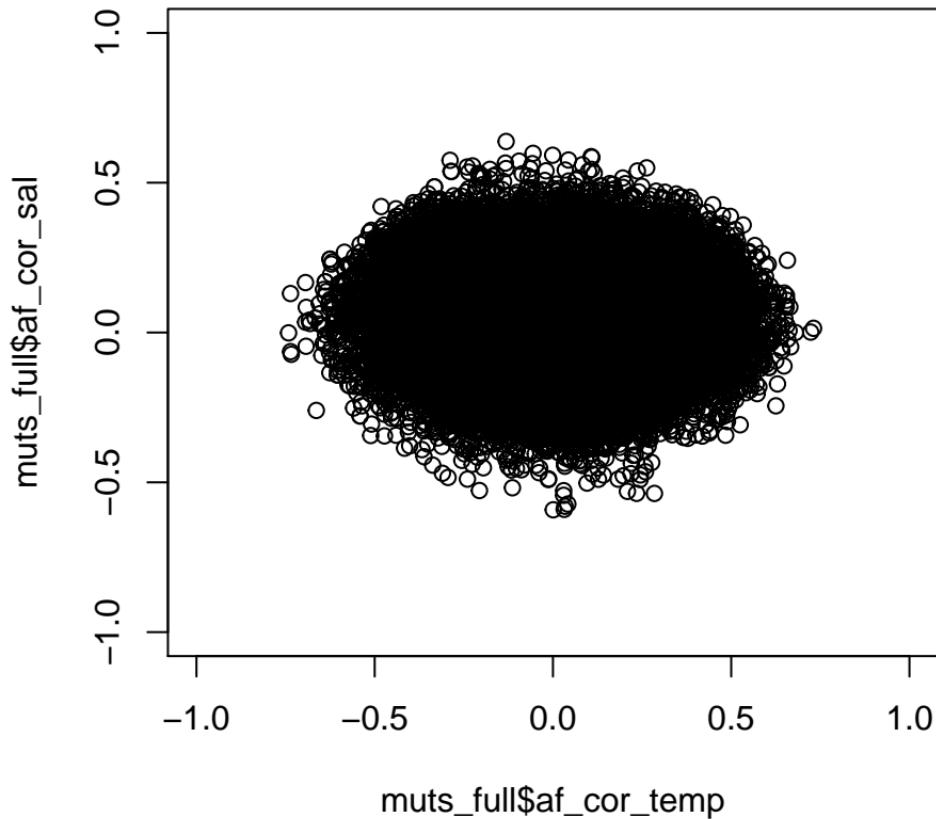
genic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-eq
1231122



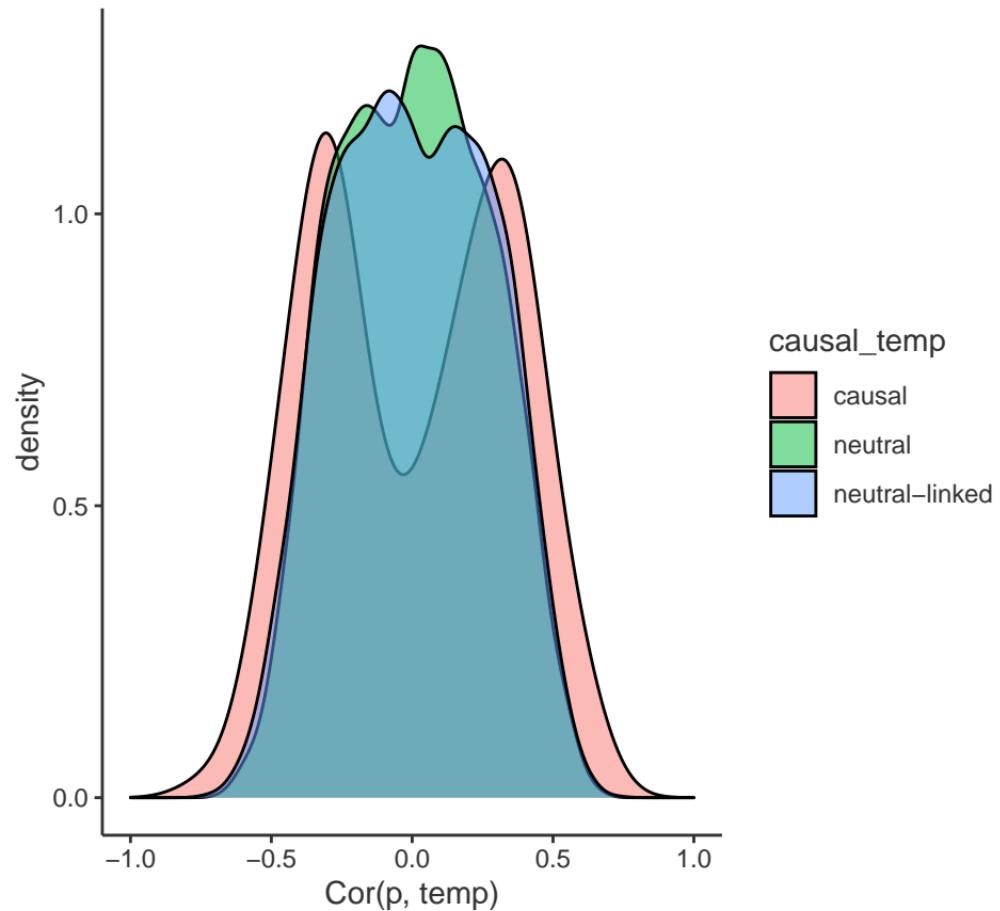
SiM Correlation (AF, ENV) all samples



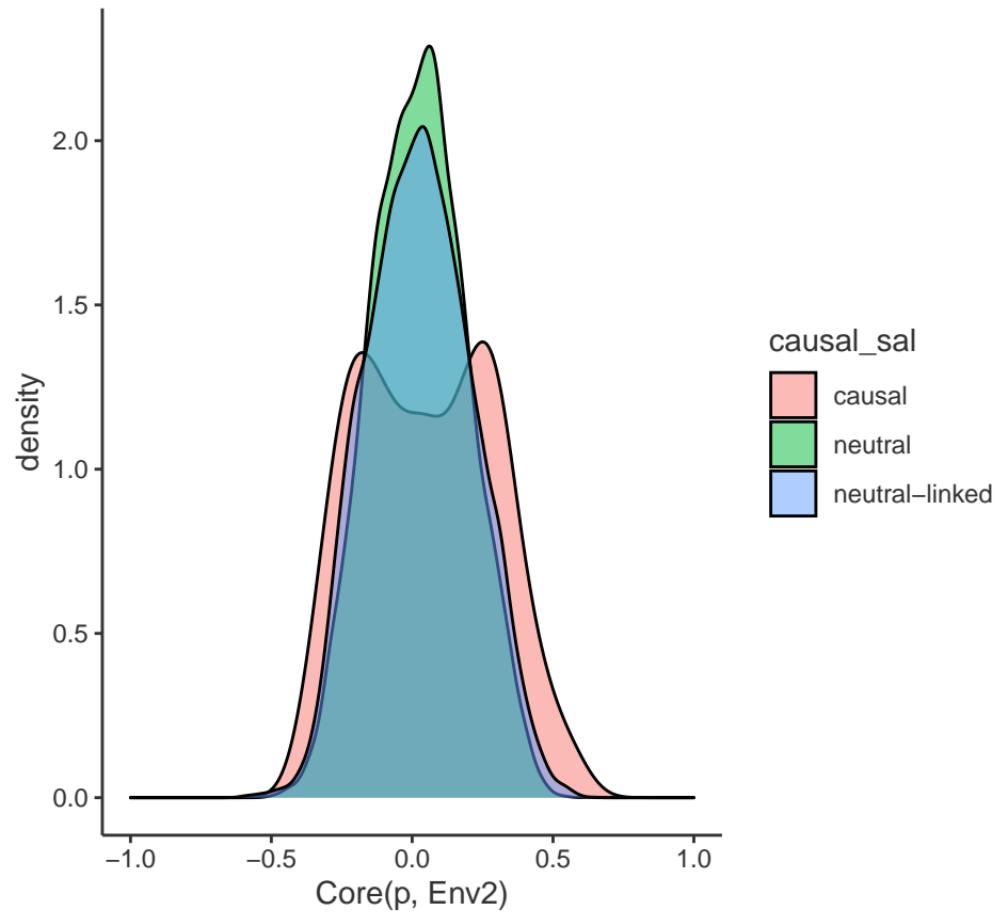
Subsample Correlation (AF, ENV) all samples



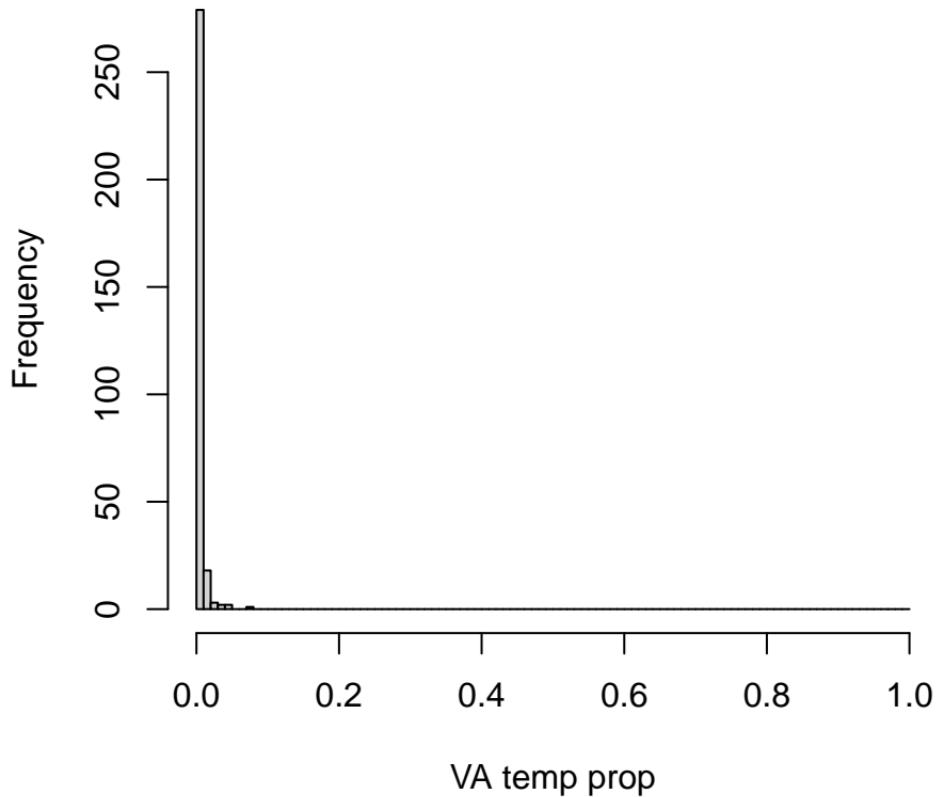
highly-polygenic_2-trait-no-pleiotropy-equal-S_SS-M
1231122



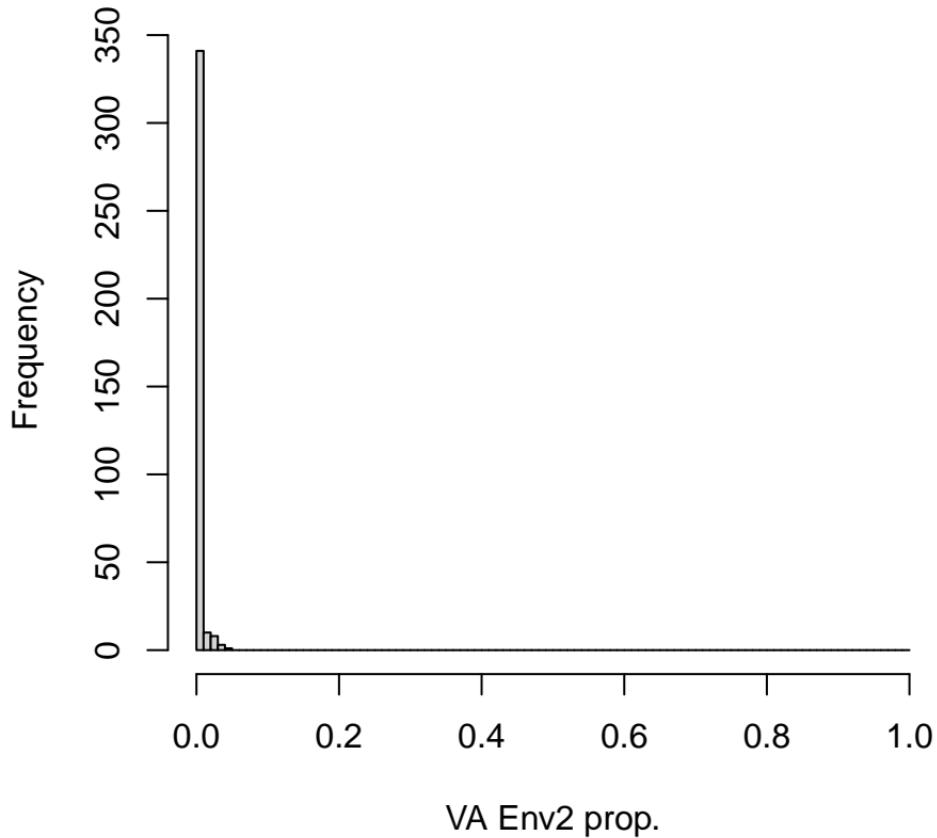
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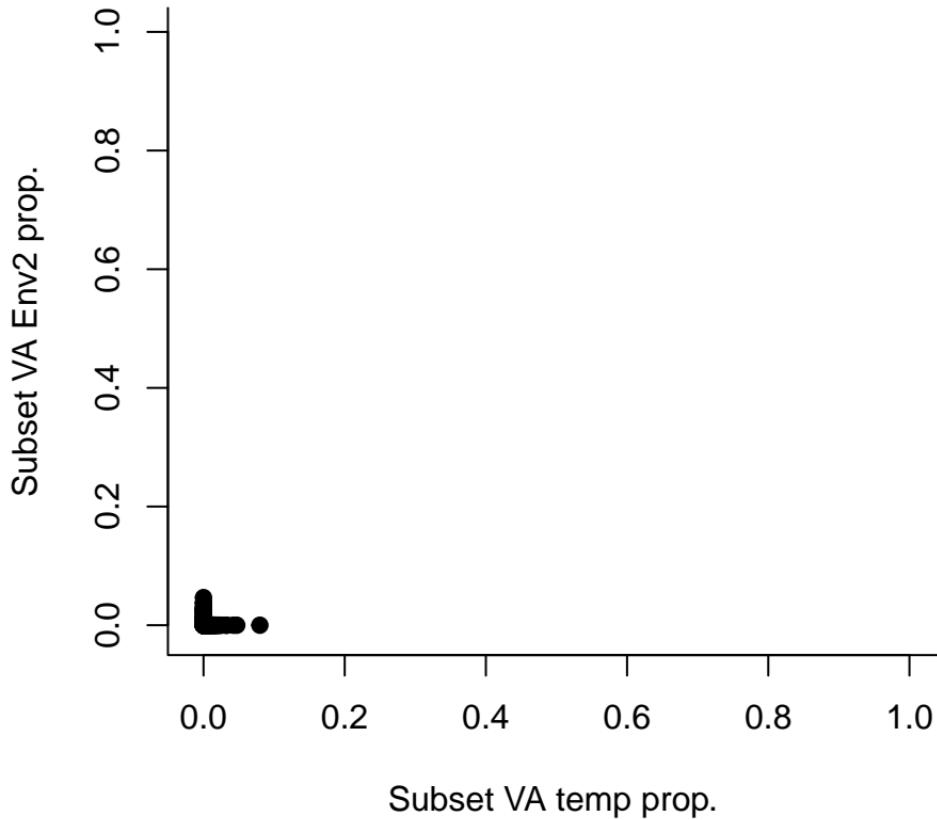
genic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-eq
1231122



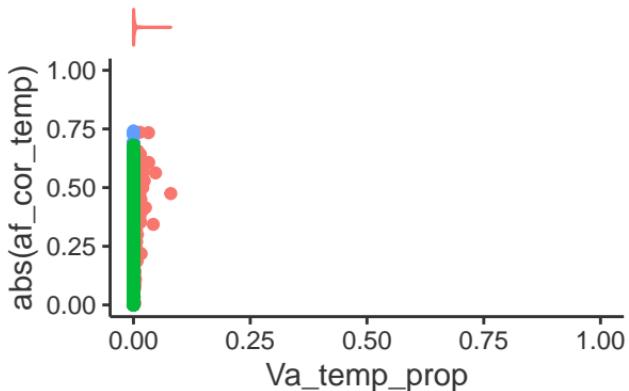
genic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-eq
1231122



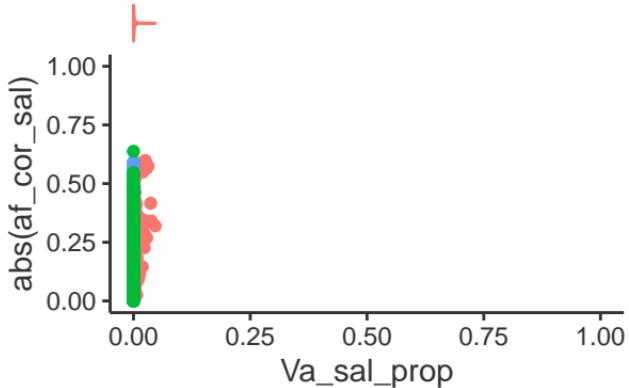
genic_2-trait-no-pleiotropy-equal-S_SS-Mtn_N-eq
1231122



highly-polygenic_2-trait-no-pl
1231122



highly-polygenic_2-trait-no-pl
1231122



highly-polygenic_2-trait-no-pl
1231122

