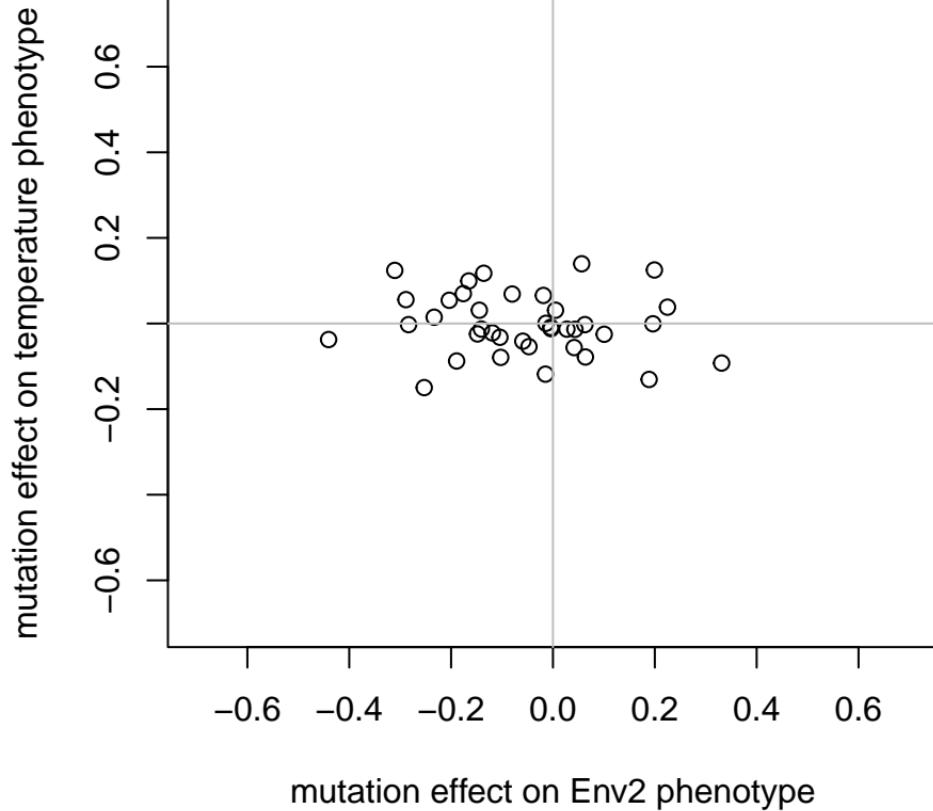
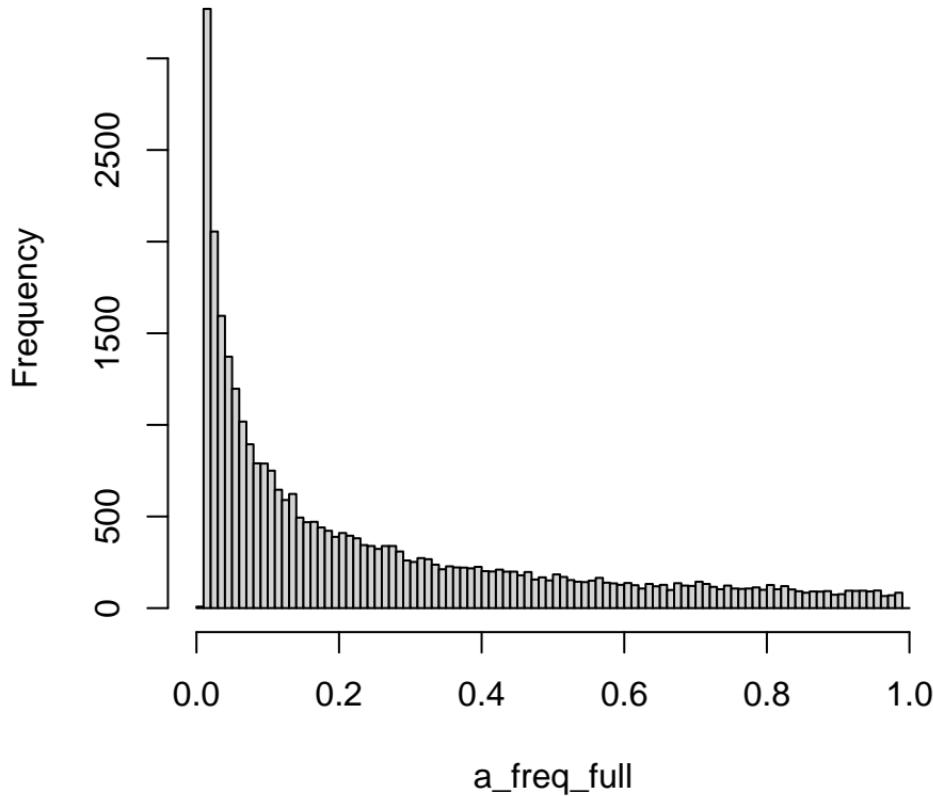


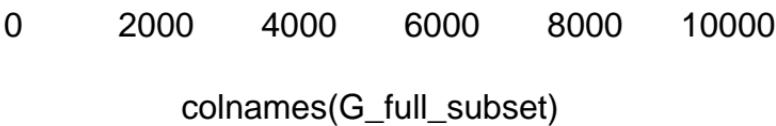
mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-equal_m-constant
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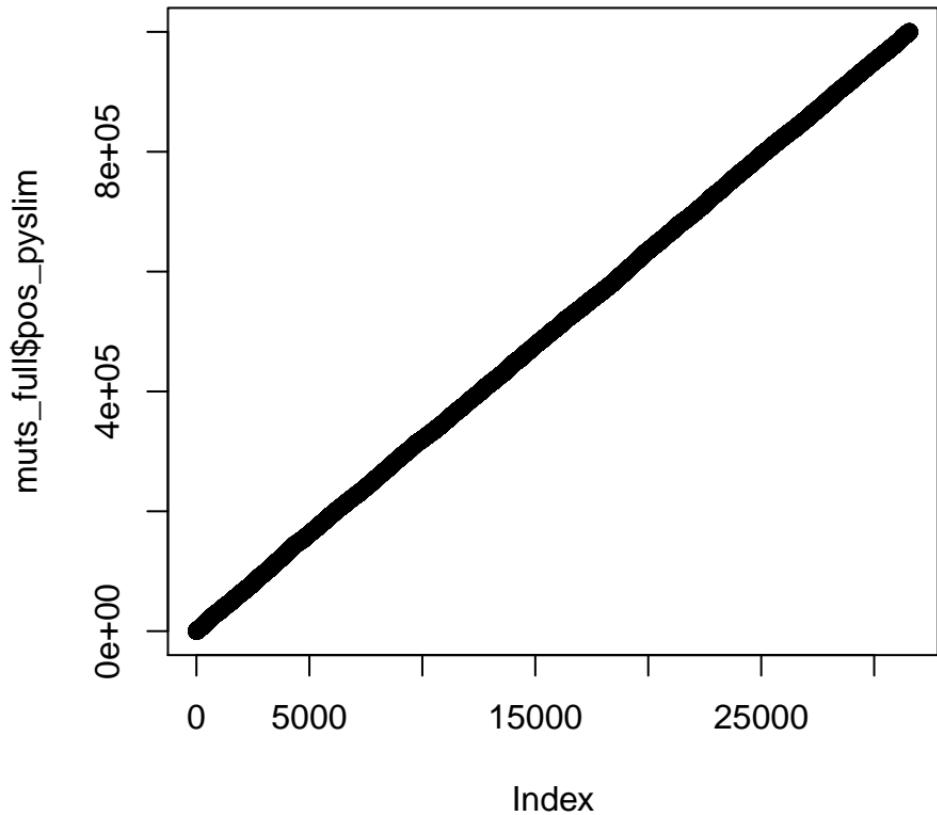


Histogram of a_freq_full

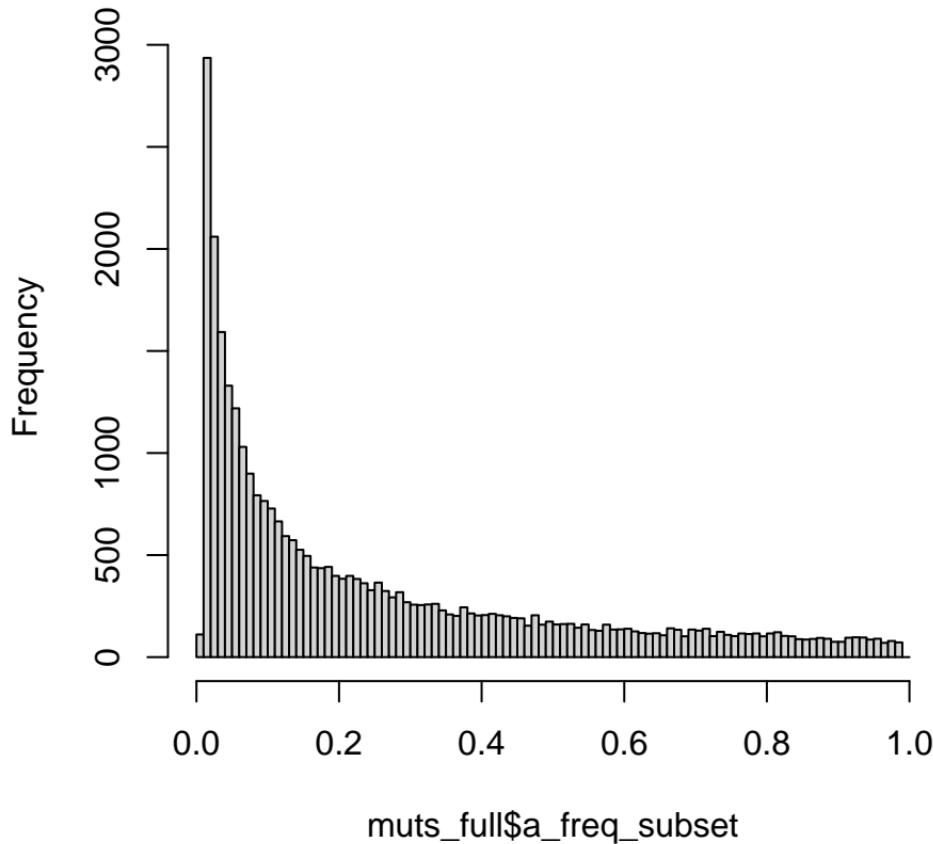


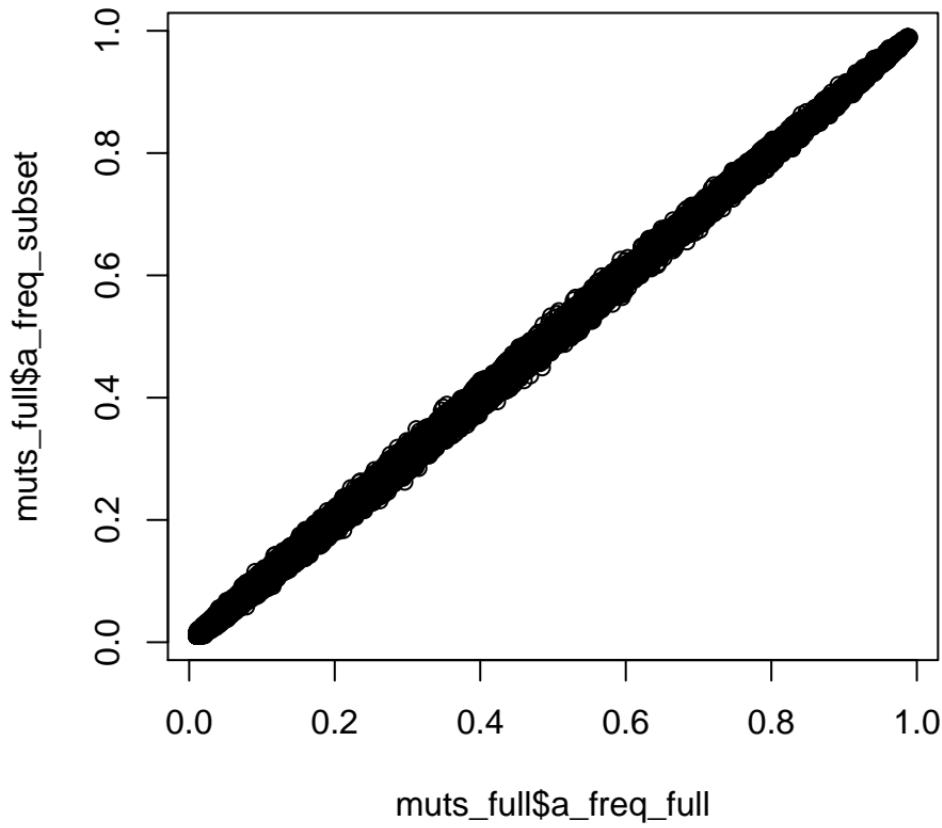
indPhen_df\$indID[which(indPhen_df\$subset)]



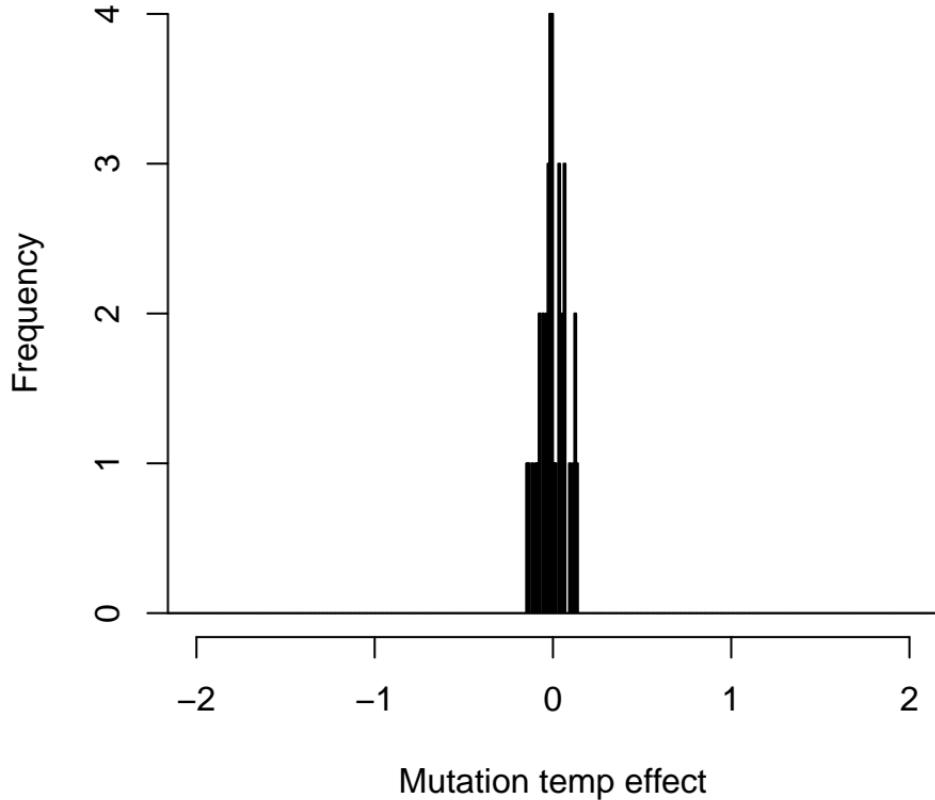


Histogram of muts_full\$a_freq_subset

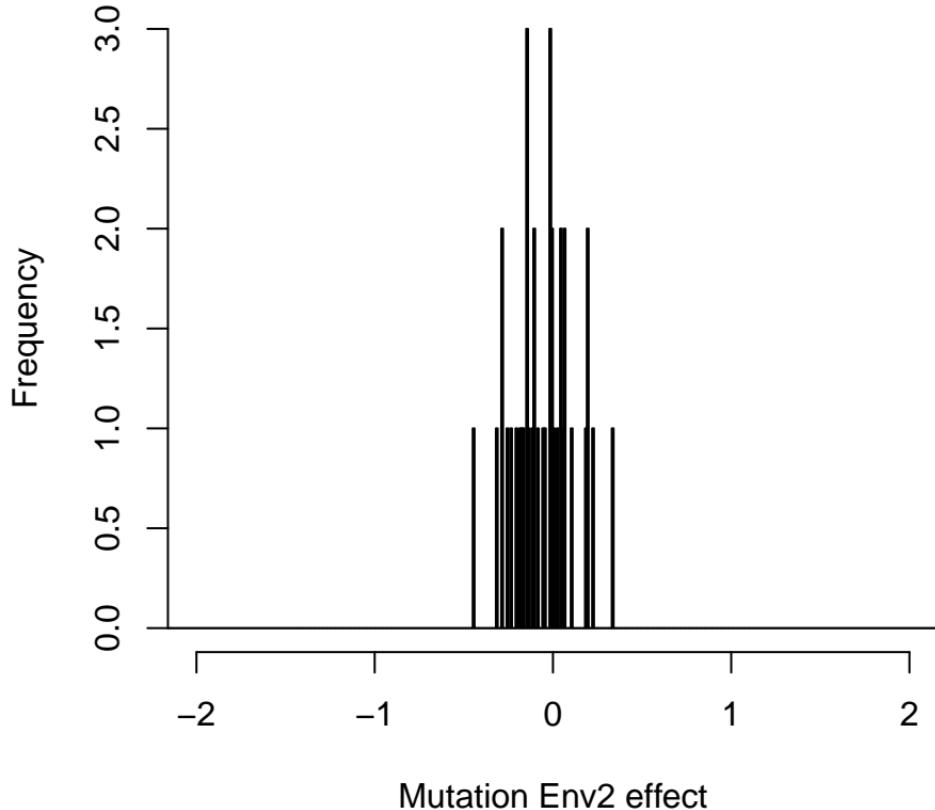




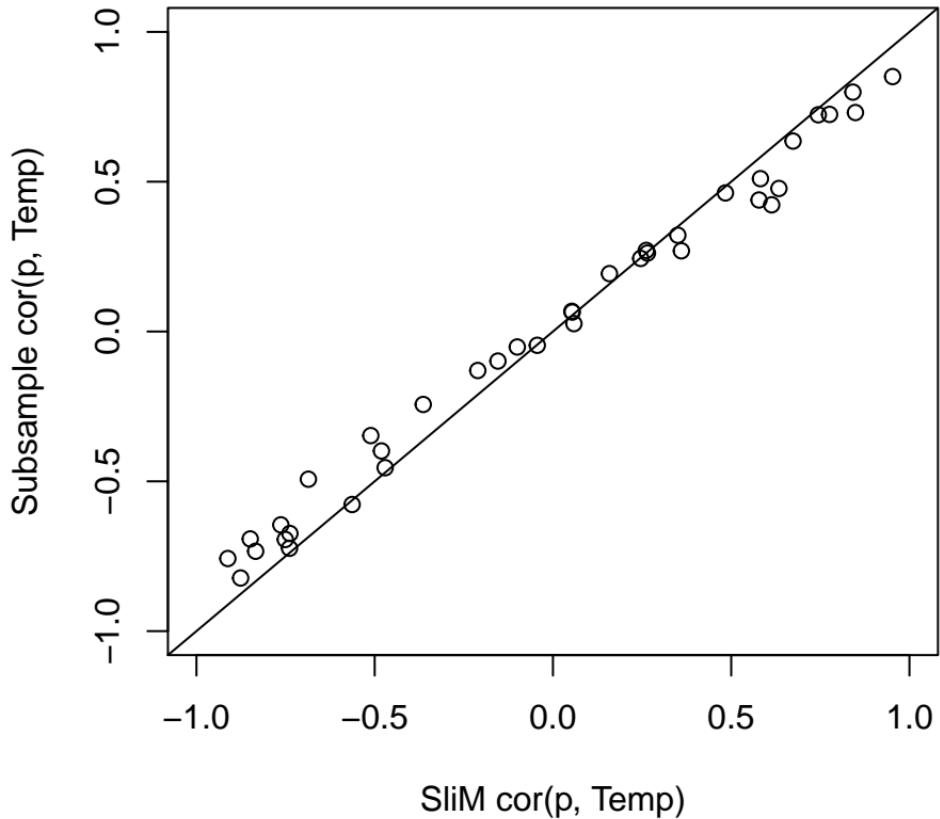
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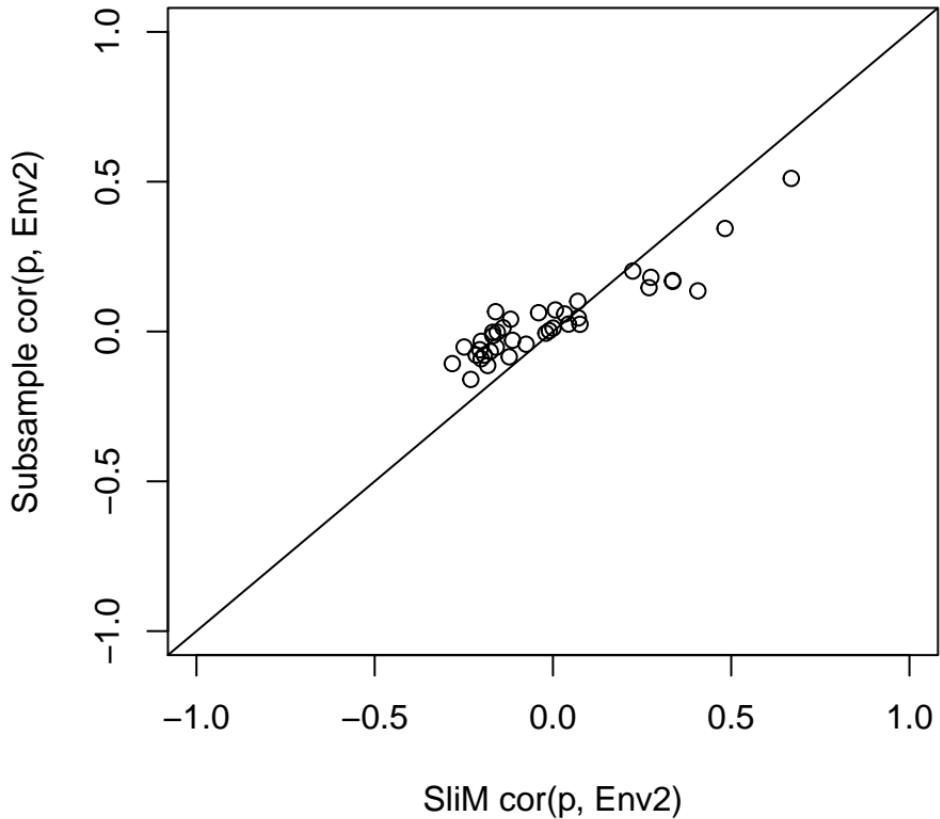
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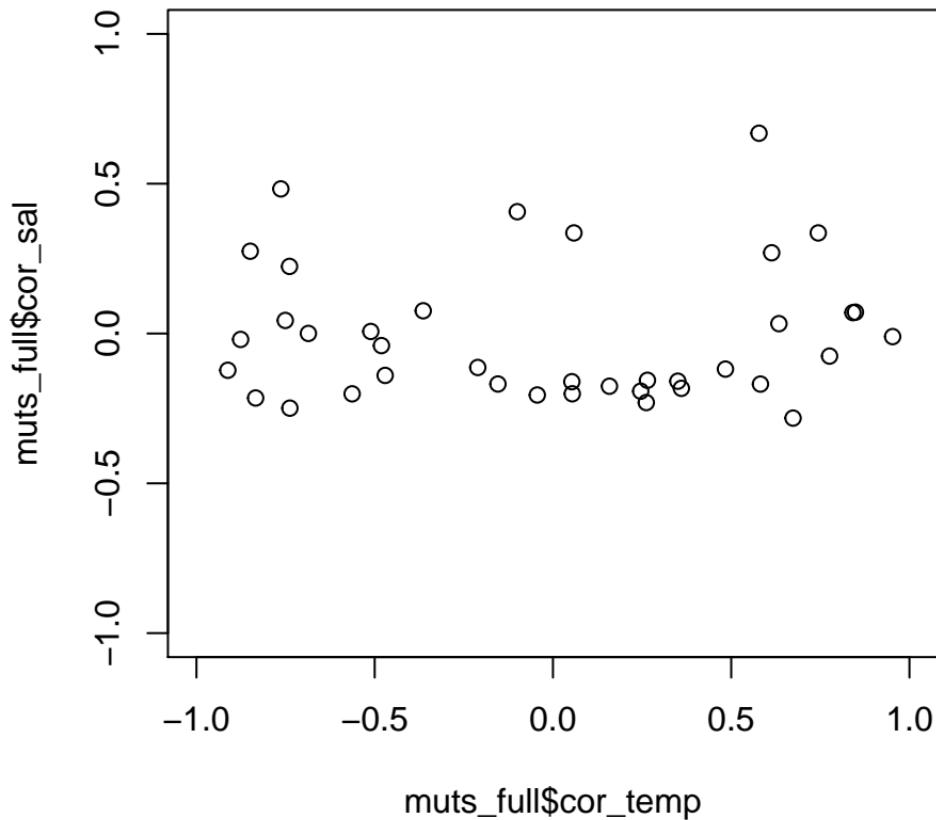
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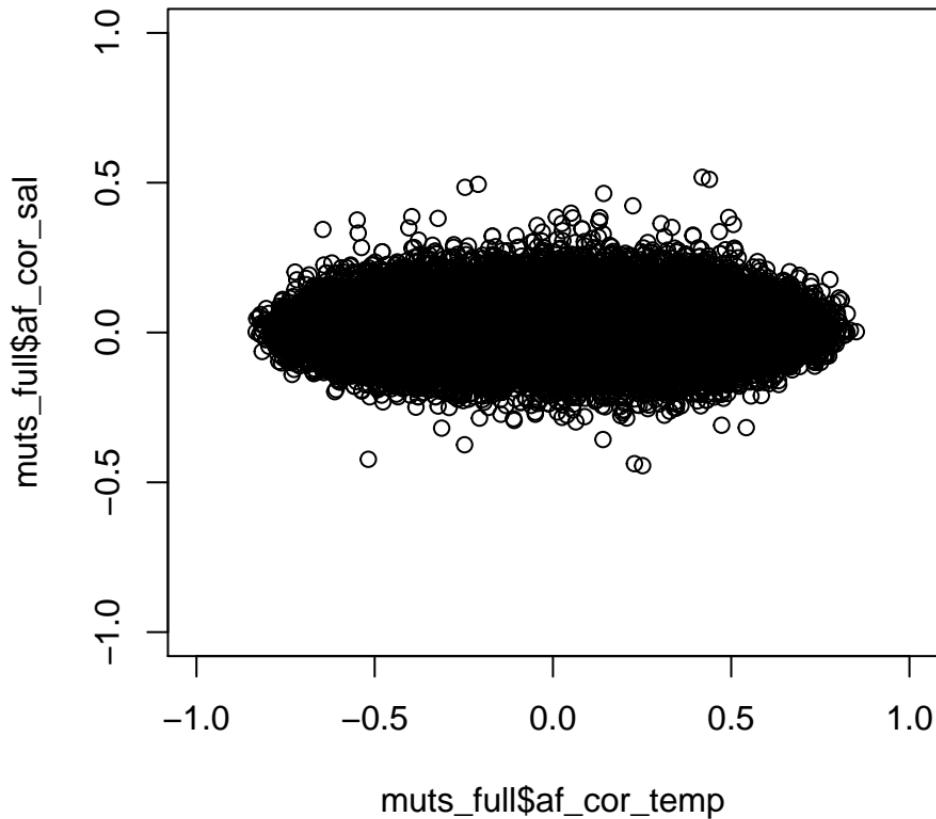
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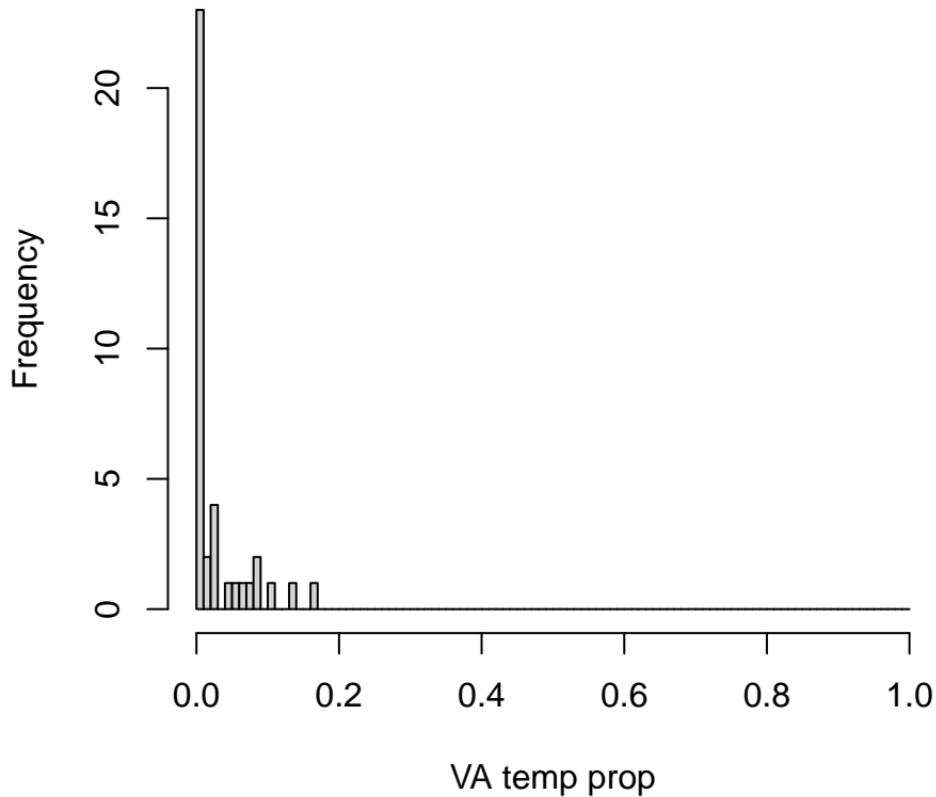
SiM Correlation (AF, ENV) all samples



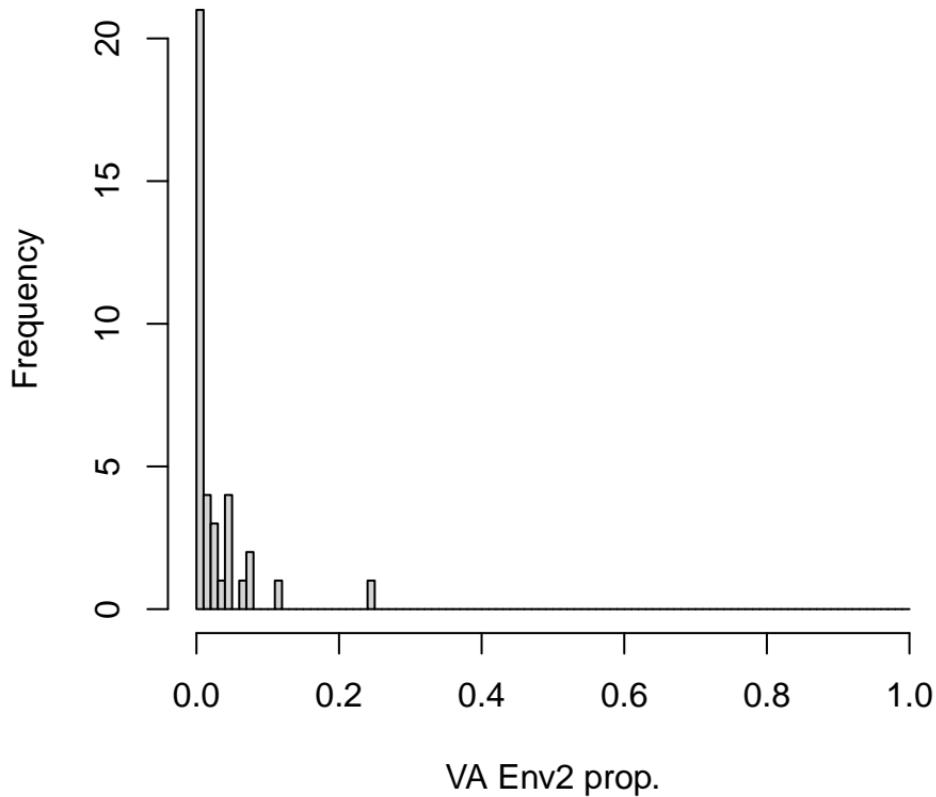
Subsample Correlation (AF, ENV) all samples

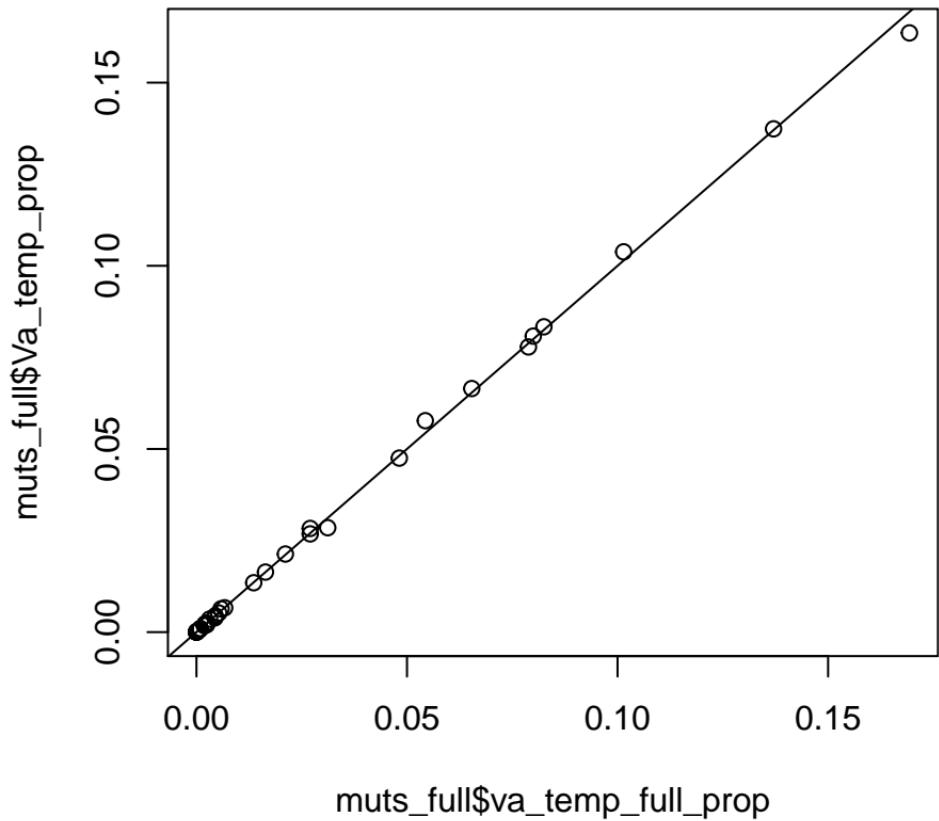


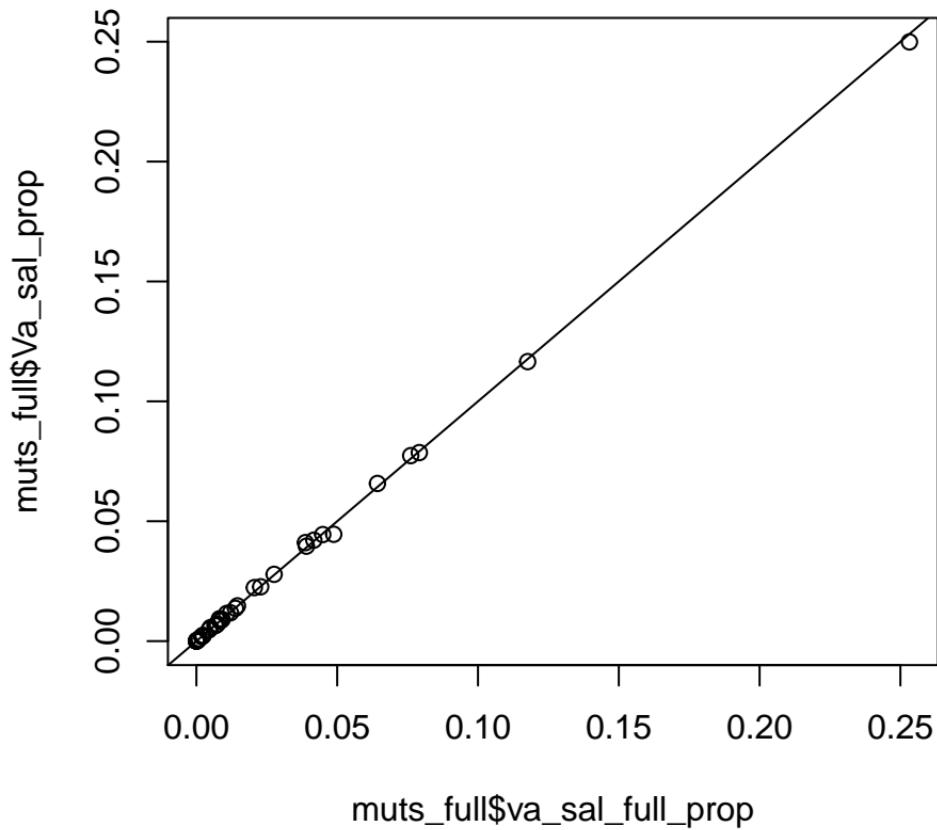
genic_2-trait-pleiotropy-equal-S_Est-Clines_N-equa
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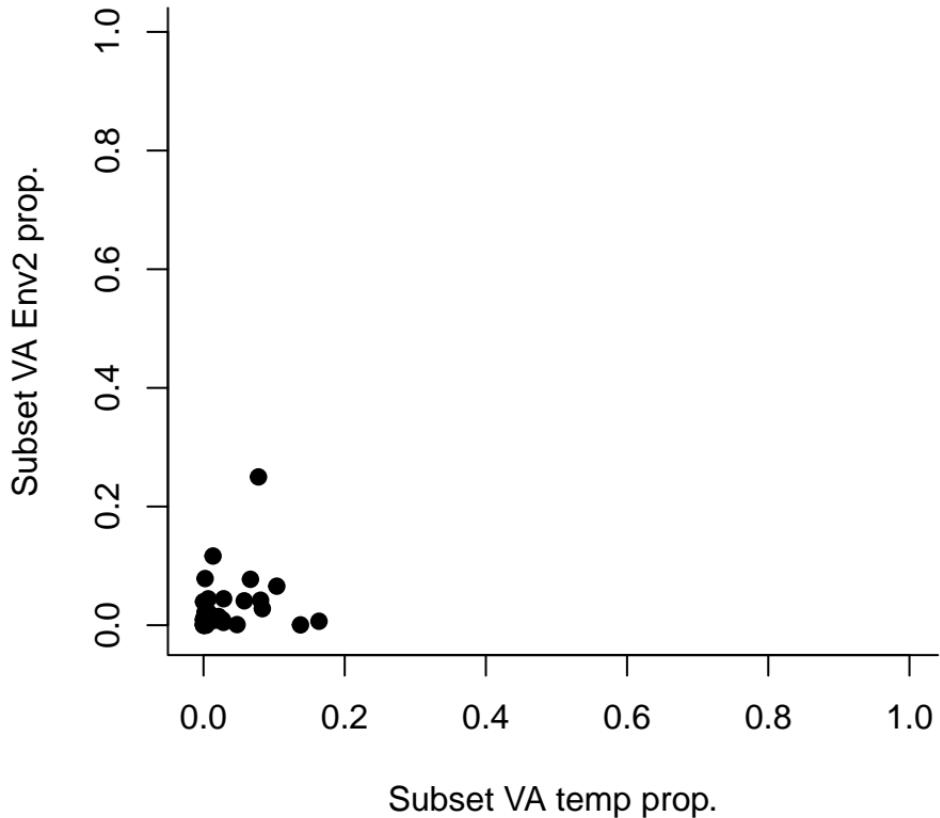
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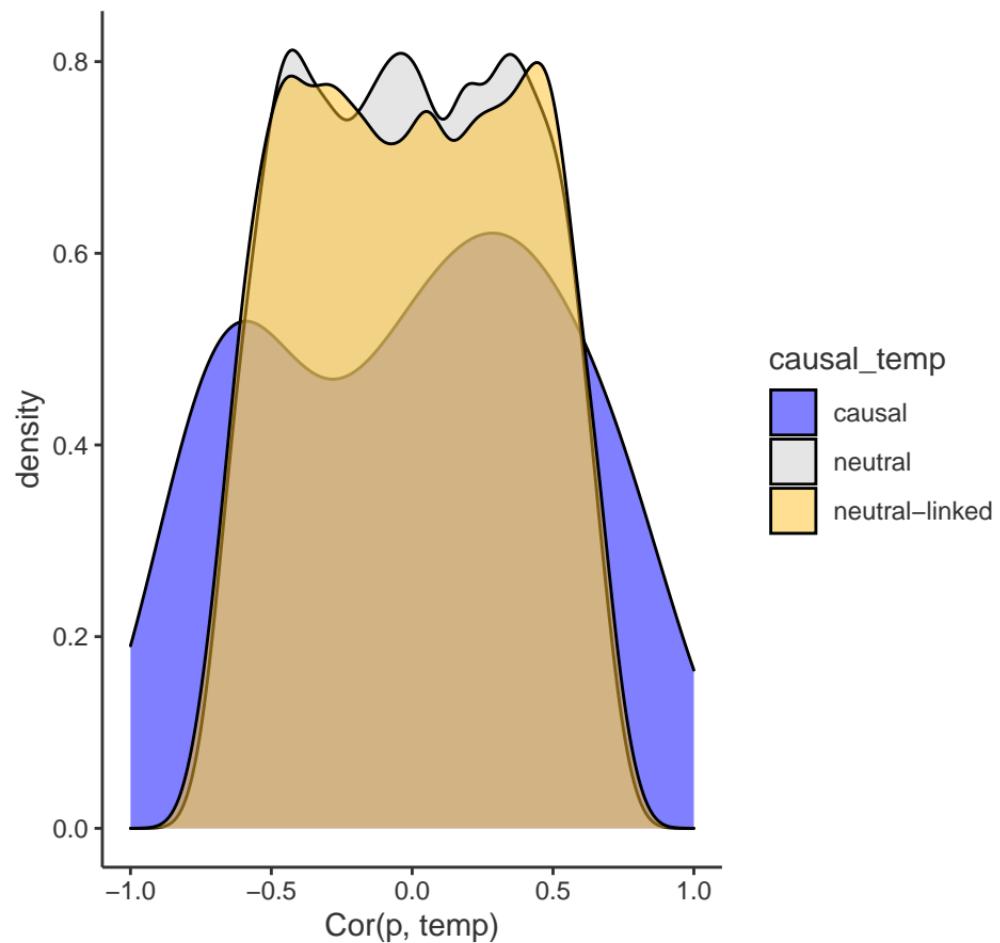




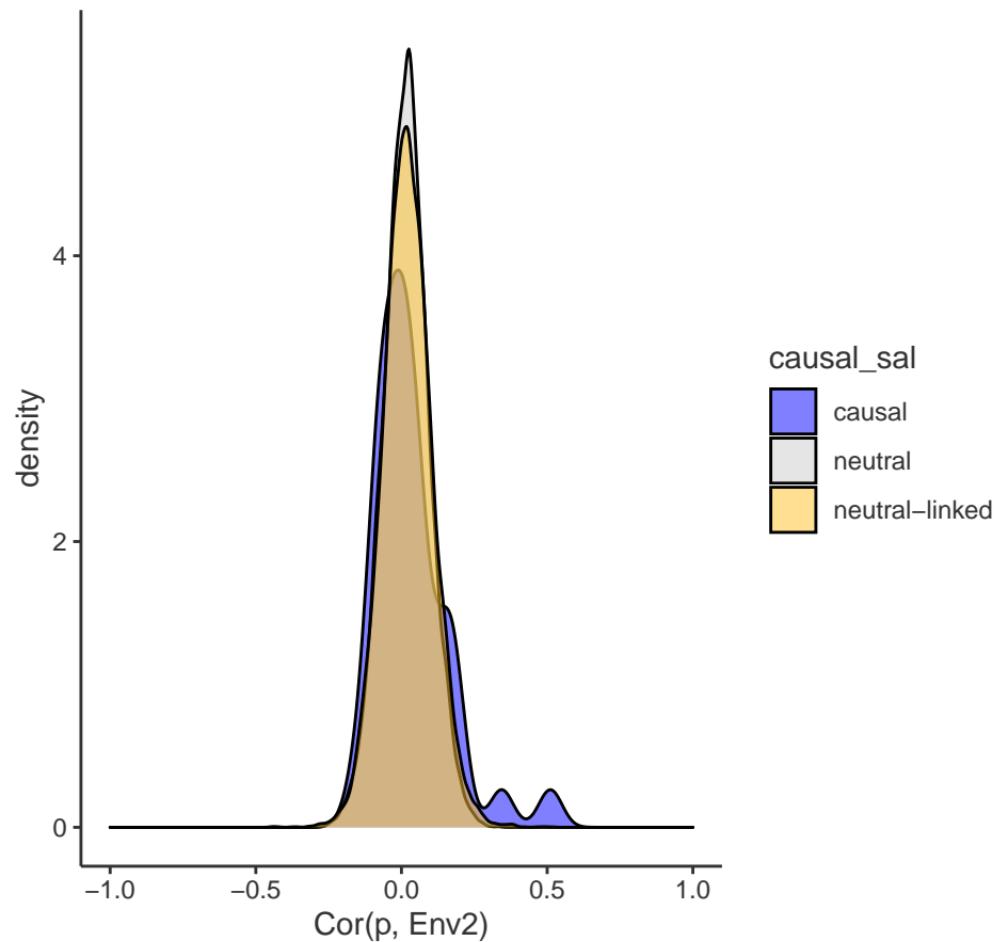
genic_2-trait-pleiotropy-equal-S_Est-Clines_N-equa
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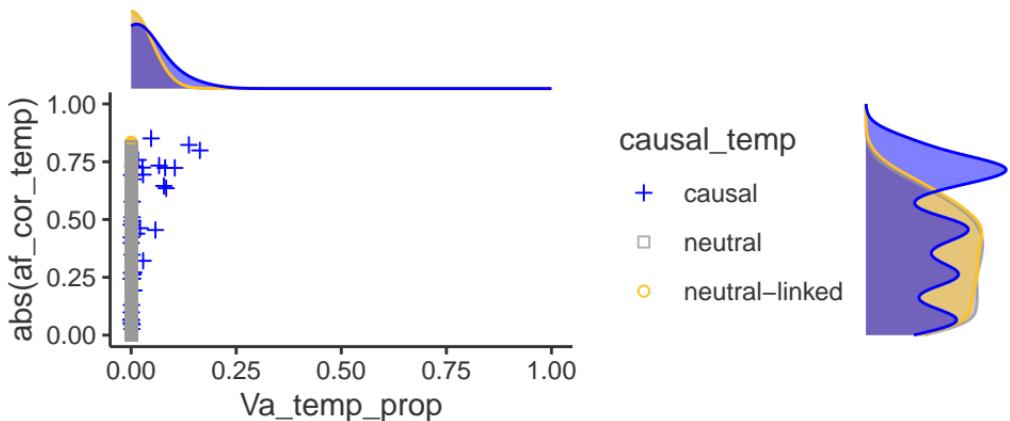
mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-equal_m-constant
1232342



mod-polygenic_2-trait-pleiotropy-equal-S_Est-Clines_N-equal_m-constant
1232342



mod-polygenic_2-trait-pleiotropy-equal-S__
1232342



mod-polygenic_2-trait-pleiotropy-equal-S__
1232342

