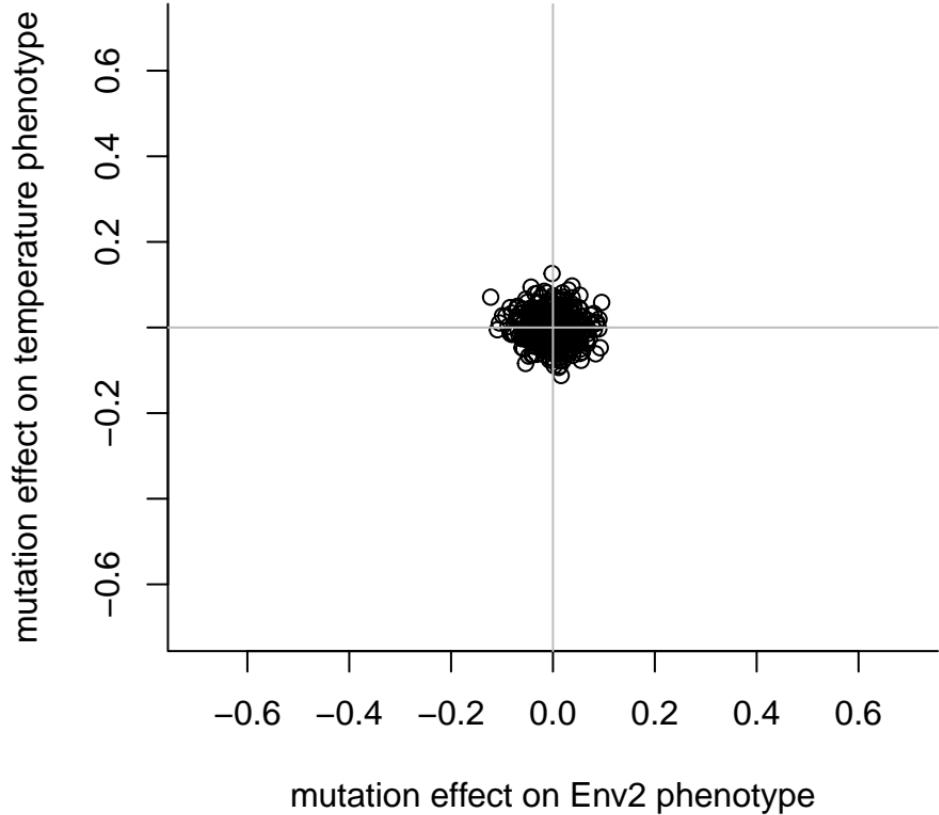
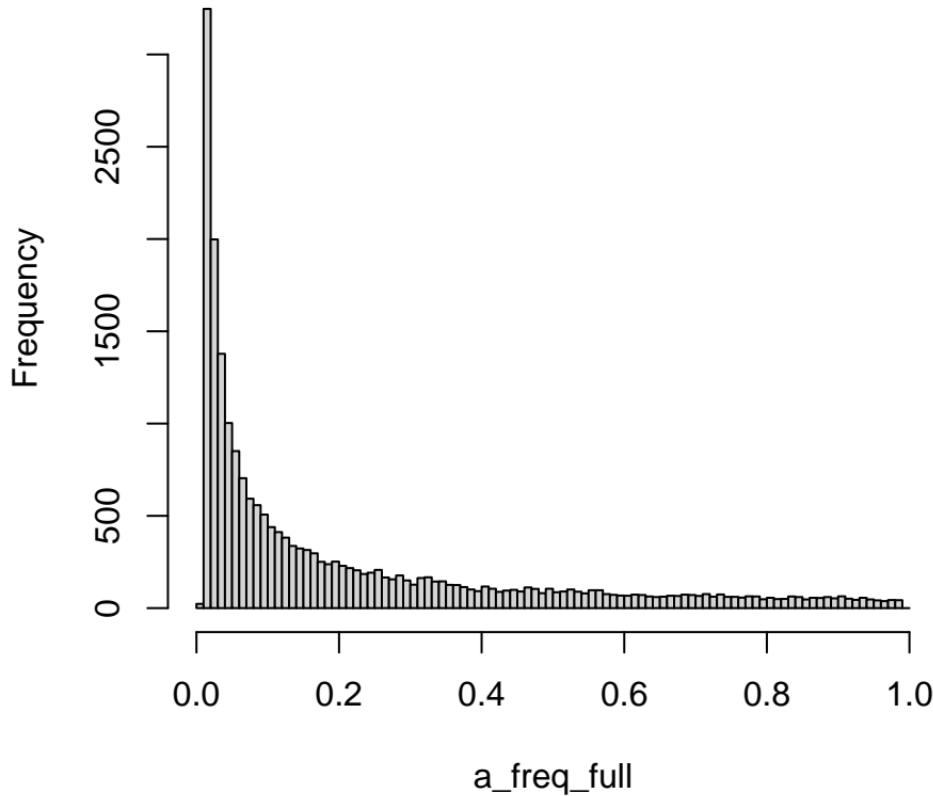


polygenic_2-trait-pleiotropy-equal-S_SS-Clines_N-equal_m-constant
1231222



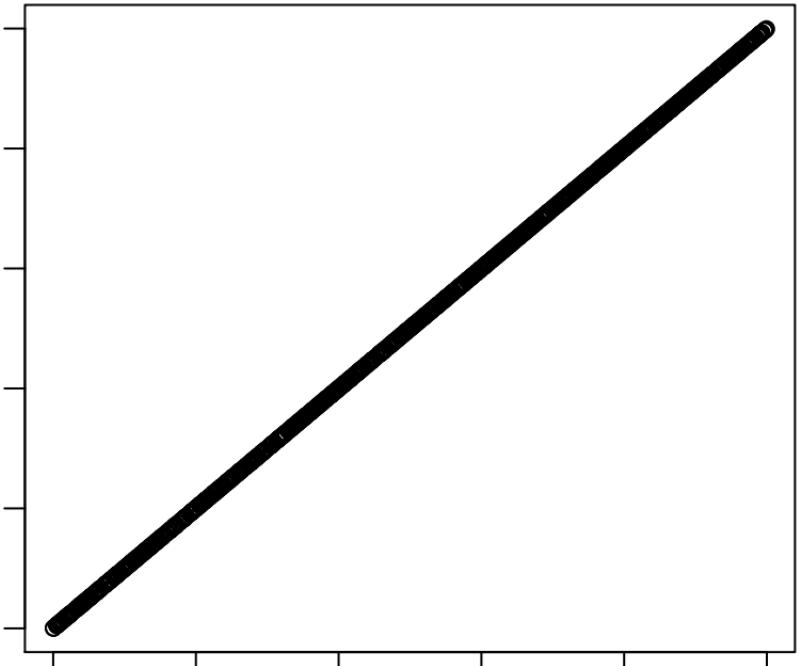
Histogram of a_freq_full

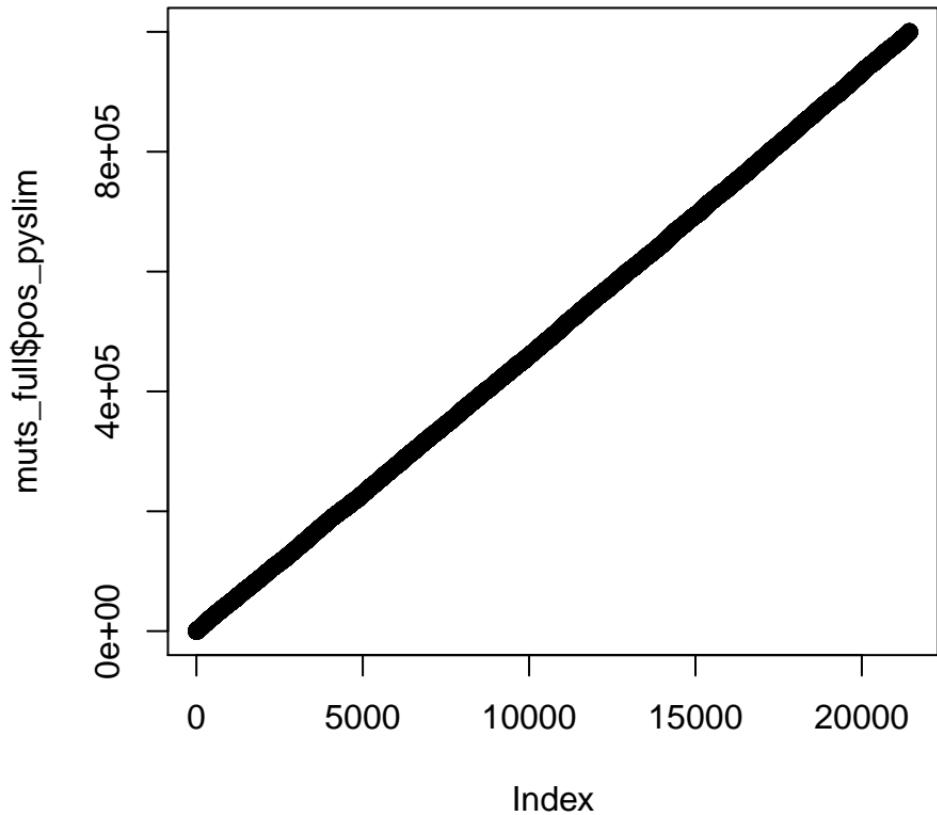


indPhen_df\$indID[which(indPhen_df\$subset)]

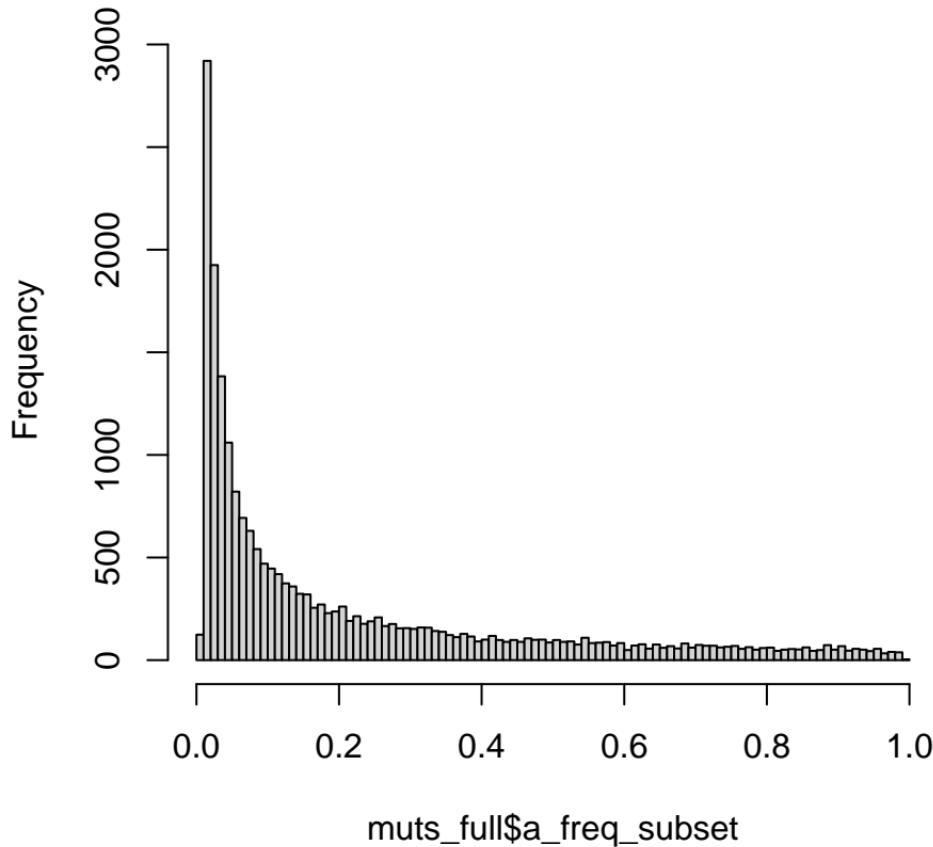
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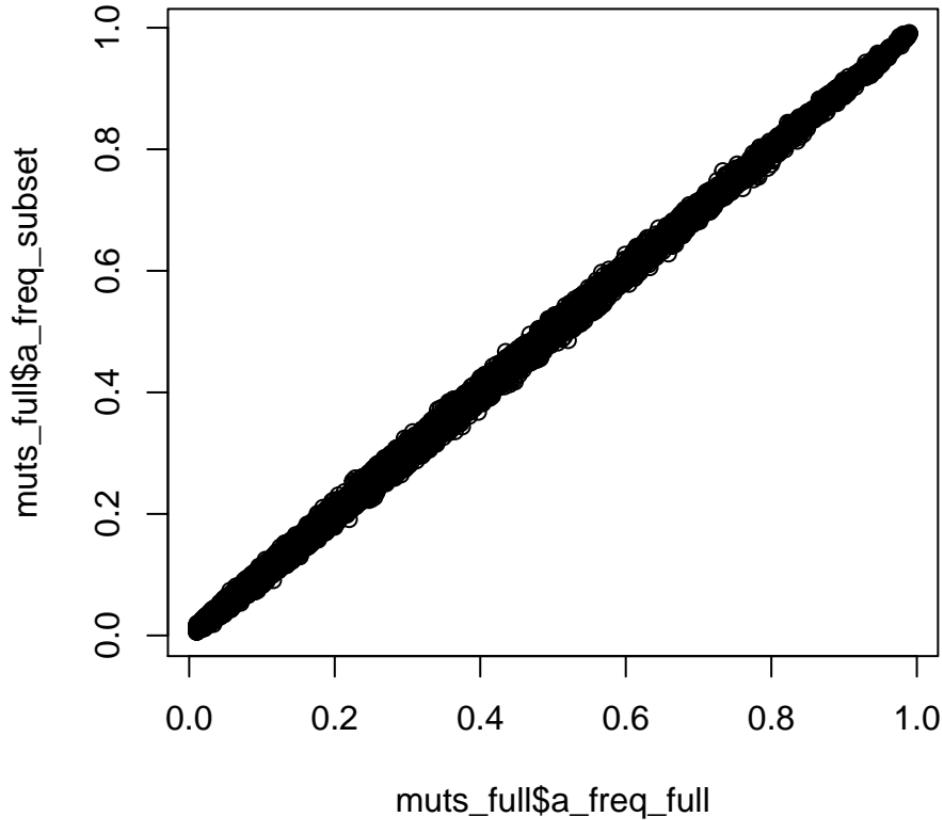
colnames(G_full_subset)



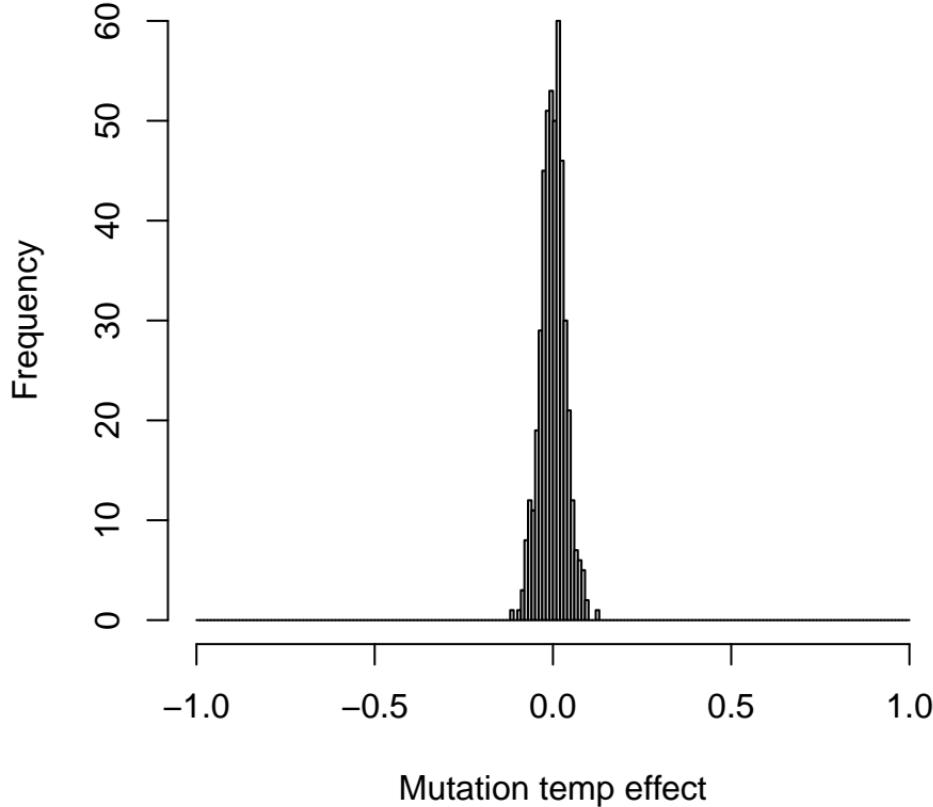


Histogram of muts_full\$a_freq_subset

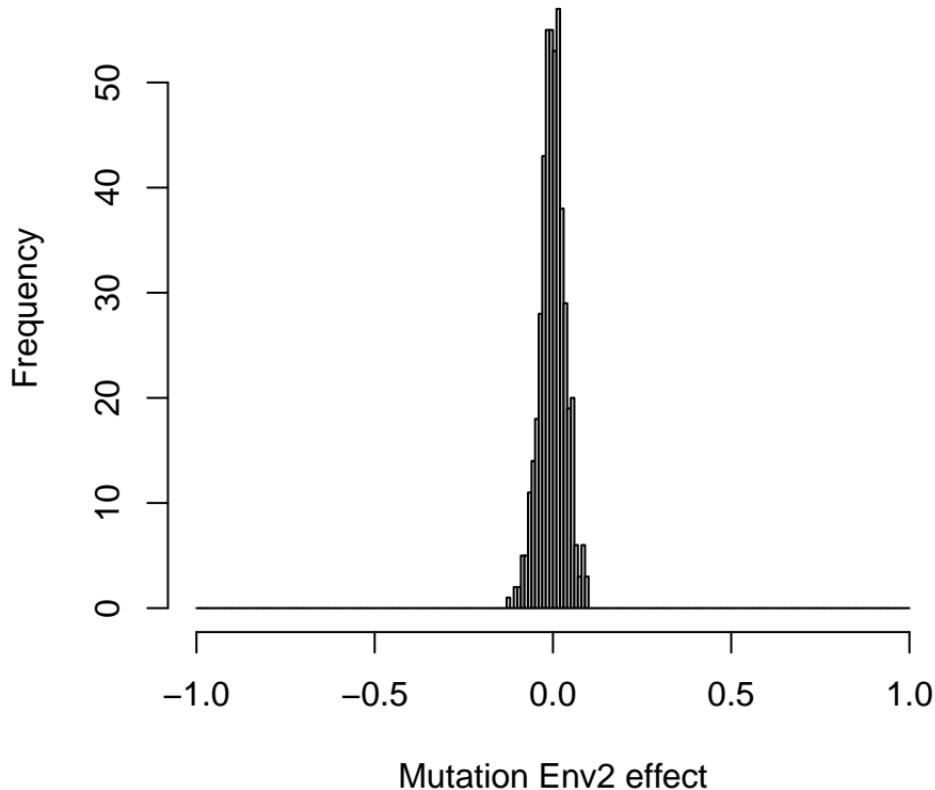




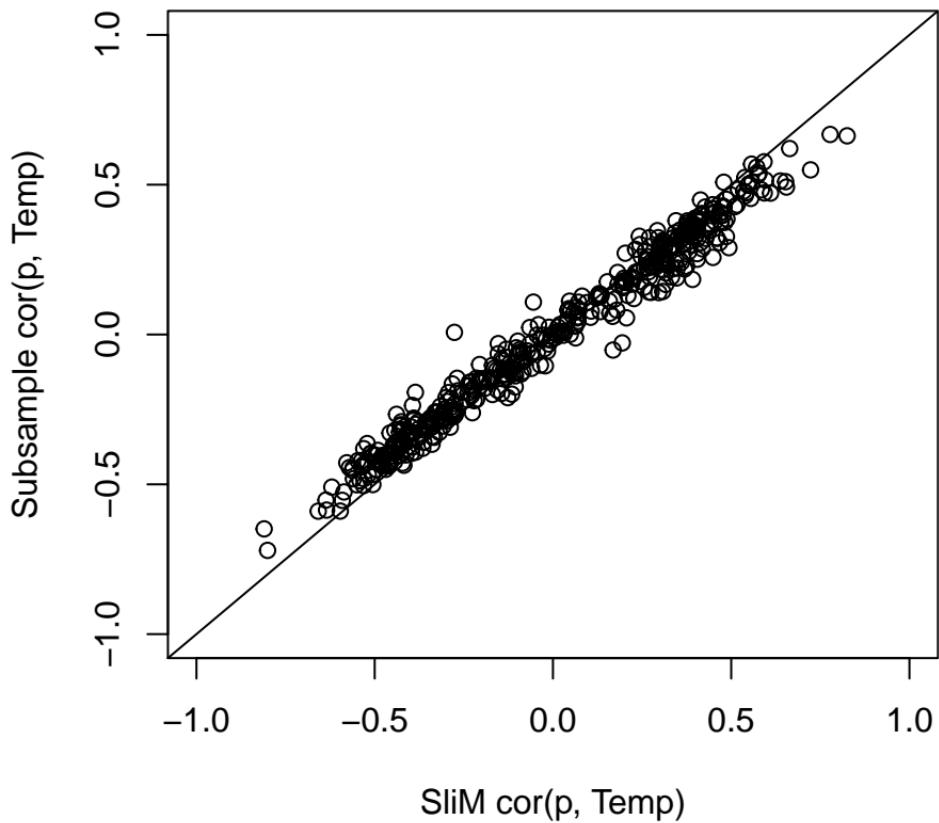
polygenic_2-trait-pleiotropy-equal-S_SS-Clines_N-equal_m-constant
1231222



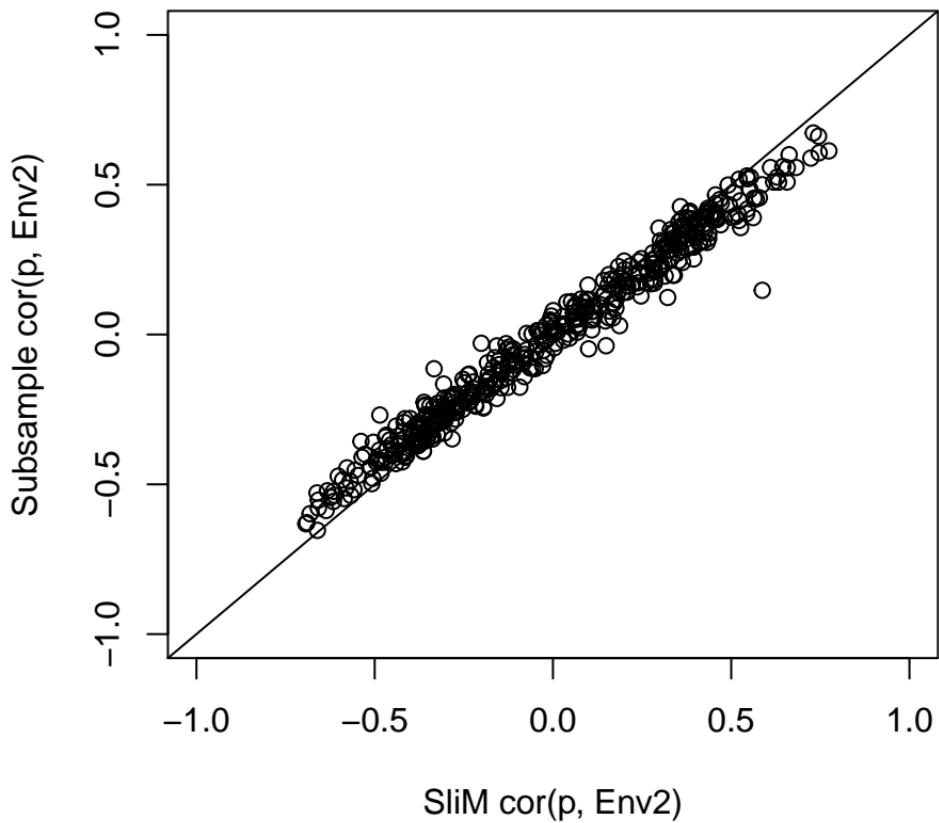
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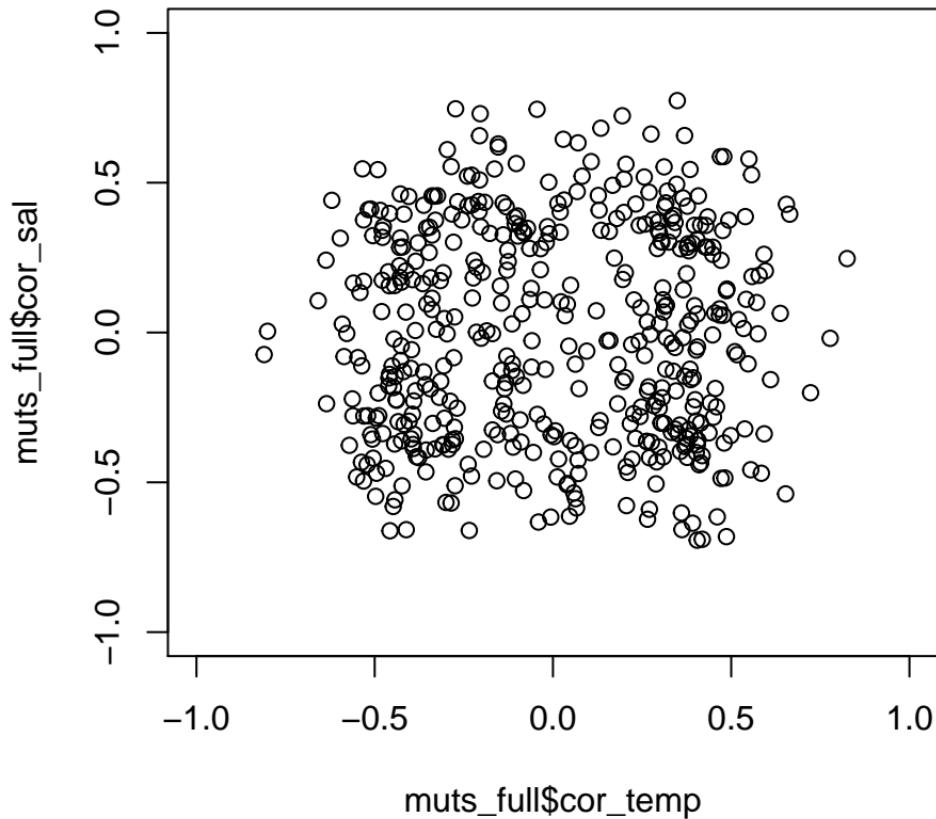
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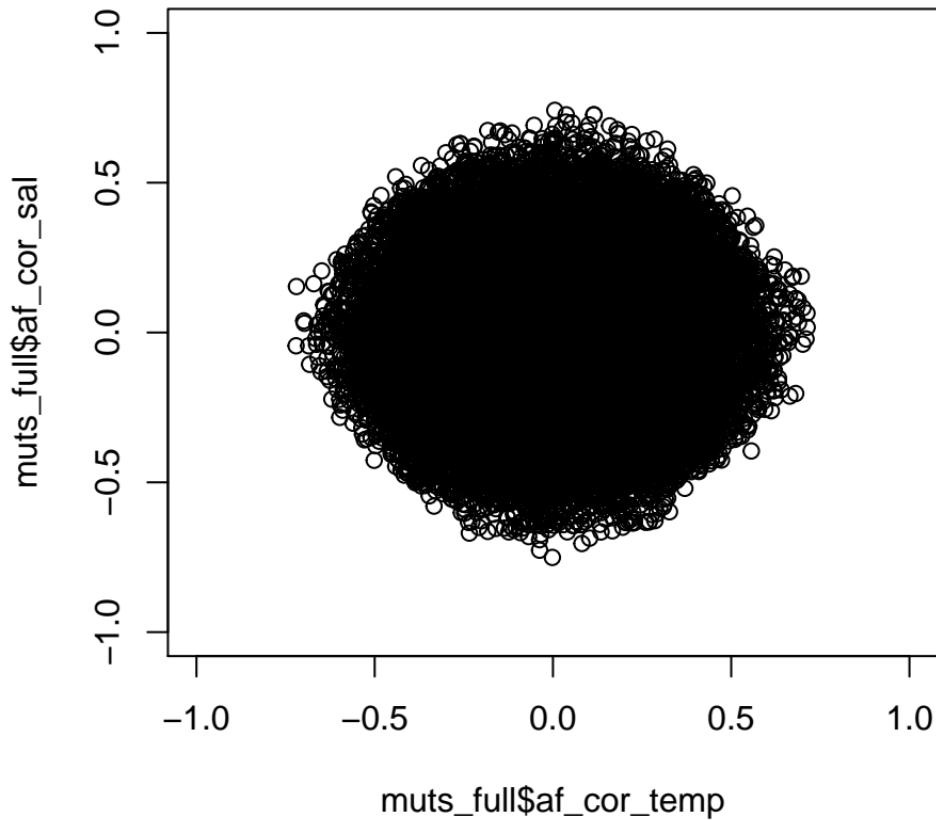
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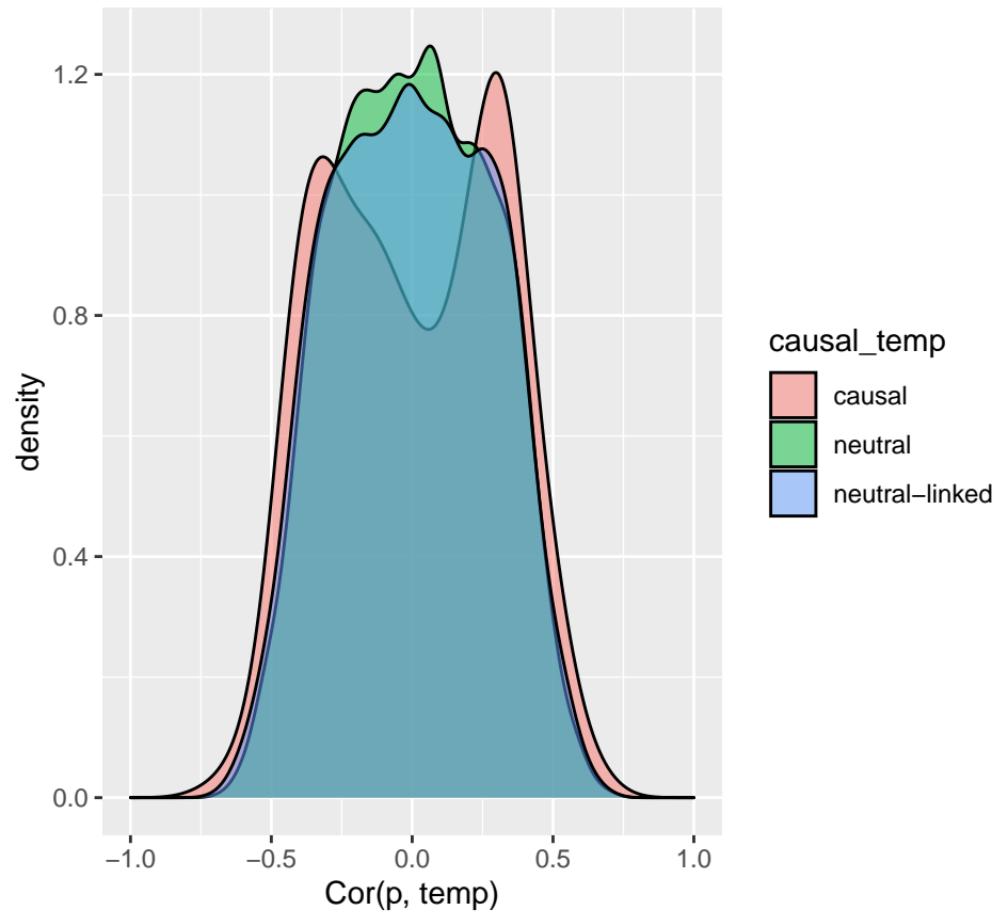
SiM Correlation (AF, ENV) all samples



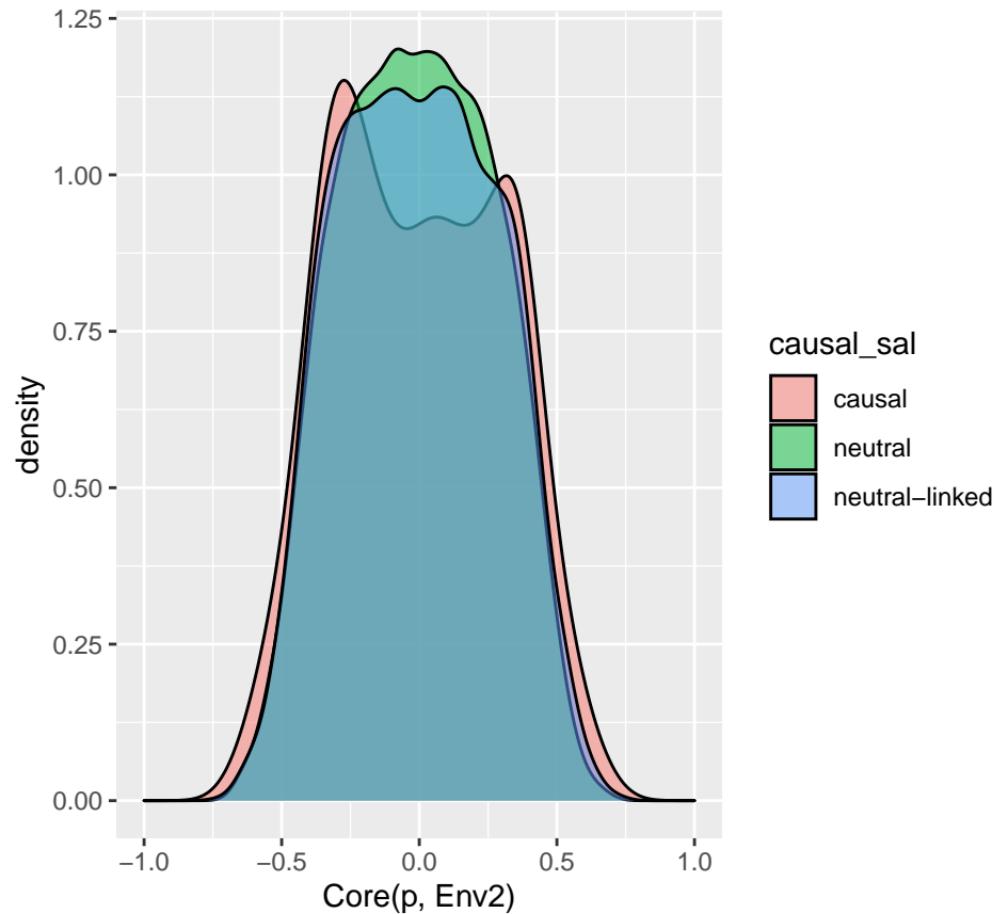
Subsample Correlation (AF, ENV) all samples



polygenic_2-trait-pleiotropy-equal-S_SS-Clines_N-e
1231222

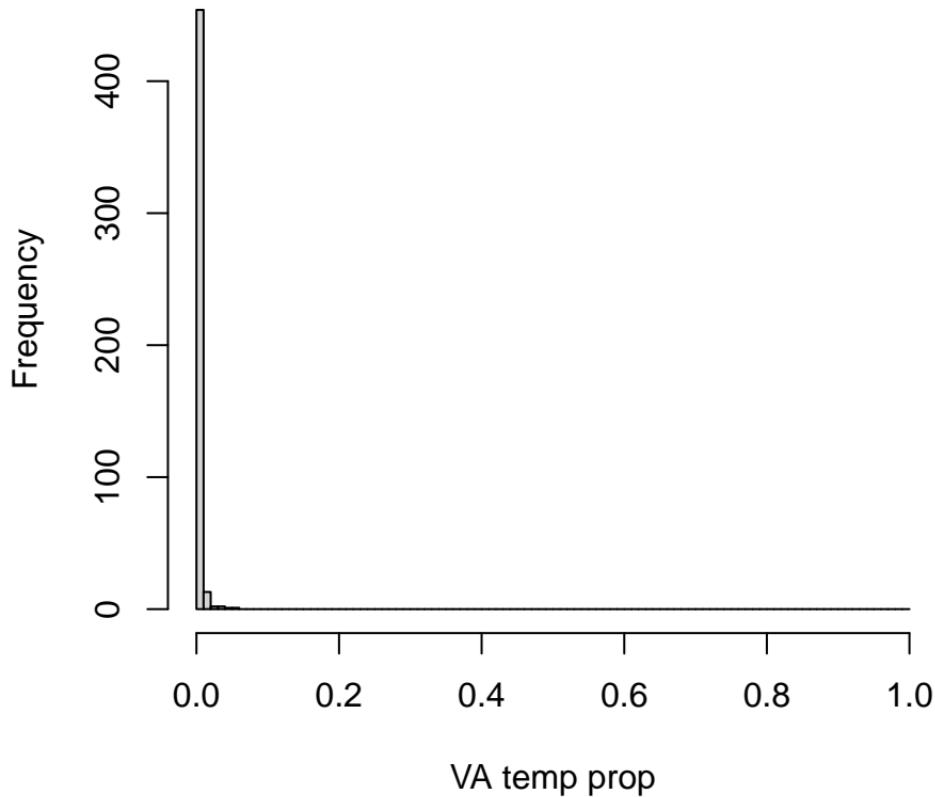


polygenic_2-trait-pleiotropy-equal-S_SS-Clines_N-
1231222



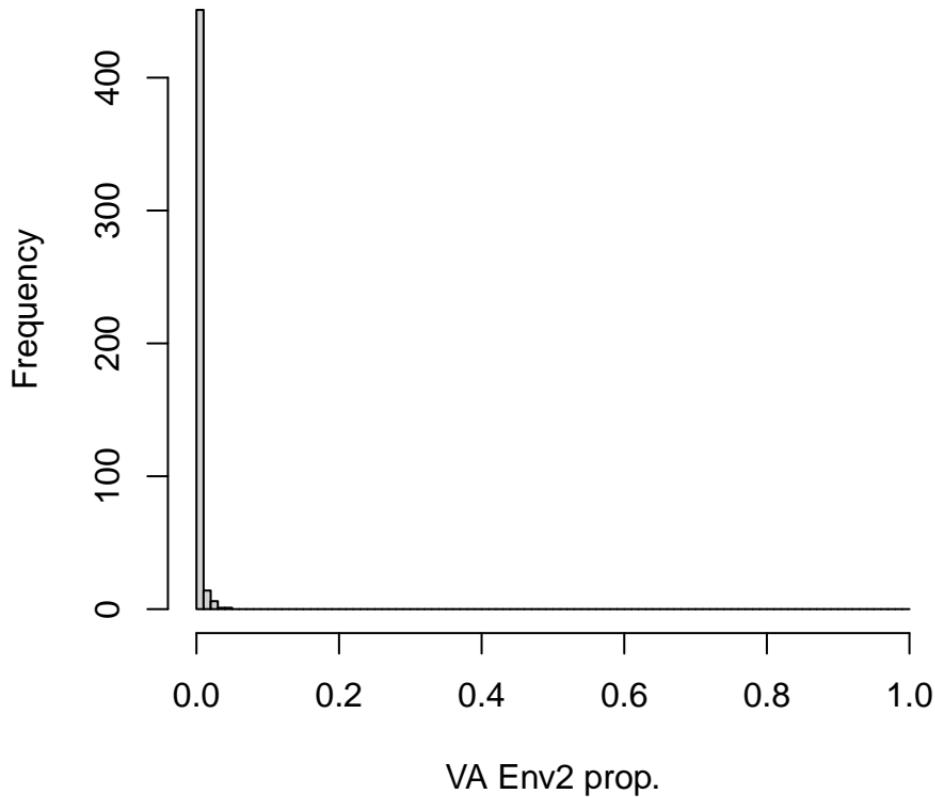
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1231222

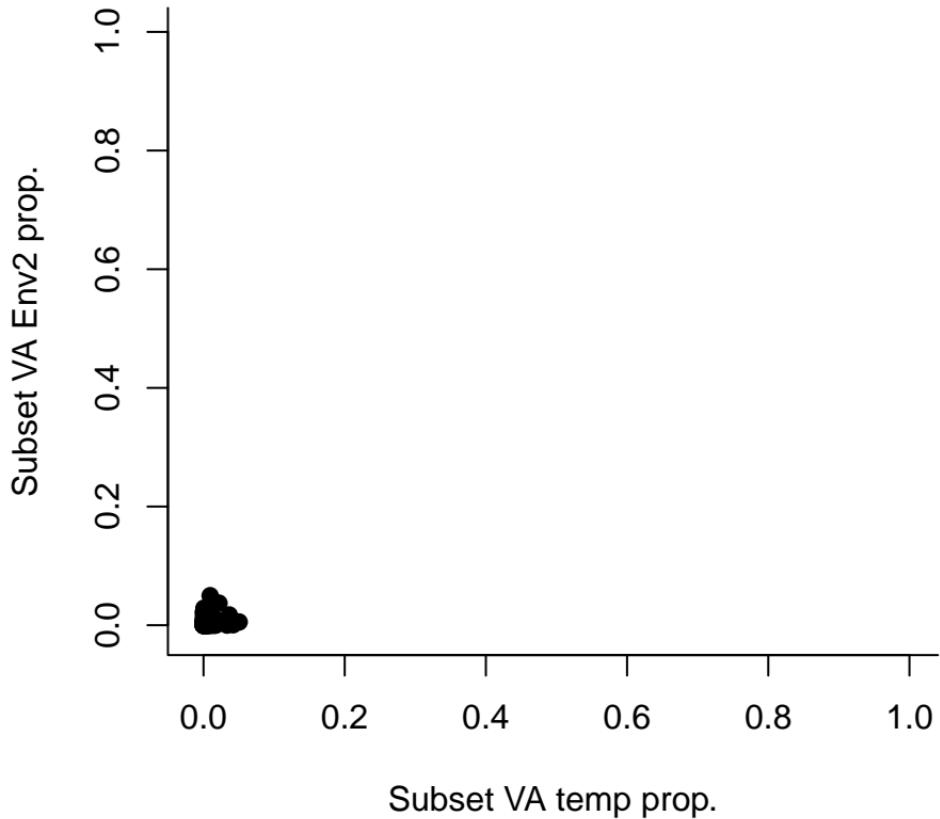


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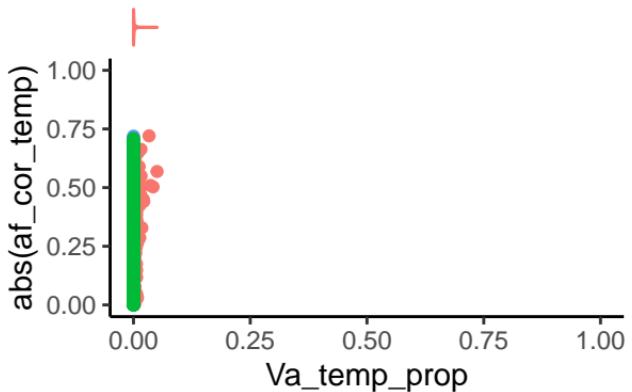
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nic_2-trait-pleiotropy-equal-S_SS-Clines_N-equal_1231222

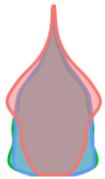


polygenic_2-trait-pleiotropy-ec
1231222

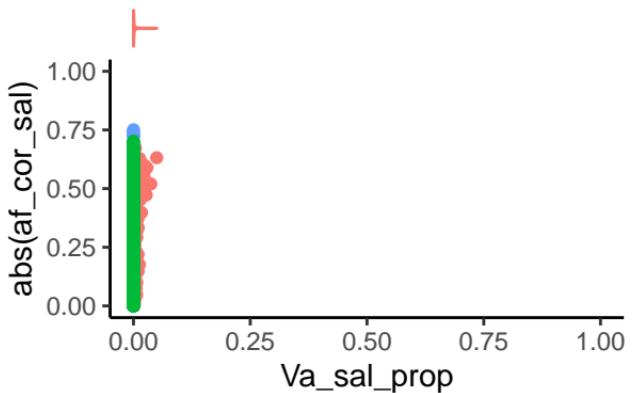


causal_temp

- causal
- neutral
- neutral-linked



polygenic_2-trait-pleiotropy-ec
1231222



causal_sal

- causal
- neutral
- neutral-linked

