





highly-polygenic_2-trait-no-pleiotropy-equal-S__Est-Clines_N-variable_m-variable 1231113 mutations - Env2 0.3 Env2 0.2 Env2 VA prop. 0.02 0.04 0.1 0.06 RDA 2 mut Env2 effect Temp 0.0 0.1 0.0 -0.1 -0.1 -0.2 -0.25 0.00 0.25 0.50 RDA 1

