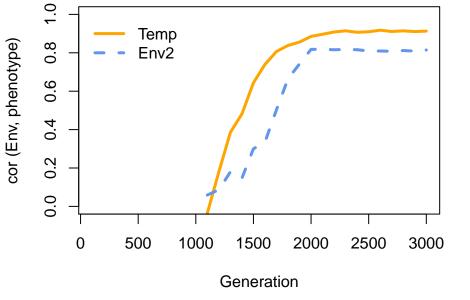
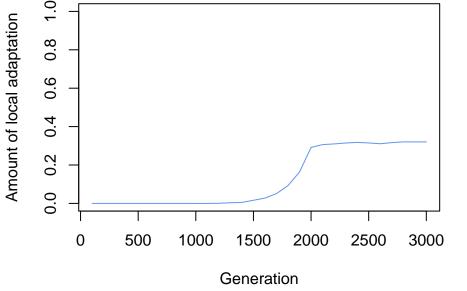


2–trait–no–pleiotropy–unequal–S__Est–Clines_N–clin 1231125

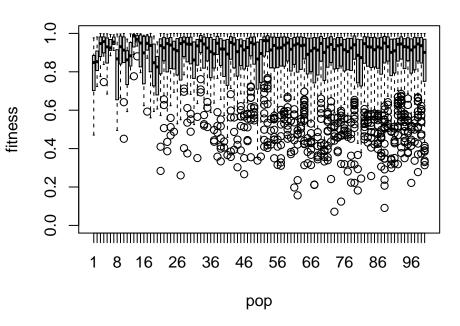


2–trait–no–pleiotropy–unequal–S__Est–Clines_N–clin 1231125

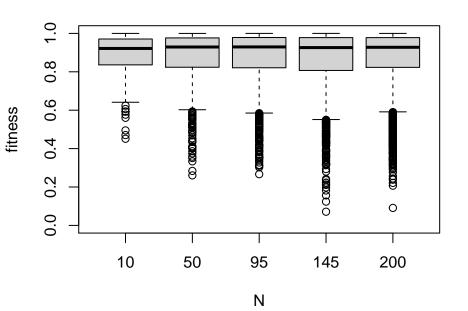


2-trait-no-pleiotropy-unequal-S__Est-Clines_N-clin 1231125 Mean metapopulation phenotype 0.5 0.0 -0.50 500 1000 1500 2000 2500 3000

Generation







subset

