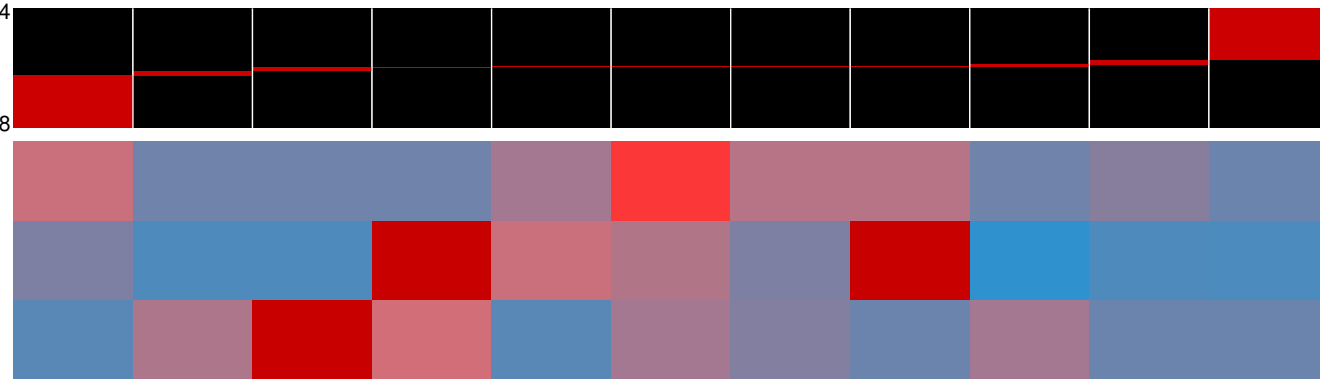




5.34
-5.58



GNF2_MSH2

MORF_ACTG1

MORF_RPA2