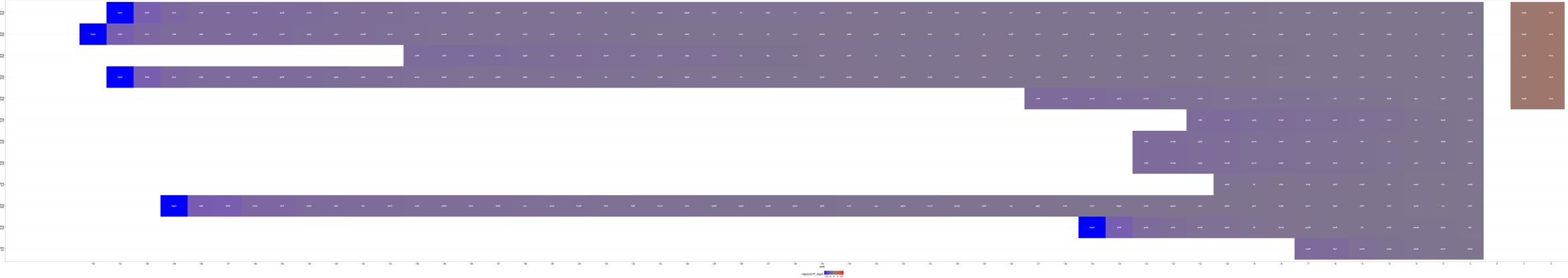
genes_P0.05Fdr02_protein_sT_set00_StcNasMgh_S_baseMaybgHdg_baseNa_Exp_noFilter_p15f_noNorm__Mg_mM_Levels_mf
metal ion binding
pval(3.3e-6)fdr(0.74) *12 *13 640
ion binding
pval(3.2e-6)fdr(0.74) *11 *13 640
transition metal ion binding
pval(3.2e-6)fdr(0.74) *11 *12 640
cation binding
pval(3.2e-6)fdr(0.74) *11 *12 654
iron ion binding
pval(3.8e-6)fdr(0.0001) *11 *12 230
4 iron, 4 sulfur cluster binding
pval(3.8e-6)fdr(0.0001) *11 *12 227
iron sulfur cluster binding
pval(3.8e-6)fdr(0.0001) *11 *12 243
metal cluster binding
pval(3.8e-6)fdr(0.0001) *11 *12 243
manganese ion binding
pval(3.6e-6)fdr(0.74) *11 *12 57
nucleoside binding
pval(3.6e-6)fdr(0.0101) *9 *10 *11 660
magnesium ion binding
pval(3.2e-6)fdr(0.74) *10 *11 *12 212
hydrolase activity, acting on carbon-nitrogen (but not peptide) bonds, in linear amidines
pval(3.2e-6)fdr(0.74) *11 *12 52
-log10(corrP_Fdr)
0 2 4 6 8 10
-log10
-52 -51 -50 -49 -48 -47 -46 -45 -44 -43 -42 -41 -40 -39 -38 -37 -36 -35 -34 -33 -32 -31 -30 -29 -28 -27 -26 -25 -24 -23 -22 -21 -20 -19 -18 -17 -16 -15 -14 -13 -12 -11 -10 -9 -8 -7 -6 -5 -4 -3 -2 -1 0 1 2