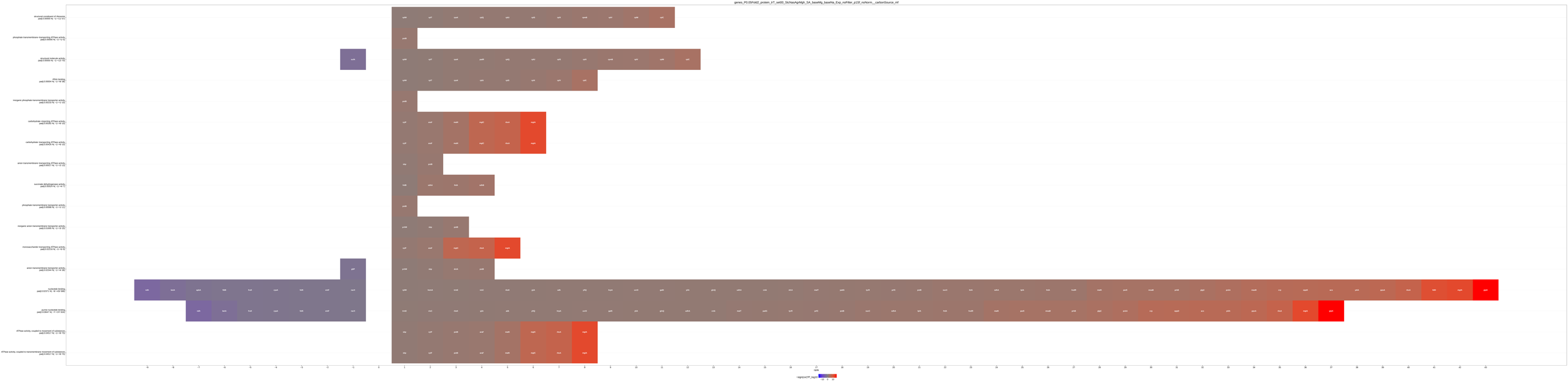
genes_P0.05Fdr82_protein_b'f_set00_StcNasAgrMgh_SA_baseMdg_baseNa_Exp_noFilter_p15f_noNorm__carbonSource_mf
structural constituent of ribosome
padj:0.00000 NL =2 =Ln/ 571
phosphate transmembrane-transporting ATPase activity
padj:0.00000 NL =Ln =2 8
structural molecule activity
padj:0.00000 NL =2 =Ln/ 731
rRNA binding
padj:0.00004 NL =Ln =Ir/ 38
inorganic phosphate transmembrane transporter activity
padj:0.00033 NL =Ln =Ir/ 103
carbohydrate-importing ATPase activity
padj:0.00082 NL =Ir/ =Ir/ 131
carbohydrate-transporting ATPase activity
padj:0.00424 NL =Ir/ =Ir/ 215
anion transmembrane-transporting ATPase activity
padj:0.00812 NL =Ir/ =Ir/ 315
fumarate dehydratase activity
padj:0.00535 NL =Ir/ =Ir/ 7
phosphate transmembrane transporter activity
padj:0.00595 NL =Ir/ =Ir/ 112
inorganic anion transmembrane transporter activity
padj:0.00895 NL =Ir/ =Ir/ 152
monosaccharide-transporting ATPase activity
padj:0.01216 NL =Ln =Ir/ 6
anion transmembrane transporter activity
padj:0.01944 NL =Ir/ =Ir/ 96
nucleoside binding
padj:0.02371 NL =Ir/ =Ir/2 5965
purine nucleoside binding
padj:0.03647 NL =Ir/ =Ir/2 5936
ATPase activity, coupled to movement of substances
padj:0.04217 NL =Ir/ =Ir/ 70
ATPase activity, coupled to transmembrane movement of substances
padj:0.04812 NL =Ir/ =Ir/ 70
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