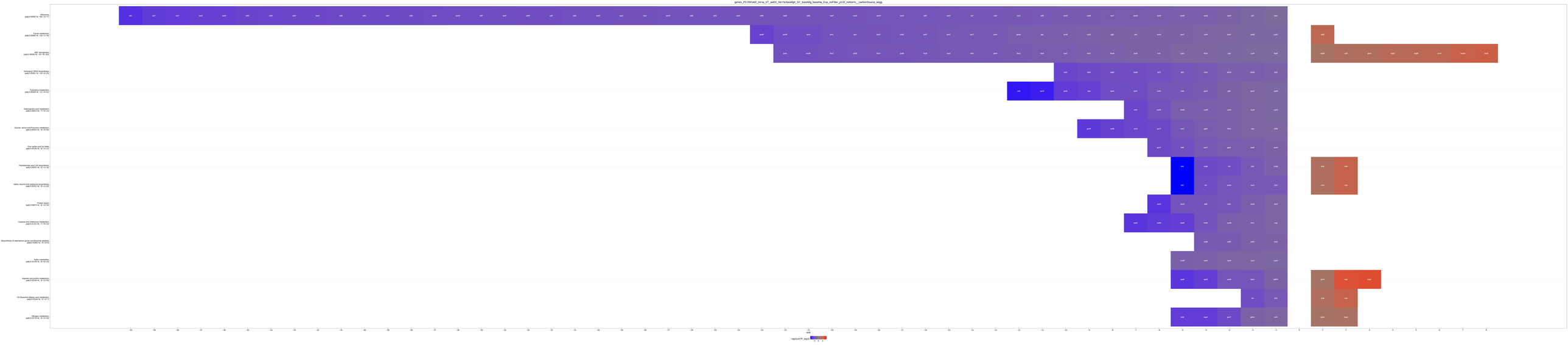
genes_PD.DSFdiff2_mrna_trT_setD2_filtYtNotdiffgh_SV_baseMq_baseNq_Exp_noFilter_jc1SI_noNcorrs__carbonSource_kegg
Ribosome
padj:0.00000 NL -Inf +Inf TL
Purine metabolism
padj:0.00000 NL -119 +3 TL
ABC transporters
padj:0.00000 NL -124 +8 TL
Aminoacyl-tRNA biosynthesis
padj:0.00000 NL -122 +9 TL
Pyrimidine metabolism
padj:0.00000 NL -123 +9 TL
Selenocompound metabolism
padj:0.00001NL -77 +9 TL
Glycine, serine and threonine metabolism
padj:0.00001 NL -58 +9 TL
Fine carbon sink fix fields
padj:0.00120 NL -54 +9 TL
Pantothenate and CoA biosynthesis
padj:0.00001 NL -51 +9 TL
Valine, leucine and isoleucine biosynthesis
padj:0.00001 NL -52 +9 TL
Protein export
padj:0.00079 NL -52 +9 TL
Cysteine and methionine metabolism
padj:0.01101 NL -11 +9 TL
Biosynthesis of siderophore group nonribosomal peptides
padj:0.01082 NL -11 +9 TL
Sulfur metabolism
padj:0.01104 NL -10 +9 TL
Arginine and proline metabolism
padj:0.01104 NL -10 +9 TL
C5-Branched dibasic acid metabolism
padj:0.01104 NL -12 +12 TL
Nitrogen metabolism
padj:0.01104 NL -10 +9 TL
-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 3 4 5 6 7 8
rank
-log10(corVP_log10)
-8 -6 -4 -2