genes_P0.05Fold2_protein_trT_set00_StcNasAgrMgh_SA_baseMg_baseNa_Exp_noFilter_p1Sf_noNorm__carbonSource_mf rplW rplT rpsK yadR rplQ rplU rplS rplX rpmB rplV rplM rplC rRNA binding padj:0.00004 N(-1/ +8/ 38) rplW rplT rpsK rplU rplS rplX rplV rplC araF malK mglC rbsA frdB sdhA frdA sdhB ychM sbp dctA pstB ndh betA apbA fldB fruK cyaA folK entF narX rpIW hemA glac trmE visC rbsK yjiA glgC rbsK yjiA glgC rbsA malK prkB moaB trmE visC rbsA glnQ udhA cmk sfcA malK prkB moaB trmE sucC folA moaB trmE sucC moaB wigC rbsA glnQ udhA cmk sfcA malK prkB wisC rbsA wigC rbs trmE visC rbsK yjiA acs yfiQ prkB yjiA sucC sdhA lpIA glpK yliA glpK yliA glpK yliA glpK yliA glpK yliA glpK yliA sucC sdhA bryR glpK yliA glpK yl ndh betA fruK cyaA folK entF narX sbp xylF pstB araF malK mglC rbsA mg sbp xyIF pstB araF malK mgIC rbsA mgIA