



- 1: 2.41 GO:0043228~non-membrane-bounded organelle
 2: 2.41 GO:0043232~intracellular non-membrane-bounded organelle
 3: 3.49 ribosomal protein
 4: 3.23 GO:0005840~ribosome
 5: 3.13 GO:0030529~ribonucleoprotein complex
 6: 3.29 ribosome
 7: 3.07 ecp03010:Ribosome
 8: 3.17 ribonucleoprotein
 9: 2.47 protein biosynthesis
 10: 2.87 GO:0003735~structural constituent of ribosome
 11: 2.46 GO:0005198~structural molecule activity
 12: 1.92 GO:0006412~translation
 13: 18.47 SM00411:BHL
 14: 17.65 IPR000119:Histone-like bacterial DNA-binding protein
 15: 17.48 PIRSF002095:DNA-binding protein HU,alpha/beta subunits
 16: 16.95 GO:0015768~maltose transport
 17: 12.71 GO:0000017~alpha-glucoside transport
 18: 12.71 GO:0015766~disaccharide transport
 19: 12.71 GO:0042946~glucoside transport
 20: 12.71 GO:0015772~oligosaccharide transport
 21: 2.82 GO:0046395~carboxylic acid catabolic process
 22: 2.82 GO:0016054~organic acid catabolic process
 23: 3.92 GO:0006970~response to osmotic stress
 24: 2.02 GO:0009628~response to abiotic stimulus
 25: 10.82 ecj00430:Taurine and hypotaurine metabolism
 26: 6.57 translation regulation
 27: 3.99 GO:0006417~regulation of translation
 28: 3.99 GO:0032268~regulation of cellular protein metabolic process
 29: 3.57 GO:0010608~posttranscriptional regulation of gene expression
 30: 2.93 rRNA-binding
 31: 2.65 GO:0019843~rRNA binding
 32: 2.01 sugar transport
 33: 1.66 GO:0008643~carbohydrate transport
 34: 2.72 GO:0016407~acetyltransferase activity
 35: 1.98 Acyltransferase
 36: 3.27 GO:0016410~N-acyltransferase activity
 37: 4.15 IPR016181:Acy-CoA N-acyltransferase
 38: 5.83 Translation,ribosomal structure and biogenesis
 39: 3.62 domain:N-acetyltransferase
 40: 5.53 IPR000182:GCN5-related N-acetyltransferase
 41: 2.97 GO:0008080~N-acetyltransferase activity