	S1	S28	S6
p__Human Diseases_PWY55; Staphylococcus aureus infection	0.7	3.6	2.3
p__Human Diseases_PWY49; Bacterial invasion of epithelial cells	0.2	1.2	3.2
p__Genetic Information Processing_PWY37; Ribosome	1	1.6	2
p__Organismal Systems_PWY142; Mineral absorption	1	1.6	1.7
p__Human Diseases_PWY53; Pertussis	1.8	0.7	1.6
p__Cellular Processes_PWY1; Apoptosis	0.8	2.5	0.4
p__Genetic Information Processing_PWY33; RNA polymerase	0.7	1.5	1.3
p__Genetic Information Processing_PWY38; Ribosome biogenesis in eukaryotes	1.1	0.8	1.4
p__Genetic Information Processing_PWY23; Protein export	0.8	1.1	1.4
p__Environmental Information Processing_PWY18; Bacterial secretion system	1.3	0.9	1.1
p__Human Diseases_PWY45; Cationic antimicrobial peptide (CAMP) resistance	0.6	1.6	1
p__Metabolism_PWY114; Biotin metabolism	0.9	1.1	1.3
p__Human Diseases_PWY44; beta−Lactam resistance	0.7	1.6	0.9
p__Human Diseases_PWY40; Chemical carcinogenesis − reactive oxygen species	0.9	1	1.2
p__Metabolism_PWY107; Teichoic acid biosynthesis	0.6	1.5	1
p__Cellular Processes_PWY12; Quorum sensing	0.8	1.2	1.1
p__Metabolism_PWY80; Galactose metabolism	0.5	1.4	1.2
p__Organismal Systems_PWY140; Longevity regulating pathway − multiple species	1.1	1.1	0.9
p__Cellular Processes_PWY13; Autophagy − yeast	1	0.8	1.2
p__Metabolism_PWY115; Pantothenate and CoA biosynthesis	0.7	1	1.3
p__Metabolism_PWY130; Pyrimidine metabolism	0.6	1	1.3
p__Metabolism_PWY112; Glycerophospholipid metabolism	0.6	1.1	1.2
p__Cellular Processes_PWY3; Cell cycle − Caulobacter	0.8	1	1.2
p__Environmental Information Processing_PWY17; ABC transporters	0.7	1	1.2
p__Metabolism_PWY99; Nucleotide metabolism	0.7	1	1.1
p__Cellular Processes_PWY11; Biofilm formation − Vibrio cholerae	0.8	1	1.1
p__Metabolism_PWY72; Valine, leucine and isoleucine biosynthesis	0.7	1	1.2
p__Genetic Information Processing_PWY35; Aminoacyl−tRNA biosynthesis	0.8	0.9	1.2
p__Metabolism_PWY118; Riboflavin metabolism	0.8	1	1.1
p__Cellular Processes_PWY7; Flagellar assembly	1	0.7	1.2
p__Cellular Processes_PWY9; Biofilm formation − Escherichia coli	0.8	0.8	1.2
p__Genetic Information Processing_PWY25; RNA degradation	0.8	1	1.1
p__Metabolism_PWY121; Phosphonate and phosphinate metabolism	0.9	1.1	0.8
p__Environmental Information Processing_PWY21; Two−component system	0.7	1.1	0.9
p__Human Diseases_PWY50; Epithelial cell signaling in Helicobacter pylori infection	1.2	1	0.6

S1 S28 S6

% Read Abundance

3.0

1.0

0.3

0.1