



- 1: 2.13 GO:0042597~periplasmic space
- 2: 1.71 periplasm
- 3: 6.21 IPR003173:response regulator,OmpR type
- 4: 4.56 IPR001867:Signal transduction response regulator,C-terminal
- 5: 2.2 GO:0009165~nucleotide biosynthetic process
- 6: 1.98 GO:0034654~nucleobase,nucleoside and nucleic acid biosynthetic process
- 7: 1.98 GO:0034404~nucleobase,nucleoside and nucleotide biosynthetic process
- 8: 5.65 ecz00670:One carbon pool by folate
- 9: 3.3 domain:Response regulatory
- 10: 3.22 IPR001789:Signal transduction response regulator,receiver region
- 11: 2.82 SM00448:REC
- 12: 2.34 two-component regulatory system
- 13: 2.47 GO:0000156~two-component response regulator activity
- 14: 5.5 purine biosynthesis
- 15: 2.46 GO:0006164~purine nucleotide biosynthetic process
- 16: 2.4 GO:0006163~purine nucleotide metabolic process
- 17: 3.39 ecs00230:Purine metabolism
- 18: 2.28 ece00230:Purine metabolism
- 19: 1.62 transcription regulation
- 20: 1.49 Transcription
- 21: 1.41 GO:0006350~transcription
- 22: 1.33 GO:0045449~regulation of transcription
- 23: 1.3 GO:0030528~transcription regulator activity
- 24: 4.4 potassium
- 25: 3.14 GO:0031420~alkali metal ion binding
- 26: 3.99 GO:0030955~potassium ion binding
- 27: 7.26 ecf00380:Tryptophan metabolism
- 28: 6.24 purine nucleotide biosynthesis
- 29: 5.03 GO:0009161~ribonucleoside monophosphate metabolic process
- 30: 5.03 GO:0009156~ribonucleoside monophosphate biosynthetic process
- 31: 4.14 GO:0009123~nucleoside monophosphate metabolic process
- 32: 4.14 GO:0009124~nucleoside monophosphate biosynthetic process
- 33: 4.8 GO:0009126~purine nucleoside monophosphate metabolic process
- 34: 4.8 GO:0009167~purine ribonucleoside monophosphate metabolic process
- 35: 4.8 GO:0009168~purine ribonucleoside monophosphate biosynthetic process
- 36: 4.8 GO:0009127~purine nucleoside monophosphate biosynthetic process
- 37: 2.45 GO:0009260~ribonucleotide biosynthetic process
- 38: 2.38 GO:0009259~ribonucleotide metabolic process
- 39: 2.13 GO:0009152~purine ribonucleotide biosynthetic process
- 40: 2.07 GO:0009150~purine ribonucleotide metabolic process

- group 1: 1.92
- group 2: 1.46
- group 3: 1.38
- group 4: 1.36
- group 5: 1.36
- group 6: 1.34
- group 7: 1.28
- group 8: 1.25
- group 9: 1.18
- group 10: 1.18
- group 11: 1.03
- group 12: 1.01