



group 1: 2.45
group 2: 1.68
group 3: 1.11
group 4: 1.38
group 5: 1.31
group 6: 1.25
group 7: 1.18
group 8: 1.22
group 9: 1.19
group 10: 1.18
group 11: 1.13
group 12: 1.12

1-4.23 GO:00090931-branched chain family amino acid biosynthetic process
2-4.15 GO:00090931-branched chain family amino acid metabolic process
3-4.15 GO:00090931-branched chain family amino acid biosynthetic process
4-1.68 GO:0030313-cell envelope
5-1.68 GO:0030313-cell envelope
6-1.68 GO:00090931-cell envelope
7-1.68 GO:00090931-cell envelope
8-1.68 GO:00090931-cell envelope
9-1.37 GO:00090931-cell envelope
10-1.36 GO:00424802-identical protein binding
11-1.36 GO:00424802-identical protein binding
12-1.28 GO:0030312-external encapsulating structure
13-1.28 GO:0030312-external encapsulating structure
14-1.18 GO:000974-peptidoglycan-based cell wall
15-1.18 GO:000974-peptidoglycan-based cell wall
16-1.05 grp-binding
17-1.05 grp-binding
18-1.75 GO:0019901-guanosine nucleotide binding
19-1.75 GO:0019901-guanosine nucleotide binding
20-1.99 GO:0019867-outer membrane
21-1.99 GO:0019867-outer membrane
22-4.3 GO:0006551-leucine metabolic process
23-4.3 GO:0006551-leucine metabolic process
24-4.3 GO:0045934-negative regulation of nucleobase/nucleoside/nucleotide and nucleic acid metabolic process
25-4.3 GO:0045934-negative regulation of nucleobase/nucleoside/nucleotide and nucleic acid metabolic process
26-4.3 GO:0010558-negative regulation of macromolecule biosynthetic process
27-4.3 GO:0010558-negative regulation of macromolecule biosynthetic process
28-4.05 GO:0009890-negative regulation of biosynthetic process
29-3.19 GO:0046477-colanic acid biosynthetic process
30-3.19 GO:0046477-colanic acid metabolic process
31-3.19 GO:0046477-colanic acid biosynthetic process
32-1.33 signal transduction
33-1.33 signal transduction
34-1.41 dna-binding
35-1.33 dna-binding