| Term | Annotated | Significant | Expected | Rank in classic Fisher | classic Fisher | elimFisher |
|---|---|---|---|---|---|---|
| lipid transport | 6 | 6 | 2.3 | 1 | 0.0025 | 0.0025 |
| proteolysis | 15 | 11 | 5.74 | 2 | 0.0037 | 0.0037 |
| embryonic development ending in birth or... | 17 | 12 | 6.5 | 3 | 0.0038 | 0.0038 |
| embryonic development | 18 | 12 | 6.89 | 4 | 0.008 | 1 |
| developmental process | 37 | 20 | 14.16 | 5 | 0.0145 | 0.2978 |
| multicellular organismal development | 35 | 18 | 13.39 | 6 | 0.044 | 0.575 |
| protein metabolic process | 19 | 11 | 7.27 | 7 | 0.049 | 1 |
| cellular protein metabolic process | 19 | 11 | 7.27 | 8 | 0.049 | 1 |
| cellular biopolymer metabolic process | 22 | 12 | 8.42 | 9 | 0.0676 | 0.9457 |
| cellular macromolecule metabolic process | 22 | 12 | 8.42 | 10 | 0.0676 | 0.9457 |
| multicellular organismal process | 39 | 19 | 14.92 | 11 | 0.074 | 0.6319 |
| macromolecule metabolic process | 23 | 12 | 8.8 | 12 | 0.0986 | 0.965 |
| biopolymer metabolic process | 23 | 12 | 8.8 | 13 | 0.0986 | 0.965 |
| regulation of growth rate | 19 | 10 | 7.27 | 14 | 0.1252 | 0.1252 |
| positive regulation of growth rate | 19 | 10 | 7.27 | 15 | 0.1252 | 0.1252 |
| biological regulation | 24 | 12 | 9.18 | 16 | 0.1373 | 0.1373 |
| regulation of growth | 20 | 10 | 7.65 | 17 | 0.1744 | 0.1744 |
| positive regulation of growth | 20 | 10 | 7.65 | 18 | 0.1744 | 0.1744 |
| positive regulation of biological proces... | 20 | 10 | 7.65 | 19 | 0.1744 | 0.1744 |
| growth | 23 | 11 | 8.8 | 20 | 0.2065 | 0.2065 |
| regulation of biological process | 23 | 11 | 8.8 | 21 | 0.2065 | 0.2065 |
| anatomical structure development | 7 | 4 | 2.68 | 22 | 0.251 | 0.251 |
| anatomical structure morphogenesis | 5 | 3 | 1.91 | 23 | 0.2843 | 0.2843 |
| transport | 17 | 8 | 6.5 | 24 | 0.292 | 0.9476 |
| localization | 17 | 8 | 6.5 | 25 | 0.292 | 0.9476 |
| establishment of localization | 17 | 8 | 6.5 | 26 | 0.292 | 0.9476 |
| body morphogenesis | 3 | 2 | 1.15 | 27 | 0.3257 | 0.3257 |
| transition metal ion transport | 1 | 1 | 0.38 | 28 | 0.3826 | 0.3826 |
| cell fate specification | 1 | 1 | 0.38 | 29 | 0.3826 | 0.3826 |
| morphogenesis of an epithelium | 1 | 1 | 0.38 | 30 | 0.3826 | 0.3826 |
| iron ion transport | 1 | 1 | 0.38 | 31 | 0.3826 | 0.3826 |
| cellular ion homeostasis | 1 | 1 | 0.38 | 32 | 0.3826 | 0.3826 |
| cellular iron ion homeostasis | 1 | 1 | 0.38 | 33 | 0.3826 | 0.3826 |
| pattern specification process | 1 | 1 | 0.38 | 34 | 0.3826 | 0.3826 |
| protein-based cuticle development | 1 | 1 | 0.38 | 35 | 0.3826 | 0.3826 |
| embryonic pattern specification | 1 | 1 | 0.38 | 36 | 0.3826 | 0.3826 |
| di-, tri-valent inorganic cation transpo... | 1 | 1 | 0.38 | 37 | 0.3826 | 0.3826 |
| cellular homeostasis | 1 | 1 | 0.38 | 38 | 0.3826 | 0.3826 |
| cellular cation homeostasis | 1 | 1 | 0.38 | 39 | 0.3826 | 0.3826 |
| cellular di-, tri-valent inorganic catio... | 1 | 1 | 0.38 | 40 | 0.3826 | 0.3826 |
| cell differentiation | 1 | 1 | 0.38 | 41 | 0.3826 | 0.3826 |
| collagen and cuticulin-based cuticle dev... | 1 | 1 | 0.38 | 42 | 0.3826 | 0.3826 |
| regulation of locomotion | 1 | 1 | 0.38 | 43 | 0.3826 | 0.3826 |
| positive regulation of locomotion | 1 | 1 | 0.38 | 44 | 0.3826 | 0.3826 |
| cuticle development | 1 | 1 | 0.38 | 45 | 0.3826 | 0.3826 |
| homeostatic process | 1 | 1 | 0.38 | 46 | 0.3826 | 0.3826 |
| cell fate commitment | 1 | 1 | 0.38 | 47 | 0.3826 | 0.3826 |
| cellular developmental process | 1 | 1 | 0.38 | 48 | 0.3826 | 0.3826 |
| chemical homeostasis | 1 | 1 | 0.38 | 49 | 0.3826 | 0.3826 |
| ion homeostasis | 1 | 1 | 0.38 | 50 | 0.3826 | 0.3826 |
| di-, tri-valent inorganic cation homeost... | 1 | 1 | 0.38 | 51 | 0.3826 | 0.3826 |
| iron ion homeostasis | 1 | 1 | 0.38 | 52 | 0.3826 | 0.3826 |
| cation homeostasis | 1 | 1 | 0.38 | 53 | 0.3826 | 0.3826 |
| cellular chemical homeostasis | 1 | 1 | 0.38 | 54 | 0.3826 | 0.3826 |
| transmembrane transport | 1 | 1 | 0.38 | 55 | 0.3826 | 0.3826 |
| regulation of biological quality | 1 | 1 | 0.38 | 56 | 0.3826 | 0.3826 |
| aging | 11 | 5 | 4.21 | 57 | 0.4171 | 0.4171 |
| determination of adult life span | 11 | 5 | 4.21 | 58 | 0.4171 | 0.4171 |
| multicellular organismal aging | 11 | 5 | 4.21 | 59 | 0.4171 | 0.4171 |
| locomotion | 11 | 5 | 4.21 | 60 | 0.4171 | 0.4171 |
| reproductive developmental process | 2 | 1 | 0.77 | 61 | 0.6209 | 0.6209 |
| cation transport | 2 | 1 | 0.77 | 62 | 0.6209 | 0.6209 |
| sex differentiation | 2 | 1 | 0.77 | 63 | 0.6209 | 0.6209 |
| metal ion transport | 2 | 1 | 0.77 | 64 | 0.6209 | 0.6209 |
| hermaphrodite genitalia development | 2 | 1 | 0.77 | 65 | 0.6209 | 0.6209 |
| organ development | 2 | 1 | 0.77 | 66 | 0.6209 | 0.6209 |
| system development | 2 | 1 | 0.77 | 67 | 0.6209 | 0.6209 |
| genitalia development | 2 | 1 | 0.77 | 68 | 0.6209 | 0.6209 |
| nematode larval development | 5 | 2 | 1.91 | 69 | 0.6366 | 0.6366 |
| larval development | 5 | 2 | 1.91 | 70 | 0.6366 | 0.6366 |
| post-embryonic development | 5 | 2 | 1.91 | 71 | 0.6366 | 0.6366 |
| behavior | 6 | 2 | 2.3 | 72 | 0.7455 | 0.7455 |

| Term | Annotated | Significant | Expected | Rank in classic Fisher | classic Fisher | elimFisher |
|---|---|---|---|---|---|---|
| oviposition | 6 | 2 | 2.3 | 73 | 0.7455 | 0.7455 |
| reproductive behavior | 6 | 2 | 2.3 | 74 | 0.7455 | 0.7455 |
| multicellular organism reproduction | 6 | 2 | 2.3 | 75 | 0.7455 | 0.7455 |
| reproductive behavior in a multicellular... | 6 | 2 | 2.3 | 76 | 0.7455 | 0.7455 |
| reproductive process in a multicellular ... | 6 | 2 | 2.3 | 77 | 0.7455 | 0.7455 |
| transcription | 3 | 1 | 1.15 | 78 | 0.7685 | 0.7685 |
| transcription, DNA-dependent | 3 | 1 | 1.15 | 79 | 0.7685 | 0.7685 |
| regulation of transcription, DNA-depende... | 3 | 1 | 1.15 | 80 | 0.7685 | 0.7685 |
| ion transport | 3 | 1 | 1.15 | 81 | 0.7685 | 0.7685 |
| regulation of biosynthetic process | 3 | 1 | 1.15 | 82 | 0.7685 | 0.7685 |
| regulation of gene expression | 3 | 1 | 1.15 | 83 | 0.7685 | 0.7685 |
| regulation of macromolecule biosynthetic... | 3 | 1 | 1.15 | 84 | 0.7685 | 0.7685 |
| RNA metabolic process | 3 | 1 | 1.15 | 85 | 0.7685 | 0.7685 |
| regulation of nucleobase, nucleoside, nu... | 3 | 1 | 1.15 | 86 | 0.7685 | 0.7685 |
| regulation of metabolic process | 3 | 1 | 1.15 | 87 | 0.7685 | 0.7685 |
| regulation of cellular metabolic process | 3 | 1 | 1.15 | 88 | 0.7685 | 0.7685 |
| regulation of cellular biosynthetic proc... | 3 | 1 | 1.15 | 89 | 0.7685 | 0.7685 |
| RNA biosynthetic process | 3 | 1 | 1.15 | 90 | 0.7685 | 0.7685 |
| regulation of transcription | 3 | 1 | 1.15 | 91 | 0.7685 | 0.7685 |
| regulation of RNA metabolic process | 3 | 1 | 1.15 | 92 | 0.7685 | 0.7685 |
| regulation of macromolecule metabolic pr... | 3 | 1 | 1.15 | 93 | 0.7685 | 0.7685 |
| metabolic process | 56 | 20 | 21.43 | 94 | 0.77 | 0.9863 |
| cellular metabolic process | 35 | 12 | 13.39 | 95 | 0.7839 | 0.9999 |
| reproduction | 13 | 4 | 4.97 | 96 | 0.8125 | 0.8125 |
| reproductive process | 7 | 2 | 2.68 | 97 | 0.8258 | 0.8258 |
| gene expression | 4 | 1 | 1.53 | 98 | 0.8595 | 0.8595 |
| cellular biopolymer biosynthetic process | 4 | 1 | 1.53 | 99 | 0.8595 | 0.8595 |
| biopolymer biosynthetic process | 4 | 1 | 1.53 | 100 | 0.8595 | 0.8595 |
| cellular process | 46 | 15 | 17.6 | 101 | 0.888 | 0.9994 |
| macromolecule biosynthetic process | 5 | 1 | 1.91 | 102 | 0.9152 | 0.9152 |
| cellular macromolecule biosynthetic proc... | 5 | 1 | 1.91 | 103 | 0.9152 | 0.9152 |
| regulation of cellular process | 5 | 1 | 1.91 | 104 | 0.9152 | 0.9152 |
| response to stimulus | 10 | 2 | 3.83 | 105 | 0.9498 | 0.9498 |
| primary metabolic process | 41 | 12 | 15.69 | 106 | 0.9544 | 1 |
| nucleobase, nucleoside, nucleotide and n... | 7 | 1 | 2.68 | 107 | 0.9696 | 0.9696 |
| biosynthetic process | 11 | 1 | 4.21 | 108 | 0.9964 | 0.9964 |
| cellular biosynthetic process | 11 | 1 | 4.21 | 109 | 0.9964 | 0.9964 |
| biological_process | 115 | 44 | 44 | 110 | 1 | 1 |
| peptidoglycan metabolic process | 2 | 0 | 0.77 | 111 | 1 | 1 |
| M phase | 1 | 0 | 0.38 | 112 | 1 | 1 |
| gastrulation with mouth forming first | 1 | 0 | 0.38 | 113 | 1 | 1 |
| carbohydrate metabolic process | 6 | 0 | 2.3 | 114 | 1 | 1 |
| polysaccharide metabolic process | 1 | 0 | 0.38 | 115 | 1 | 1 |
| monosaccharide metabolic process | 1 | 0 | 0.38 | 116 | 1 | 1 |
| fucose metabolic process | 1 | 0 | 0.38 | 117 | 1 | 1 |
| L-fucose biosynthetic process | 1 | 0 | 0.38 | 118 | 1 | 1 |
| mannose metabolic process | 1 | 0 | 0.38 | 119 | 1 | 1 |
| chitin metabolic process | 1 | 0 | 0.38 | 120 | 1 | 1 |
| chitin catabolic process | 1 | 0 | 0.38 | 121 | 1 | 1 |
| amino sugar metabolic process | 1 | 0 | 0.38 | 122 | 1 | 1 |
| glucosamine metabolic process | 1 | 0 | 0.38 | 123 | 1 | 1 |
| glucosamine catabolic process | 1 | 0 | 0.38 | 124 | 1 | 1 |
| N-acetylglucosamine metabolic process | 1 | 0 | 0.38 | 125 | 1 | 1 |
| N-acetylglucosamine catabolic process | 1 | 0 | 0.38 | 126 | 1 | 1 |
| cellular alcohol metabolic process | 1 | 0 | 0.38 | 127 | 1 | 1 |
| organic acid metabolic process | 6 | 0 | 2.3 | 128 | 1 | 1 |
| purine base metabolic process | 1 | 0 | 0.38 | 129 | 1 | 1 |
| purine nucleotide metabolic process | 3 | 0 | 1.15 | 130 | 1 | 1 |
| purine nucleotide biosynthetic process | 3 | 0 | 1.15 | 131 | 1 | 1 |
| IMP biosynthetic process | 3 | 0 | 1.15 | 132 | 1 | 1 |
| 'de novo' IMP biosynthetic process | 2 | 0 | 0.77 | 133 | 1 | 1 |
| DNA packaging | 2 | 0 | 0.77 | 134 | 1 | 1 |
| establishment or maintenance of chromati... | 2 | 0 | 0.77 | 135 | 1 | 1 |
| chromatin assembly or disassembly | 2 | 0 | 0.77 | 136 | 1 | 1 |
| nucleosome assembly | 2 | 0 | 0.77 | 137 | 1 | 1 |
| translation | 1 | 0 | 0.38 | 138 | 1 | 1 |
| protein folding | 1 | 0 | 0.38 | 139 | 1 | 1 |
| protein modification process | 3 | 0 | 1.15 | 140 | 1 | 1 |
| protein amino acid phosphorylation | 1 | 0 | 0.38 | 141 | 1 | 1 |
| protein amino acid glycosylation | 1 | 0 | 0.38 | 142 | 1 | 1 |
| cellular amino acid and derivative metab... | 2 | 0 | 0.77 | 143 | 1 | 1 |
| amino acid metabolic process | 2 | 0 | 0.77 | 144 | 1 | 1 |
| asparagine metabolic process | 1 | 0 | 0.38 | 145 | 1 | 1 |
| asparagine biosynthetic process | 1 | 0 | 0.38 | 146 | 1 | 1 |
| glutamine metabolic process | 1 | 0 | 0.38 | 147 | 1 | 1 |
| glutamine biosynthetic process | 1 | 0 | 0.38 | 148 | 1 | 1 |
| lipid metabolic process | 9 | 0 | 3.44 | 149 | 1 | 1 |

| Term | Annotated | Significant | Expected | Rank in classic Fisher | classic Fisher | elimFisher |
|---|---|---|---|---|---|---|
| fatty acid metabolic process | 3 | 0 | 1.15 | 150 | 1 | 1 |
| fatty acid biosynthetic process | 1 | 0 | 0.38 | 151 | 1 | 1 |
| acyl-CoA metabolic process | 1 | 0 | 0.38 | 152 | 1 | 1 |
| membrane lipid metabolic process | 2 | 0 | 0.77 | 153 | 1 | 1 |
| phospholipid metabolic process | 2 | 0 | 0.77 | 154 | 1 | 1 |
| sphingolipid metabolic process | 2 | 0 | 0.77 | 155 | 1 | 1 |
| sphingomyelin metabolic process | 2 | 0 | 0.77 | 156 | 1 | 1 |
| sphingomyelin catabolic process | 2 | 0 | 0.77 | 157 | 1 | 1 |
| cellular aromatic compound metabolic pro... | 2 | 0 | 0.77 | 158 | 1 | 1 |
| coenzyme metabolic process | 2 | 0 | 0.77 | 159 | 1 | 1 |
| group transfer coenzyme metabolic proces... | 1 | 0 | 0.38 | 160 | 1 | 1 |
| nucleoside phosphate metabolic process | 3 | 0 | 1.15 | 161 | 1 | 1 |
| folic acid and derivative metabolic proc... | 1 | 0 | 0.38 | 162 | 1 | 1 |
| phosphorus metabolic process | 2 | 0 | 0.77 | 163 | 1 | 1 |
| phosphate metabolic process | 2 | 0 | 0.77 | 164 | 1 | 1 |
| nitrogen compound metabolic process | 2 | 0 | 0.77 | 165 | 1 | 1 |
| sodium ion transport | 1 | 0 | 0.38 | 166 | 1 | 1 |
| phosphate transport | 1 | 0 | 0.38 | 167 | 1 | 1 |
| anion transport | 1 | 0 | 0.38 | 168 | 1 | 1 |
| response to stress | 4 | 0 | 1.53 | 169 | 1 | 1 |
| defense response | 2 | 0 | 0.77 | 170 | 1 | 1 |
| ER-nuclear signaling pathway | 2 | 0 | 0.77 | 171 | 1 | 1 |
| response to unfolded protein | 2 | 0 | 0.77 | 172 | 1 | 1 |
| organelle organization | 2 | 0 | 0.77 | 173 | 1 | 1 |
| cell wall organization | 2 | 0 | 0.77 | 174 | 1 | 1 |
| cell cycle | 1 | 0 | 0.38 | 175 | 1 | 1 |
| chromosome segregation | 1 | 0 | 0.38 | 176 | 1 | 1 |
| meiosis | 1 | 0 | 0.38 | 177 | 1 | 1 |
| cell communication | 3 | 0 | 1.15 | 178 | 1 | 1 |
| cell adhesion | 1 | 0 | 0.38 | 179 | 1 | 1 |
| cell-matrix adhesion | 1 | 0 | 0.38 | 180 | 1 | 1 |
| signal transduction | 2 | 0 | 0.77 | 181 | 1 | 1 |
| intracellular signaling cascade | 2 | 0 | 0.77 | 182 | 1 | 1 |
| gastrulation | 1 | 0 | 0.38 | 183 | 1 | 1 |
| lipid biosynthetic process | 1 | 0 | 0.38 | 184 | 1 | 1 |
| amino acid biosynthetic process | 2 | 0 | 0.77 | 185 | 1 | 1 |
| catabolic process | 5 | 0 | 1.91 | 186 | 1 | 1 |
| glutamine family amino acid metabolic pr... | 1 | 0 | 0.38 | 187 | 1 | 1 |
| aspartate family amino acid metabolic pr... | 1 | 0 | 0.38 | 188 | 1 | 1 |
| aspartate family amino acid biosynthetic... | 1 | 0 | 0.38 | 189 | 1 | 1 |
| glutamine family amino acid biosynthetic... | 1 | 0 | 0.38 | 190 | 1 | 1 |
| glycoprotein metabolic process | 1 | 0 | 0.38 | 191 | 1 | 1 |
| glycoprotein biosynthetic process | 1 | 0 | 0.38 | 192 | 1 | 1 |
| coenzyme biosynthetic process | 1 | 0 | 0.38 | 193 | 1 | 1 |
| nucleobase metabolic process | 1 | 0 | 0.38 | 194 | 1 | 1 |
| purine base biosynthetic process | 1 | 0 | 0.38 | 195 | 1 | 1 |
| nucleotide metabolic process | 3 | 0 | 1.15 | 196 | 1 | 1 |
| nucleoside monophosphate metabolic proce... | 3 | 0 | 1.15 | 197 | 1 | 1 |
| nucleoside monophosphate biosynthetic pr... | 3 | 0 | 1.15 | 198 | 1 | 1 |
| purine nucleoside monophosphate metaboli... | 3 | 0 | 1.15 | 199 | 1 | 1 |
| purine nucleoside monophosphate biosynth... | 3 | 0 | 1.15 | 200 | 1 | 1 |
| purine ribonucleotide metabolic process | 3 | 0 | 1.15 | 201 | 1 | 1 |
| purine ribonucleotide biosynthetic proce... | 3 | 0 | 1.15 | 202 | 1 | 1 |
| ribonucleoside monophosphate biosyntheti... | 3 | 0 | 1.15 | 203 | 1 | 1 |
| ribonucleoside monophosphate metabolic p... | 3 | 0 | 1.15 | 204 | 1 | 1 |
| nucleotide biosynthetic process | 3 | 0 | 1.15 | 205 | 1 | 1 |
| purine ribonucleoside monophosphate meta... | 3 | 0 | 1.15 | 206 | 1 | 1 |
| purine ribonucleoside monophosphate bios... | 3 | 0 | 1.15 | 207 | 1 | 1 |
| nucleotide-sugar metabolic process | 1 | 0 | 0.38 | 208 | 1 | 1 |
| nucleotide-sugar biosynthetic process | 1 | 0 | 0.38 | 209 | 1 | 1 |
| peptidoglycan catabolic process | 2 | 0 | 0.77 | 210 | 1 | 1 |
| ribonucleotide metabolic process | 3 | 0 | 1.15 | 211 | 1 | 1 |
| ribonucleotide biosynthetic process | 3 | 0 | 1.15 | 212 | 1 | 1 |
| cellular amine metabolic process | 2 | 0 | 0.77 | 213 | 1 | 1 |
| amine biosynthetic process | 2 | 0 | 0.77 | 214 | 1 | 1 |
| phospholipid catabolic process | 2 | 0 | 0.77 | 215 | 1 | 1 |
| folic acid and derivative biosynthetic p... | 1 | 0 | 0.38 | 216 | 1 | 1 |
| response to biotic stimulus | 2 | 0 | 0.77 | 217 | 1 | 1 |
| gas transport | 1 | 0 | 0.38 | 218 | 1 | 1 |
| oxygen transport | 1 | 0 | 0.38 | 219 | 1 | 1 |
| monovalent inorganic cation transport | 1 | 0 | 0.38 | 220 | 1 | 1 |
| inorganic anion transport | 1 | 0 | 0.38 | 221 | 1 | 1 |
| lipid catabolic process | 2 | 0 | 0.77 | 222 | 1 | 1 |
| cellular component organization | 4 | 0 | 1.53 | 223 | 1 | 1 |
| carbohydrate biosynthetic process | 1 | 0 | 0.38 | 224 | 1 | 1 |
| carbohydrate catabolic process | 3 | 0 | 1.15 | 225 | 1 | 1 |

| Term | Annotated | Significant | Expected | Rank in classic Fisher | classic Fisher | elimFisher |
|---|---|---|---|---|---|---|
| organic acid biosynthetic process | 1 | 0 | 0.38 | 226 | 1 | 1 |
| phosphorylation | 1 | 0 | 0.38 | 227 | 1 | 1 |
| protein processing | 1 | 0 | 0.38 | 228 | 1 | 1 |
| intein-mediated protein splicing | 1 | 0 | 0.38 | 229 | 1 | 1 |
| cell wall catabolic process | 2 | 0 | 0.77 | 230 | 1 | 1 |
| molting cycle, protein-based cuticle | 1 | 0 | 0.38 | 231 | 1 | 1 |
| molting cycle, collagen and cuticulin-ba... | 1 | 0 | 0.38 | 232 | 1 | 1 |
| hexose metabolic process | 1 | 0 | 0.38 | 233 | 1 | 1 |
| hexose biosynthetic process | 1 | 0 | 0.38 | 234 | 1 | 1 |
| GDP-mannose metabolic process | 1 | 0 | 0.38 | 235 | 1 | 1 |
| carboxylic acid metabolic process | 6 | 0 | 2.3 | 236 | 1 | 1 |
| cell cycle process | 1 | 0 | 0.38 | 237 | 1 | 1 |
| cell cycle phase | 1 | 0 | 0.38 | 238 | 1 | 1 |
| cellular component assembly | 2 | 0 | 0.77 | 239 | 1 | 1 |
| biological adhesion | 1 | 0 | 0.38 | 240 | 1 | 1 |
| sphingolipid catabolic process | 2 | 0 | 0.77 | 241 | 1 | 1 |
| protein splicing | 1 | 0 | 0.38 | 242 | 1 | 1 |
| endoplasmic reticulum unfolded protein r... | 2 | 0 | 0.77 | 243 | 1 | 1 |
| chromatin assembly | 2 | 0 | 0.77 | 244 | 1 | 1 |
| cell-substrate adhesion | 1 | 0 | 0.38 | 245 | 1 | 1 |
| monocarboxylic acid metabolic process | 4 | 0 | 1.53 | 246 | 1 | 1 |
| cellular response to stress | 2 | 0 | 0.77 | 247 | 1 | 1 |
| nucleobase, nucleoside and nucleotide bi... | 1 | 0 | 0.38 | 248 | 1 | 1 |
| cellular response to unfolded protein | 2 | 0 | 0.77 | 249 | 1 | 1 |
| cellular macromolecular complex subunit ... | 2 | 0 | 0.77 | 250 | 1 | 1 |
| cellular macromolecular complex assembly | 2 | 0 | 0.77 | 251 | 1 | 1 |
| cellular carbohydrate biosynthetic proce... | 1 | 0 | 0.38 | 252 | 1 | 1 |
| cellular nitrogen compound metabolic pro... | 2 | 0 | 0.77 | 253 | 1 | 1 |
| nucleobase, nucleoside, nucleotide and n... | 1 | 0 | 0.38 | 254 | 1 | 1 |
| nucleosome organization | 2 | 0 | 0.77 | 255 | 1 | 1 |
| response to endoplasmic reticulum stress | 2 | 0 | 0.77 | 256 | 1 | 1 |
| multicellular organism growth | 1 | 0 | 0.38 | 257 | 1 | 1 |
| regulation of multicellular organism gro... | 1 | 0 | 0.38 | 258 | 1 | 1 |
| positive regulation of multicellular org... | 1 | 0 | 0.38 | 259 | 1 | 1 |
| response to chemical stimulus | 2 | 0 | 0.77 | 260 | 1 | 1 |
| molting cycle | 1 | 0 | 0.38 | 261 | 1 | 1 |
| GDP-L-fucose biosynthetic process | 1 | 0 | 0.38 | 262 | 1 | 1 |
| fucose biosynthetic process | 1 | 0 | 0.38 | 263 | 1 | 1 |
| L-fucose metabolic process | 1 | 0 | 0.38 | 264 | 1 | 1 |
| pigment metabolic process | 1 | 0 | 0.38 | 265 | 1 | 1 |
| biopolymer modification | 3 | 0 | 1.15 | 266 | 1 | 1 |
| biopolymer glycosylation | 1 | 0 | 0.38 | 267 | 1 | 1 |
| post-translational protein modification | 2 | 0 | 0.77 | 268 | 1 | 1 |
| macromolecular complex subunit organizat... | 2 | 0 | 0.77 | 269 | 1 | 1 |
| cell wall metabolic process | 2 | 0 | 0.77 | 270 | 1 | 1 |
| cellular lipid catabolic process | 2 | 0 | 0.77 | 271 | 1 | 1 |
| cellular catabolic process | 2 | 0 | 0.77 | 272 | 1 | 1 |
| cellular lipid metabolic process | 5 | 0 | 1.91 | 273 | 1 | 1 |
| cellular carbohydrate metabolic process | 1 | 0 | 0.38 | 274 | 1 | 1 |
| nitrogen compound biosynthetic process | 2 | 0 | 0.77 | 275 | 1 | 1 |
| meiotic chromosome segregation | 1 | 0 | 0.38 | 276 | 1 | 1 |
| external encapsulating structure organiz... | 2 | 0 | 0.77 | 277 | 1 | 1 |
| IMP metabolic process | 3 | 0 | 1.15 | 278 | 1 | 1 |
| nucleobase biosynthetic process | 1 | 0 | 0.38 | 279 | 1 | 1 |
| pigment biosynthetic process | 1 | 0 | 0.38 | 280 | 1 | 1 |
| alcohol biosynthetic process | 1 | 0 | 0.38 | 281 | 1 | 1 |
| amino sugar catabolic process | 1 | 0 | 0.38 | 282 | 1 | 1 |
| monosaccharide biosynthetic process | 1 | 0 | 0.38 | 283 | 1 | 1 |
| GDP-L-fucose metabolic process | 1 | 0 | 0.38 | 284 | 1 | 1 |
| carboxylic acid biosynthetic process | 1 | 0 | 0.38 | 285 | 1 | 1 |
| membrane lipid catabolic process | 2 | 0 | 0.77 | 286 | 1 | 1 |
| heterocycle metabolic process | 2 | 0 | 0.77 | 287 | 1 | 1 |
| embryonic morphogenesis | 1 | 0 | 0.38 | 288 | 1 | 1 |
| cofactor metabolic process | 2 | 0 | 0.77 | 289 | 1 | 1 |
| cofactor biosynthetic process | 1 | 0 | 0.38 | 290 | 1 | 1 |
| regulation of multicellular organismal p... | 1 | 0 | 0.38 | 291 | 1 | 1 |
| positive regulation of multicellular org... | 1 | 0 | 0.38 | 292 | 1 | 1 |
| chromosome organization | 2 | 0 | 0.77 | 293 | 1 | 1 |
| meiotic cell cycle | 1 | 0 | 0.38 | 294 | 1 | 1 |
| M phase of meiotic cell cycle | 1 | 0 | 0.38 | 295 | 1 | 1 |
| cellular response to stimulus | 2 | 0 | 0.77 | 296 | 1 | 1 |
| response to protein stimulus | 2 | 0 | 0.77 | 297 | 1 | 1 |
| nucleobase, nucleoside and nucleotide me... | 3 | 0 | 1.15 | 298 | 1 | 1 |
| oxidation reduction | 1 | 0 | 0.38 | 299 | 1 | 1 |
| macromolecular complex assembly | 2 | 0 | 0.77 | 300 | 1 | 1 |
| protein-DNA complex assembly | 2 | 0 | 0.77 | 301 | 1 | 1 |
| glycosylation | 1 | 0 | 0.38 | 302 | 1 | 1 |