



- 19/377 nucleobase–containing compound catabolic process
- 21/577 organic cyclic compound catabolic process
- 18/208 RNA catabolic process
- 18/75 nuclear–transcribed mRNA catabolic process, nonsense–mediated decay
- 24/964 macromolecule catabolic process
- 23/533 mRNA metabolic process
- 41/424 amide biosynthetic process
- 20/111 translational initiation
- 44/658 cellular amide metabolic process
- 40/412 peptide metabolic process
- 38/316 peptide biosynthetic process
- 32/96 cytoplasmic translation
- 20/192 establishment of protein localization to membrane
- 25/378 protein localization to membrane
- 32/655 single–organism cellular localization
- 38/976 membrane organization
- 11/222 plasma membrane organization
- 20/89 protein localization to endoplasmic reticulum
- 25/642 protein localization to organelle
- 20/75 establishment of protein localization to endoplasmic reticulum
- 20/365 establishment of protein localization to organelle
- 20/107 protein targeting to membrane
- 20/278 protein targeting
- 27/655 intracellular protein transport
- 29/1060 establishment of protein localization
- 28/951 cytoplasmic transport
- 19/412 ribonucleoprotein complex biogenesis
- 5/32 ribosomal large subunit assembly
- 18/245 ribosome biogenesis
- 9/55 ribosome assembly
- 11/72 ribosomal large subunit biogenesis
- 9/211 rRNA metabolic process
- 4/27 maturation of LSU–rRNA
- 11/245 ribonucleoprotein complex subunit organization
- 4/18 ribosomal small subunit assembly
- 4/32 cellular response to cAMP
- 9/151 response to purine–containing compound
- 9/104 response to organophosphorus
- 7/127 regulation of tube architecture, open tracheal system
- 11/295 open tracheal system development
- 17/514 respiratory system development
- 13/378 wing disc development
- 20/738 imaginal disc development
- 4/26 head involution
- 3/17 focal adhesion assembly
- 5/52 cell–substrate junction assembly
- 13/261 cell junction organization
- 9/117 adherens junction organization
- 11/167 establishment of cell polarity
- 18/364 establishment or maintenance of cell polarity
- 3/9 establishment of neuroblast polarity
- 8/164 asymmetric cell division
- 12/322 protein complex localization
- 3/12 protein localization to cell cortex
- 21/593 actin filament–based process
- 11/230 actin filament organization
- 8/61 actin filament bundle organization
- 17/418 fibril organization
- 10/93 actin cytoskeleton reorganization
- 29/1028 cytoskeleton organization
- 19/525 microtubule cytoskeleton organization
- 22/707 microtubule–based process
- 7/118 cortical cytoskeleton organization
- 24/844 organelle assembly
- 5/67 sarcomere organization
- 17/509 positive regulation of cellular component biogenesis
- 28/1020 regulation of cellular component biogenesis
- 6/75 regulation of cell junction assembly
- 5/50 regulation of cell–substrate junction assembly
- 7/98 regulation of cell–matrix adhesion
- 24/898 regulation of cellular component movement
- 5/43 regulation of embryonic cell shape
- 12/159 regulation of cell shape
- 20/686 regulation of cell morphogenesis
- 19/402 regulation of actin filament–based process
- 8/172 regulation of protein polymerization
- 18/484 regulation of cytoskeleton organization
- 5/63 regulation of microtubule polymerization or depolymerization
- 8/138 negative regulation of cytoskeleton organization
- 5/57 regulation of protein depolymerization
- 7/80 regulation of protein complex disassembly
- 4/32 lamellipodium organization
- 5/64 Rho protein signal transduction
- 9/198 small GTPase mediated signal transduction
- 10/234 regulation of small GTPase mediated signal transduction
- 4/28 axon choice point recognition
- 17/537 neuron projection guidance
- 24/989 regulation of growth
- 15/481 regulation of cell growth
- 8/158 maintenance of location in cell
- 3/15 cytoskeletal anchoring at plasma membrane
- 8/161 stress–activated protein kinase signaling cascade
- 7/126 stress–activated MAPK cascade
- 3/18 renal filtration
- 3/18 inductive cell migration
- 3/14 energy coupled proton transport, down electrochemical gradient
- 6/68 nucleoside triphosphate biosynthetic process

p < 0.01
p < 0.05
p < 0.1