27/100 regulation of mitochondrion organization 220/996 regulation of phosphorylation 41/204 regulation of autophagy 23/55 positive regulation of peptide hormone secretion 163/958 regulation of cell development 36/257 regulation of synapse structure or 106/607 regulation of growth 143/519 regulation of immune response 18/38 cytosolic pattern recognition receptor signaling pathway 25/115 immune response-regulating cell surface receptor signaling pathway 85/274 activation of immune response 8/46 antigen receptor-mediated signaling pathway 30/98 regulation of I-kappaB kinase/NF-kappaB signaling 191/878 positive regulation of signal transduction 11/24 cholesterol efflux 85/441 immune system development 48/200 protein polyubiquitination 4/22 regulation of cytokinetic process 4/34 mitotic cytokinesis checkpoint signaling 13/54 cell cycle DNA replication 5/23 plant organ development 8/29 response to high light intensity 5/18 plant epidermis development 184/1120 neuron differentiation 21/102 response to osmotic stress 4/50 non-motile cilium assembly 46/391 cell projection assembly 4/53 intraciliary transport 54/343 microtubule-based movement 4/47 axonemal dynein complex assembly 23/186 cilium movement 7/95 cilium or flagellum-dependent cell motility 5/41 regulation of microtubule-based movement 111/517 obsolete cofactor metabolic process 43/214 pyridine-containing compound metabolic process 171/926 organophosphate metabolic process 4/23 sorocarp development 9/43 'de novo' protein folding 39/202 protein folding 21/88 chaperone-mediated protein folding 6/18 protein folding in endoplasmic reticulum 9/55 peptidyl-proline modification 95/275 RNA catabolic process 24/127 mitochondrial gene expression

p < 1e-05 p < 1e-04 p < 0.001