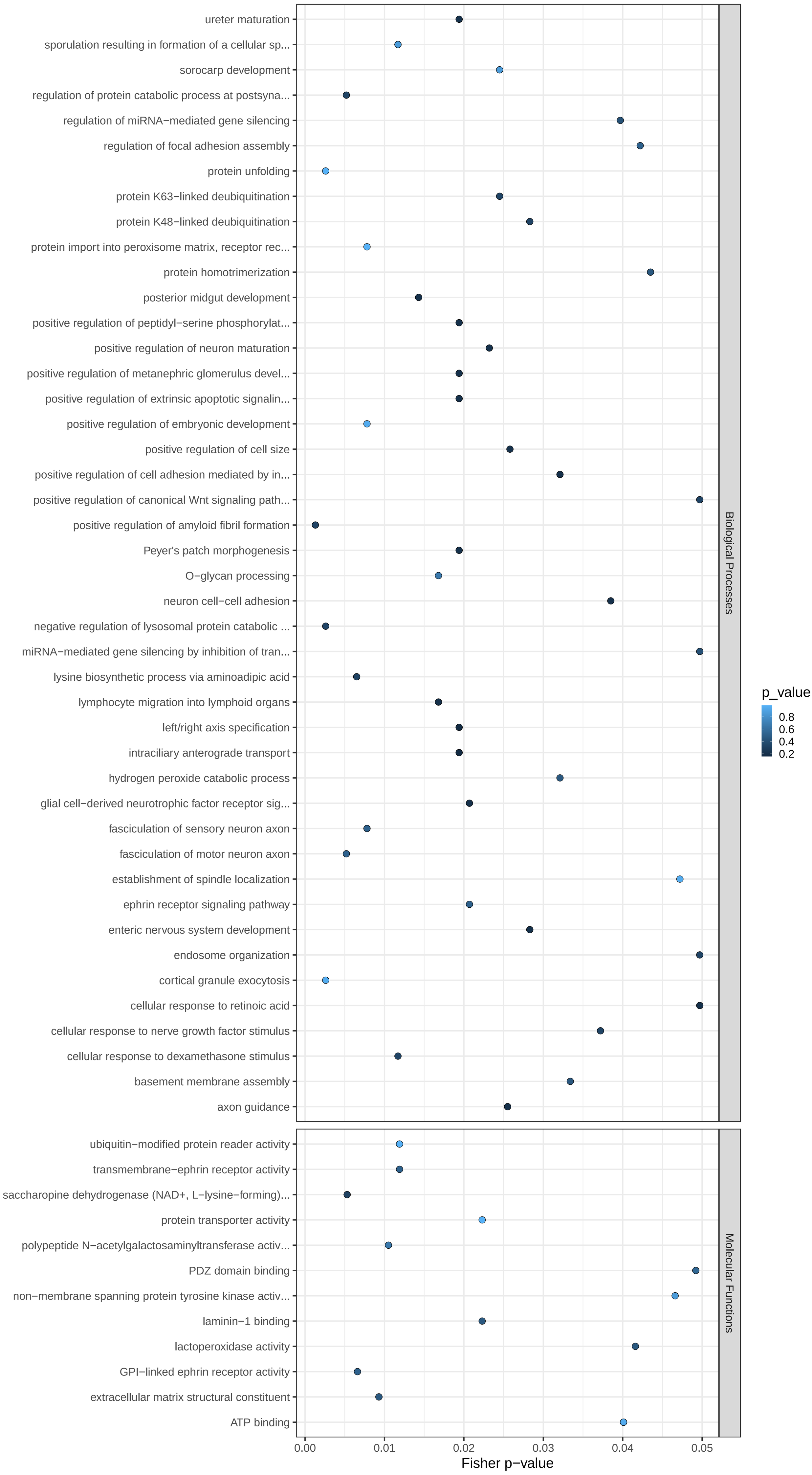
ureter maturation
sporulation resulting in formation of a cellular sp...
sorocarp development
regulation of protein catabolic process at postsyna...
regulation of miRNA−mediated gene silencing
regulation of focal adhesion assembly
protein unfolding
protein K63−linked deubiquitination
protein K48−linked deubiquitination
protein import into peroxisome matrix, receptor rec...
protein homotrimerization
posterior midgut development
positive regulation of peptidyl−serine phosphorylat...
positive regulation of neuron maturation
positive regulation of metanephric glomerulus devel...
positive regulation of extrinsic apoptotic signalin...
positive regulation of embryonic development
positive regulation of cell size
positive regulation of cell adhesion mediated by in...
positive regulation of canonical Wnt signaling path...
positive regulation of amyloid fibril formation
Peyer's patch morphogenesis
O−glycan processing
neuron cell−cell adhesion
negative regulation of lysosomal protein catabolic ...
miRNA−mediated gene silencing by inhibition of tran...
lysine biosynthetic process via aminoadipic acid
lymphocyte migration into lymphoid organs
left/right axis specification
intraciliary anterograde transport
hydrogen peroxide catabolic process
glial cell−derived neurotrophic factor receptor sig...
fasciculation of sensory neuron axon
fasciculation of motor neuron axon
establishment of spindle localization
ephrin receptor signaling pathway
enteric nervous system development
endosome organization
cortical granule exocytosis
cellular response to retinoic acid
cellular response to nerve growth factor stimulus
cellular response to dexamethasone stimulus
basement membrane assembly
axon guidance
Biological Processes
ubiquitin−modified protein reader activity
transmembrane−ephrin receptor activity
saccharopine dehydrogenase (NAD+, L−lysine−forming)...
protein transporter activity
polypeptide N−acetylgalactosaminyltransferase activ...
PDZ domain binding
non−membrane spanning protein tyrosine kinase activ...
laminin−1 binding
lactoperoxidase activity
GPI−linked ephrin receptor activity
extracellular matrix structural constituent
ATP binding
Molecular Functions
0.00
0.01
0.02
0.03
0.04
0.05
Fisher p−value
p_value
0.8
0.6
0.4
0.2