

regulation of macroautophagy, GO:0016241

- proton transport, GO:0015992
- GDP binding, GO:0019003
- fatty acid biosynthetic process, GO:0006633
- ATP hydrolysis coupled proton transport, GO:0015991
- cell growth, GO:0016049
- ADP binding, GO:0043531
- cholesterol biosynthetic process, GO:0006695
- proton-transporting ATPase activity, GO:0046961
- extracellular matrix disassembly, GO:0022617
- positive regulation of Wnt signaling pathway, GO:0030177
- positive regulation of endothelial cell proliferation, GO:0001938
- microvillus, GO:0005902
- sterol biosynthetic process, GO:0016126
- hippocampus development, GO:0021766
- misfolded protein binding, GO:0051787
- lipid biosynthetic process, GO:0008610
- axon extension, GO:0048675
- regulation of androgen receptor signaling pathway, GO:0060765
- ruffle organization, GO:0031529
- negative regulation of B cell proliferation, GO:0030889
- vacuolar acidification, GO:0007035
- vacuole, GO:0005773
- cysteine-type endopeptidase activator activity involved in apoptotic process, GO:0008656
- regulation of ryanodine-sensitive calcium-release channel activity, GO:0060314
- positive regulation of receptor activity, GO:2000273
- vacuolar membrane, GO:0005774
- cornification, GO:0070268
- visual learning, GO:0008542
- exocyst, GO:0000145
- cyclin-dependent protein serine/threonine kinase inhibitor activity, GO:0004861
- NAD-dependent histone deacetylase activity (H3-K14 specific), GO:0032041
- positive regulation of glycolytic process, GO:0045821
- calcium ion import, GO:0070509
- cellular zinc ion homeostasis, GO:0006882
- AMP binding, GO:0016208
- retromer, GO:0030906
- axonal fasciculation, GO:0007413
- protein serine/threonine kinase activator activity, GO:0043539
- high-density lipoprotein particle assembly, GO:0034380
- calcium-transporting ATPase activity, GO:0005388
- blood vessel morphogenesis, GO:0048514
- MAP kinase activity, GO:0004707
- vacuolar proton-transporting V-type ATPase complex assembly, GO:0070072
- proton-transporting V-type ATPase, GO:0033179
- neuron cell-cell adhesion, GO:0007158
- proton-transporting V-type ATPase, GO:0033180
- regulation of chaperone-mediated autophagy, GO:1904714
- proton-transporting two-sector ATPase complex, GO:0016469
- self proteolysis, GO:0097264