

centriole, GO:0005814

- cilium assembly, GO:0060271
- cell projection organization, GO:0030030
- ciliary basal body, GO:0036064
- intraciliary transport involved in cilium assembly, GO:0035735
- cell leading edge, GO:0031252
- axoneme, GO:0005930
- neural tube closure, GO:0001843
- ciliary tip, GO:0097542
- tubulin binding, GO:0015631
- spindle assembly, GO:0051225
- regulation of protein localization, GO:0032880
- pericentriolar material, GO:0000242
- protein export from nucleus, GO:0006611
- establishment of protein localization, GO:0045184
- motile cilium, GO:0031514
- cytoplasmic microtubule organization, GO:0031122
- smoothened signaling pathway, GO:0007224
- negative regulation of phosphatase activity, GO:0010923
- neuron migration, GO:0001764
- non-motile cilium assembly, GO:1905515
- kinesin binding, GO:0019894
- centriolar satellite, GO:0034451
- alpha-tubulin binding, GO:0043014
- microtubule-based process, GO:0007017
- ATP-dependent microtubule motor activity, GO:0008574
- ciliary base, GO:0097546
- determination of left/right symmetry, GO:0007368
- intraciliary transport particle B, GO:0030992
- intraciliary transport, GO:0042073
- gamma-tubulin binding, GO:0043015
- histone H4 deacetylation, GO:0070933
- ciliary transition zone, GO:0035869
- positive regulation of intracellular protein transport, GO:0090316
- centriole replication, GO:0007099
- aggresome, GO:0016235
- maintenance of protein location in nucleus, GO:0051457
- neural tube development, GO:0021915
- regulation of exocytosis, GO:0017157
- NAD+ binding, GO:0070403
- protein polyglutamylation, GO:0018095
- iron-sulfur cluster assembly, GO:0016226
- structural constituent of cytoskeleton, GO:0005200
- cardiac muscle cell differentiation, GO:0055007
- photoreceptor cell maintenance, GO:0045494
- BBSome, GO:0034464
- WASH complex, GO:0071203
- DNA binding, GO:0008301
- social behavior, GO:0035176
- myosin V binding, GO:0031489
- Flemming body, GO:0090543
- regulation of cilium assembly, GO:1902017
- positive regulation of interleukin-8 production, GO:0032757
- bleb, GO:0032059
- Rap protein signal transduction, GO:0032486
- regulation of smoothened signaling pathway, GO:0008589
- regulation of fat cell differentiation, GO:0045598
- hepatocyte growth factor receptor signaling pathway, GO:0048012
- neural tube formation, GO:0001841
- plus-end-directed vesicle transport along microtubule, GO:0072383
- XY body, GO:0001741
- Schmidt-Lanterman incisure, GO:0043220
- paranodal junction, GO:0033010
- epithelial structure maintenance, GO:0010669
- MKS complex, GO:0036038
- left/right axis specification, GO:0070986
- melanosome transport, GO:0032402
- positive regulation of focal adhesion assembly, GO:0051894
- protein ADP-ribosylation, GO:0006471
- COPI-coated vesicle, GO:0030137
- negative regulation of neurogenesis, GO:0050768
- motile cilium assembly, GO:0044458
- transport along microtubule, GO:0010970
- negative regulation of multicellular organism growth, GO:0040015
- face development, GO:0060324
- metallocarboxypeptidase activity, GO:0004181
- microtubule anchoring at centrosome, GO:0034454
- interphase microtubule nucleation by interphase microtubule organizing center, GO:0051415
- equatorial microtubule organizing center, GO:0000923
- negative regulation of dendritic spine development, GO:0061000
- dynein light chain binding, GO:0045503
- cochlea development, GO:0090102
- myoblast migration, GO:0051451
- negative regulation of protein catabolic process, GO:0042177
- microtubule minus-end binding, GO:0051011