

G2/M transition of mitotic cell cycle, GO:0000086

- cilium assembly, GO:0060271
- cell projection organization, GO:0030030
- protein dephosphorylation, GO:0006470
- ciliary basal body, GO:0036064
- peptidyl-threonine phosphorylation, GO:0018107
- negative regulation of protein binding, GO:0032091
- SCF ubiquitin ligase complex, GO:0019005
- animal organ regeneration, GO:0031100
- negative regulation of catalytic activity, GO:0043086
- intercellular bridge, GO:0045171
- positive regulation of intracellular protein transport, GO:0090316
- spindle assembly, GO:0051225
- positive regulation of cyclin-dependent protein serine/threonine kinase activity, GO:0045737
- substantia nigra development, GO:0021762
- ciliary base, GO:0097546
- centriolar satellite, GO:0034451
- negative regulation of phosphatase activity, GO:0010923
- regulation of heart rate by cardiac conduction, GO:0086091
- ciliary transition zone, GO:0035869
- cyclin-dependent protein serine/threonine kinase regulator activity, GO:0016538
- protein localization to centrosome, GO:0071539
- response to radiation, GO:0009314
- negative regulation of cyclin-dependent protein serine/threonine kinase activity, GO:0045736
- non-motile cilium assembly, GO:1905515
- pericentriolar material, GO:0000242
- natural killer cell mediated cytotoxicity, GO:0042267
- gamma-tubulin binding, GO:0043015
- cellular response to glucagon stimulus, GO:0071377
- dynactin binding, GO:0034452
- centriole replication, GO:0007099
- photoreceptor connecting cilium, GO:0032391
- cellular response to extracellular stimulus, GO:0031668
- tau-protein kinase activity, GO:0050321
- cochlea development, GO:0090102
- protein kinase A regulatory subunit binding, GO:0034237
- oocyte maturation, GO:0001556
- mitotic cell cycle arrest, GO:0071850
- regulation of glycogen biosynthetic process, GO:0005979
- regulation of synaptic vesicle exocytosis, GO:2000300
- intraciliary retrograde transport, GO:0035721
- protein phosphatase inhibitor activity, GO:0004864
- positive regulation of Ras protein signal transduction, GO:0046579
- activation of protein kinase A activity, GO:0034199
- protein localization to cilium, GO:0061512
- MKS complex, GO:0036038
- adult heart development, GO:0007512
- positive regulation of phosphorylation of RNA polymerase II C-terminal domain, GO:1901409