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| GO:1903047 – mitotic cell cycle process | |
| GO:0043604 – amide biosynthetic process | |
| GO:0043603 – cellular amide metabolic process | |
| GO:0043043 – peptide biosynthetic process | |
| GO:0034248 – regulation of cellular amide metabolic process | |
| GO:0010608 – posttranscriptional regulation of gene expression | |
| GO:0006518 – peptide metabolic process | |
| GO:0006417 – regulation of translation | |
| GO:0006412 – translation | |
| GO:0000278 – mitotic cell cycle | |
| GO:2000134 – negative regulation of G1/S transition of mitotic cell cycle | |
| GO:2000036 – regulation of stem cell population maintenance | |
| GO:1905515 – non-motile cilium assembly | |
| GO:1902857 – positive regulation of non-motile cilium assembly | |
| GO:1902855 – regulation of non-motile cilium assembly | |
| GO:1902807 – negative regulation of cell cycle G1/S phase transition | |
| GO:1902403 – signal transduction involved in mitotic DNA integrity checkpoint | |
| GO:1902402 – signal transduction involved in mitotic DNA damage checkpoint | |
| GO:1902400 – intracellular signal transduction involved in G1 DNA damage checkpoint | |
| GO:1902117 – positive regulation of organelle assembly | |
| GO:1902017 – regulation of cilium assembly | |
| GO:0120162 – positive regulation of cold-induced thermogenesis | |
| GO:0120034 – positive regulation of plasma membrane bounded cell projection assembly | |
| GO:0090307 – mitotic spindle assembly | |
| GO:0072431 – signal transduction involved in mitotic G1 DNA damage checkpoint | |
| GO:0072422 – signal transduction involved in DNA damage checkpoint | |
| GO:0072413 – signal transduction involved in mitotic cell cycle checkpoint | |
| GO:0072401 – signal transduction involved in DNA integrity checkpoint | |
| GO:0072395 – signal transduction involved in cell cycle checkpoint | |
| GO:0071158 – positive regulation of cell cycle arrest | |
| GO:0071156 – regulation of cell cycle arrest | |
| GO:0051225 – spindle assembly | |
| GO:0045948 – positive regulation of translational initiation | |
| GO:0045727 – positive regulation of translation | |
| GO:0045724 – positive regulation of cilium assembly | |
| GO:0044819 – mitotic G1/S transition checkpoint | |
| GO:0044783 – G1 DNA damage checkpoint | |
| GO:0044774 – mitotic DNA integrity checkpoint | |
| GO:0044773 – mitotic DNA damage checkpoint | |
| GO:0031929 – TOR signaling | |
| GO:0031571 – mitotic G1 DNA damage checkpoint | |
| GO:0030330 – DNA damage response, signal transduction by p53 class mediator | |
| GO:0007052 – mitotic spindle organization | |
| GO:0006977 – DNA damage response, signal transduction by p53 class mediator resulting in cell cycle arrest | |
| GO:0006446 – regulation of translational initiation | |
| GO:0001829 – trophectodermal cell differentiation | |
| GO:0001825 – blastocyst formation | |
| GO:0001824 – blastocyst development | |
| GO:0000289 – nuclear-transcribed mRNA poly(A) tail shortening | |
| GO:0000288 – nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay | |

