
P-body, GO:0000932
gene silencing by miRNA, GO:0035195
interleukin-7-mediated signaling pathway, GO:0038111
phosphatidylinositol-mediated signaling, GO:0048015
RNA phosphodiester bond hydrolysis, GO:0090502
double-stranded RNA binding, GO:0003725
gene silencing by RNA, GO:0031047
neurogenesis, GO:0022008
RNA phosphodiester bond hydrolysis, GO:0090501
positive regulation of viral transcription, GO:0050434
B cell homeostasis, GO:0001782
RNA phosphodiester bond hydrolysis, GO:0090503
3′-5′-exoribonuclease activity, GO:0000175
somatic stem cell population maintenance, GO:0035019
regulation of proteolysis, GO:0030162
DNA-directed RNA polymerase II, GO:0005665
ribonuclease activity, GO:0004540
miRNA metabolic process, GO:0010586
poly(A)-specific ribonuclease activity, GO:0004535
posttranscriptional regulation of gene expression, GO:0010608
CCR4-NOT complex, GO:0030014
transcription factor TFTC complex, GO:0033276
T cell homeostasis, GO:0043029
lymph node development, GO:0048535
positive regulation of regulatory T cell differentiation, GO:0045591
negative regulation of programmed cell death, GO:0043069
cytoplasmic exosome (RNase complex), GO:0000177
nuclear-transcribed mRNA catabolic process, GO:0000288
DNA demethylation, GO:0080111
viral RNA genome replication, GO:0039694
regulation of stem cell population maintenance, GO:2000036