

proteasome complex, GO:0000502

- antigen processing and presentation of exogenous peptide antigen via MHC class I, GO:0002479
- regulation of cellular amino acid metabolic process, GO:0006521
- endopeptidase activity, GO:0004175
- proteolysis involved in cellular protein catabolic process, GO:0051603
- proteasome core complex, GO:0005839
- threonine-type endopeptidase activity, GO:0004298
- proteasome accessory complex, GO:0022624
- proteasome assembly, GO:0043248
- integrator complex, GO:0032039
- blastocyst development, GO:0001824
- proteasome core complex, GO:0019774
- positive regulation of proteasomal protein catabolic process, GO:1901800
- regulation of proteasomal ubiquitin-dependent protein catabolic process, GO:0032434
- proteasome regulatory particle, GO:0005838
- endopeptidase inhibitor activity, GO:0004866
- regulation of I-kappaB kinase/NF-kappaB signaling, GO:0043122
- proteasome binding, GO:0070628
- proteasome core complex, GO:0019773
- positive regulation of calcium ion transport, GO:0051928
- positive regulation of tumor necrosis factor biosynthetic process, GO:0042535
- proteasomal ubiquitin-independent protein catabolic process, GO:0010499
- calcium ion homeostasis, GO:0055074
- positive regulation of blood vessel endothelial cell migration, GO:0043536
- lateral ventricle development, GO:0021670
- lipopolysaccharide binding, GO:0001530
- Lys63-specific deubiquitinase activity, GO:0061578