

GO:0044085 – cellular component biogenesis	
GO:0022607 – cellular component assembly	
GO:0006508 – proteolysis	
GO:0075522 – IRES-dependent viral translational initiation	
GO:0072666 – establishment of protein localization to vacuole	
GO:0072665 – protein localization to vacuole	
GO:0072594 – establishment of protein localization to organelle	
GO:0071985 – multivesicular body sorting pathway	
GO:0071826 – ribonucleoprotein complex subunit organization	
GO:0070925 – organelle assembly	
GO:0070646 – protein modification by small protein removal	
GO:0061919 – process utilizing autophagic mechanism	
GO:0051603 – proteolysis involved in cellular protein catabolic process	
GO:0051345 – positive regulation of hydrolase activity	
GO:0051056 – regulation of small GTPase mediated signal transduction	
GO:0045324 – late endosome to vacuole transport	
GO:0044257 – cellular protein catabolic process	
GO:0043632 – modification-dependent macromolecule catabolic process	
GO:0043604 – amide biosynthetic process	
GO:0043547 – positive regulation of GTPase activity	
GO:0043328 – protein transport to vacuole involved in ubiquitin-dependent protein catabolic process via the multivesicular body sorting pathway	
GO:0043162 – ubiquitin-dependent protein catabolic process via the multivesicular body sorting pathway	
GO:0043087 – regulation of GTPase activity	
GO:0043043 – peptide biosynthetic process	
GO:0036258 – multivesicular body assembly	
GO:0036257 – multivesicular body organization	
GO:0032511 – late endosome to vacuole transport via multivesicular body sorting pathway	
GO:0032509 – endosome transport via multivesicular body sorting pathway	
GO:0022618 – ribonucleoprotein complex assembly	
GO:0022613 – ribonucleoprotein complex biogenesis	
GO:0019941 – modification-dependent protein catabolic process	
GO:0019081 – viral translation	
GO:0019080 – viral gene expression	
GO:0016579 – protein deubiquitination	
GO:0016236 – macroautophagy	
GO:0016197 – endosomal transport	
GO:0016050 – vesicle organization	
GO:0016032 – viral process	
GO:0010256 – endomembrane system organization	
GO:0007264 – small GTPase mediated signal transduction	
GO:0007034 – vacuolar transport	
GO:0007032 – endosome organization	
GO:0006914 – autophagy	
GO:0006518 – peptide metabolic process	
GO:0006511 – ubiquitin-dependent protein catabolic process	
GO:0006413 – translational initiation	
GO:0006412 – translation	
GO:0002183 – cytoplasmic translational initiation	
GO:0002181 – cytoplasmic translation	
GO:0001732 – formation of cytoplasmic translation initiation complex	

