azurophil granule lumen, GO:0035578

lysosomal lumen, GO:0043202 specific granule lumen, GO:0035580 RNA phosphodiester bond hydrolysis, GO:0090501 heparin binding, GO:0008201 hydrolase activity, GO:0004553 lipid storage, GO:0019915 insulin receptor substrate binding, GO:0043560 oligosaccharide catabolic process, GO:0009313 ribonuclease activity, GO:0004540 positive regulation of apoptotic signaling pathway, GO:2001235 hyaluronan catabolic process, GO:0030214 azurophil granule, GO:0042582 positive regulation of interleukin-1 beta secretion, GO:0050718 negative regulation of actin filament polymerization, GO:0030837 cell chemotaxis, GO:0060326 sulfuric ester hydrolase activity, GO:0008484 cofactor binding, GO:0048037 positive regulation of interleukin-8 secretion, GO:2000484 GTP biosynthetic process, GO:0006183 negative regulation of growth of symbiont in host, GO:0044130 purine ribonucleoside monophosphate biosynthetic process, GO:0009168 keratan sulfate catabolic process, GO:0042340 prostaglandin biosynthetic process, GO:0001516 fatty-acyl-CoA biosynthetic process, GO:0046949 oxaloacetate metabolic process, GO:0006107 hormone activity, GO:0005179 glycogen catabolic process, GO:0005980 collagen catabolic process, GO:0030574 immunoglobulin mediated immune response, GO:0016064 NLRP3 inflammasome complex, GO:0072559 exocytic vesicle, GO:0070382 positive regulation of apoptotic cell clearance, GO:2000427 interleukin-6 receptor binding, GO:0005138 peroxidase activity, GO:0004601 positive regulation of interleukin-6 secretion, GO:2000778 regulation of mast cell degranulation, GO:0043304 positive regulation of superoxide anion generation, GO:0032930 ATP-dependent microtubule motor activity, GO:0008569 antimicrobial humoral response, GO:0019730 NAD biosynthesis via nicotinamide riboside salvage pathway, GO:0034356 deaminase activity, GO:0019239 epidermis development, GO:0008544 antioxidant activity, GO:0016209 activation of phospholipase C activity, GO:0007202