

protein glycosylation, GO:0006486

- integral component of endoplasmic reticulum membrane, GO:0030176
- manganese ion binding, GO:0030145
- azurophil granule membrane, GO:0035577
- protein N-linked glycosylation, GO:0006487
- hydrolase activity, GO:0016798
- protein O-linked glycosylation, GO:0006493
- glycosphingolipid metabolic process, GO:0006687
- glycosaminoglycan metabolic process, GO:0030203
- GPI anchor biosynthetic process, GO:0006506
- glycosaminoglycan biosynthetic process, GO:0006024
- O-glycan processing, GO:0016266
- mannosyltransferase activity, GO:0000030
- galactosyltransferase activity, GO:0008378
- mannosylation, GO:0097502
- N-glycan processing, GO:0006491
- acetylglucosaminyltransferase activity, GO:0008375
- lung alveolus development, GO:0048286
- magnesium ion transmembrane transport, GO:1903830
- fucose metabolic process, GO:0006004
- protein O-linked fucosylation, GO:0036066
- polypeptide N-acetylgalactosaminyltransferase activity, GO:0004653
- oligosaccharide metabolic process, GO:0009311
- lysosomal transport, GO:0007041
- magnesium ion transmembrane transporter activity, GO:0015095
- epithelial cell development, GO:0002064
- sterol transporter activity, GO:0015248
- protein O-linked mannosylation, GO:0035269
- heparan sulfate proteoglycan biosynthetic process, GO:0015012
- chondroitin sulfate proteoglycan biosynthetic process, GO:0050650
- kidney morphogenesis, GO:0060993
- dolichyl diphosphate biosynthetic process, GO:0006489
- mesoderm formation, GO:0001707
- negative regulation of epidermal growth factor-activated receptor activity, GO:0007175
- gastrulation, GO:0007369
- oligosaccharyltransferase complex, GO:0008250
- acetylgalactosaminyltransferase activity, GO:0008376
- UDP-galactose:beta-N-acetylglucosamine beta-1, GO:0008499
- retina morphogenesis in camera-type eye, GO:0060042
- thyroid hormone metabolic process, GO:0042403
- angiogenesis involved in wound healing, GO:0060055
- cell-cell recognition, GO:0009988
- proteoglycan metabolic process, GO:0006029
- magnesium ion transport, GO:0015693
- nucleobase-containing small molecule catabolic process, GO:0034656
- ciliary rootlet, GO:0035253
- sialyltransferase activity, GO:0008373
- mannose metabolic process, GO:0006013
- alpha-N-acetylgalactosaminide alpha-2, GO:0001665
- hydrolase activity, GO:0016799
- aspartic endopeptidase activity, GO:0042500
- regulation of insulin receptor signaling pathway, GO:0046626
- sialylation, GO:0097503
- positive regulation of epithelial cell proliferation involved in wound healing, GO:0060054
- beta-N-acetylglucosaminylglycopeptide beta-1, GO:0003831