

**protein glycosylation, GO:0006486**

- integral component of endoplasmic reticulum membrane, GO:0030176
- Golgi cisterna membrane, GO:0032580
- hydrolase activity, GO:0016798
- cis-Golgi network, GO:0005801
- manganese ion binding, GO:0030145
- protein N-linked glycosylation, GO:0006487
- protein O-linked glycosylation, GO:0006493
- integral component of Golgi membrane, GO:0030173
- GPI anchor biosynthetic process, GO:0006506
- protein N-linked glycosylation via asparagine, GO:0018279
- glycosaminoglycan metabolic process, GO:0030203
- acetylglucosaminyltransferase activity, GO:0008375
- glycosaminoglycan biosynthetic process, GO:0006024
- keratan sulfate biosynthetic process, GO:0018146
- mesoderm formation, GO:0001707
- rough endoplasmic reticulum, GO:0005791
- Golgi transport complex, GO:0017119
- O-glycan processing, GO:0016266
- protein localization to organelle, GO:0033365
- galactosyltransferase activity, GO:0008378
- fucosylation, GO:0036065
- gastrulation, GO:0007369
- oligosaccharyltransferase complex, GO:0008250
- polypeptide N-acetylgalactosaminyltransferase activity, GO:0004653
- memory, GO:0007613
- heparan sulfate proteoglycan biosynthetic process, GO:0015012
- dolichyl-diphosphooligosaccharide-protein glycotransferase activity, GO:0004579
- prenyltransferase activity, GO:0004659
- fucosyltransferase activity, GO:0008417
- protein O-linked mannosylation, GO:0035269
- sialyltransferase activity, GO:0008373
- amyloid-beta metabolic process, GO:0050435
- N-glycan processing, GO:0006491
- sialylation, GO:0097503
- lung alveolus development, GO:0048286
- mannosyltransferase activity, GO:0000030
- oligosaccharide metabolic process, GO:0009311
- Notch signaling involved in heart development, GO:0061314
- carbohydrate biosynthetic process, GO:0016051
- protein O-linked fucosylation, GO:0036066
- fucose metabolic process, GO:0006004
- mannosylation, GO:0097502
- Notch receptor processing, GO:0007220
- regulation of resting membrane potential, GO:0060075
- beta-galactoside (CMP) alpha-2, GO:0003836
- amyloid precursor protein catabolic process, GO:0042987
- alpha-mannosidase activity, GO:0004559
- beta-N-acetylglucosaminylglycopeptide beta-1, GO:0003831
- protein targeting to ER, GO:0045047
- dystrophin-associated glycoprotein complex, GO:0016010
- glycosphingolipid biosynthetic process, GO:0006688
- mannosyl-oligosaccharide 1, GO:0004571
- magnesium ion transport, GO:0015693
- mannose metabolic process, GO:0006013
- alpha-N-acetylgalactosaminide alpha-2, GO:0001665
- respiratory gaseous exchange, GO:0007585
- positive regulation of organ growth, GO:0046622