

protein glycosylation, GO:0006486

- integral component of endoplasmic reticulum membrane, GO:0030176
- protein N-linked glycosylation, GO:0006487
- manganese ion binding, GO:0030145
- hydrolase activity, GO:0016798
- protein O-linked glycosylation, GO:0006493
- glycosphingolipid metabolic process, GO:0006687
- GPI anchor biosynthetic process, GO:0006506
- Notch receptor processing, GO:0007220
- glycosaminoglycan biosynthetic process, GO:0006024
- lung alveolus development, GO:0048286
- N-glycan processing, GO:0006491
- galactosyltransferase activity, GO:0008378
- oligosaccharyltransferase complex, GO:0008250
- gastrulation, GO:0007369
- UDP-galactose:beta-N-acetylglucosamine beta-1, GO:0008499
- heparan sulfate proteoglycan biosynthetic process, GO:0015012
- fucosylation, GO:0036065
- dolichyl diphosphate biosynthetic process, GO:0006489
- polypeptide N-acetylgalactosaminyltransferase activity, GO:0004653
- mesoderm formation, GO:0001707
- protein targeting to ER, GO:0045047
- dolichyl-diphosphooligosaccharide-protein glycotransferase activity, GO:0004579
- transferase activity, GO:0016765
- respiratory gaseous exchange, GO:0007585
- glycosphingolipid biosynthetic process, GO:0006688
- protein O-linked mannosylation, GO:0035269
- acetylglucosaminyltransferase activity, GO:0008375
- UDP-N-acetylglucosamine metabolic process, GO:0006047
- protein O-linked fucosylation, GO:0036066
- mannosyl-oligosaccharide 1, GO:0004571
- oligosaccharyl transferase activity, GO:0004576
- mannose metabolic process, GO:0006013
- acetylgalactosaminyltransferase activity, GO:0008376
- keratan sulfate biosynthetic process, GO:0018146
- beta-N-acetylglucosaminylglycopeptide beta-1, GO:0003831