protein glycosylation, GO:0006486

integral component of endoplasmic reticulum membrane, GO:0030176 cellular protein modification process, GO:0006464 manganese ion binding, GO:0030145 hydrolase activity, GO:0016798 protein O-linked glycosylation, GO:0006493 protein N-linked glycosylation, GO:0006487 GPI anchor biosynthetic process, GO:0006506 glycosaminoglycan metabolic process, GO:0030203 N-glycan processing, GO:0006491 glycosphingolipid metabolic process, GO:0006687 oligosaccharyltransferase complex, GO:0008250 keratan sulfate biosynthetic process, GO:0018146 mesoderm formation, GO:0001707 glycosaminoglycan biosynthetic process, GO:0006024 respiratory gaseous exchange, GO:0007585 acetylglucosaminyltransferase activity, GO:0008375 oligosaccharide metabolic process, GO:0009311 mannose metabolic process, GO:0006013 protein O-linked mannosylation, GO:0035269 dolichyl-diphosphooligosaccharide-protein glycotransferase activity, GO:0004579 protein targeting to ER, GO:0045047 mannosylation, GO:0097502 galactosyltransferase activity, GO:0008378 magnesium ion transmembrane transporter activity, GO:0015095 O-glycan processing, GO:0016266 fucosyltransferase activity, GO:0008417 sialylation, GO:0097503 beta-N-acetylglucosaminylglycopeptide beta-1, GO:0003831 sialyltransferase activity, GO:0008373 magnesium ion transmembrane transport, GO:1903830 transferase activity, GO:0016765 alpha-mannosidase activity, GO:0004559 magnesium ion transport, GO:0015693 chondroitin sulfate metabolic process, GO:0030204 heparan sulfate proteoglycan biosynthetic process, GO:0015012 prenyltransferase activity, GO:0004659 mannosyltransferase activity, GO:0000030 acetylgalactosaminyltransferase activity, GO:0008376 dolichyl diphosphate biosynthetic process, GO:0006489 Notch receptor processing, GO:0007220 GDP-mannose biosynthetic process, GO:0009298 oligosaccharyl transferase activity, GO:0004576 mannosyl-oligosaccharide 1, GO:0004571 polypeptide N-acetylgalactosaminyltransferase activity, GO:0004653 cerebellar Purkinje cell layer development, GO:0021680 oxidoreductase activity, GO:0016628 UDP-N-acetylglucosamine metabolic process, GO:0006047 glycolipid metabolic process, GO:0006664