

MPSSSMKLF AALLVACLAQTSMAAIVVCKMDAQDGTLTAAACSVGVSGQPISLVGPGSGQQQLTGSQVSYTLDVNARSTL	80
FECASEDDLIIIDSSQYSSQTLNNCPEPLLELRGCSNAILSNTFISITRSTAQPGCTISKYGPCVAVVGAASQETDWSF	160
SSLANTFTSTICSSISATSGRLGGAF AFEHNDSPGAMSAVVKGSFTTSTACDFGGAIHSANASLTLTDSTFTGT LAVDGG	240
AVQFVGT NATVAPIQKLQVKSSTFTSNTAVTTGGIIQVTGGAVSIDGSTFTNGEAQIGQCVWLDKCESYTENQITGNTWT	320
GCAKPESPPI SWCKAHDGNNWTT CGMEGPRECY	
.....	80
..... N	160
.....	240
..... N	320
.....	400

NetNGlyc 1.0: predicted N-glycosylation sites in Sequence

