

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
D3ZUJ8	D3ZUJ8_RAT	Rattus norvegicus	dolichyl-phosphate-mannose--protein mannosyltransferase	23.439218	NaN	NaN	38843836	MREAAAALAEELGKDREAGGAAAKVR PPRSRSVADYDSVFCRVLLRSTVSEA FLNHLLAEPPSGEVGAGHSAAGRSS GNICVLGKLCFGAAGAPADLTFEAW LMMEGKMADINFKEVTLIVGVAAAC YWNSLFCGFVDDVSAILDNKDLHP STPLKTLFQNDFWGTPMSEERSHKS YRPLTVLTFRLNYLLELKPMSYHLL NTVFFHAVSVIFLKVCRFLDKRGSV IAALLFAVHPIHTEAVTGVVGRAELL SSVFFLAAFLSYTKSGPDNSIVWTP IALTVLLVAVATLCKEQTIVVVICCV YEVFVAQGYTLPLLCVAVAGQFLRGK GSIPFMSMLQTLIKLIVLMFSTLLVVI RVQVIQSQLPVFTRFDNPAAVSPTPT RQLTFNYLLPVNAWLLNPSLCCD WTMGTIPLIESFLDVRNLATFAFFCF LGALGIFSLRYPGDSSKTVLMAISLM ALPFIPASNLFFPVGFVVAERVLVYP SMGFCLVAHGQKISNKSVLKKS WVCLSMVILTHALKTLHRNWDWES EYTLFMSALKVNNNAKLWNNVGH ALENEKNFEKALKYFLOATHVQPDD IGAHMNVGRTYKNLNRKEAESY MLAKSLMPQIIPGKKAARIAPNHLN VYINLANLIRANESRLEEADQLYRQA ISMRFDFKQAYISRGELLLKMNKPL KAKEAYLKALELDRNNADLWYNLAI VYIELKEPDEALKNFNRALELNSRH KLALFNSAILMQESGEVLRPEARL RLLNYNINEEPQDANGYFNLGMLAM DDKDESEAEWTKRAIKLQDFRSA LFNLALLYQTAKELKALPILEELRY YPDHTKGLILKGDILMNQKKDIAGAK KCFEKILEMDPSNVQKHNLCVVYF EEKDLLKAERCLVETLALAPHEEYIQ RHLSIVRDKIASSGIVDQPLSPVDKTS GTEEKDEVPSDEVKEIISESRPAQIK TNDNRKSQSNGKSTGNADQDAPHK TTKDIKEIEKKRVAALKRLEEIERILN GE	False	False	0.382	False	False	4.366	1.142	2.365	False