	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P32855	SEC8_YEAST	Saccharomyces cerevisiae	Exocyst complex component SEC8	24.475439	NaN	S16;S19	33229814	MDYLKPAQKGRRRGLSINSLSETQQ SAMNSSLDHLQNDLNRINLQWNRI LSDNTNPLELALAFLDDTSVGLGHR YEEFNQLKSQIGSHLQDVVNEHSQV FNTNVASYGKAVSSIMQAQEQTLNL KNCLKEANEKITTDKGSLQELNDNN LKYTKMIDVLVNIEELLQIPEKIEENI RKENFHQVQILLERGFILMNNKSLK TVEILKPINQQLELQEHLLFNNLIEEI HDIMYSKSNKTNFTRVTNNDIFKIIS ISHNGFTSLENYLYNIVNIDIMEHSK TINKNLEQFIHDQSLNKGNIMLQEN AATQAPLAPSRNQENEGFNRIGFLL KTINNINKLPVAFNIITERAKEEIHNII VKSTESIRSKHPSLLKMATSLKNDN HFGLPVQDILSIILRECFWEIFLKLLY AIQCHRAIFEMSNILQPTSSAKPAFK FNKIWGKLLDEIELLLVRYINDPELIS SNNGSIKPINGATNNAPTLPKRKNP KIFSLEYNIEDNSSVKDQAFELKALL KDIFPGFSVSSNMDLDSIYVKDESFE QDEPLVPPSVFNMKVILDPFLLFTQS TSTIVPSVLTQNTISSLTFFDDYMNK SFLPKIQMTMDYLFTVEVESNNPYA LELSDENHNIFKTALDFQRLFYNLL NVFNTANTFREKISYCILDLLNHFYN YYLGLFNSLIGTSDRHLTRKIITAWL QNGILMDQEQKILNGDETLFHEESI ELFKEIPHFYQAGKGLSKSDLFNNL TLDTILQFSASVLWILNWLPGLKKAI NIDEVSQEPMLDADRLRSSWTFSES MDLNYSNPSSSPNSLGNLKILLDDK ASKKFDETIDGFKTLKFKLITILRFNI RALCIYDIGSFFQNTKIWNMDVGSIE LDQNIASLISELRRTESKLKQQLPEK EKNSIFIGLDIVNNYALIKGAKSIKVL NHNGIKKMLRNVNVLQHAYRNLSS EPSKINMNVTMNFYSLCGSSEAELF EYIKDNELPHCSVEDLKTILRLQFSE EMHRQLKRQSTSSTKGSIKPSNKRY TEALEKLSNLEKEQSKEGARTKIGKL KSKLNAVHTANEK	None	None	None	None	None	None	None	None	None