

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P32855	SEC8_YEAST	Saccharomyces cerevisiae	Exocyst complex component SEC8	9.365641	NaN	S16;S19	33229814	MDYLKPAQKGRRRGLSINSLSETQQ SAMNSSLDHLQNDLNRINLQWNRI LSDNTNPLELALAFLLDDTSVGLGHR YEEFNQLKSQIGSHLQDVNEHSQV FNTNVASYGKAVSSIMQAQEQLNL KNCLKEANEKITTDKGLQELNDNN LKYTKMIDVLVNIIEELLQIPEKIEENI RKENFHQVQILLERGFILMNNKSLK TVEILKPINQQLELQEHLLFNNLIEEI HDIMYSKSNKTNFTRVTNNDIFKIIS ISHNGFTSLENYLYNIVNIDIMEHSK TINKNLEQFIHDQSLNKGNIMLQEN AATQAPLAPSRNQENEGFNRIQFLL KTINNINKLPVAFNIITERAKEEIHNI VKSTESIRSKHPSLLKMATSLKNDN HFGLPVQDILSIILRECFWEIFLKLLY AIQCHRAIFEMSNIHQPTSSAKPAFK FNKIWGKLLDEIELLVRYINDPELIS SNNGSIKPINGATNNAPTLPKRKNP KIFSLEYNIEDNSSVKDQAFELKALL KDIFPGFSVSSNMDLDSIYVKDESFE QDEPLVPPSVFNMKVILDPFLLFTQS TSTIVPSVLTQNTISLTFDDYMNK SFLPKIQMTMDYLFTVEVESNNPYA LELSDENHNIFKTALDFQRLFYNLL NVFNTANTFREKISYILDLLNHFYN YYLGLFNSLIGTSDRHLTRKIITAWL QNGILMDQEQKILNGDETLFHEESI ELFKEIPHFYQAGKGLSKSDLFNNL TLDTILQFSASVLWILNWLPLGKKAI NIDEVSQEPMLDADRLRSSFSE MDLNYSNPSSPNSLGNLKILLDDK ASKKFDEIDGFKTLKFLITILRFNI RALCIYDIGSFFQNTKIWNMDVGSIE LDQNIASLISELRRTESKLKQQLPEK EKNSIFIGLDIVNNYALIKGAKSIKVL NHNGIKMLRNVNVLQHAYRNLS EPSKINMNVMTMNFYSLCGSSEAE EYIKDNEIPHCSVEDLKTILRLQFSE EMHRQLKRQSTSSSTKGSIKPSNKRY TEALEKLSNLEKEQSKEGARTKIGKL KSKLNAVHTANEK