

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
P32259	MED16_YEAST	Saccharomyces cerevisiae	Mediator of RNA polymerase II transcription subunit 16	24.475439	NaN	NaN	33229814	MMLGEHLMSWSKTGIIAYSDSQSS NANICLTFLESINGINWRFHTPQKYV LHPQLHEVQYQESSSTLSTHSTTTSV NGSTTAGVVGSTPNFGGNSNKSPPQF FYNISSIHWNWVSLPGDMLAVCD ELGNMTMLITGQRPRATTYEKLTM VFQDNVYKIYNHVMPLKPVDKLKPM NIERKQTRKEYNTSILEFRWLTSSKS VIVSQFCADFSSNTYRSRAQQVPPY GVYHPPFIKYACLAIKRNQIDFWYQ FSNSKDHKKITLQLLDTSNQRFKDL QWLEFARITPMNDDQCMILTYSKL SKNISFYKLVNWNLNATKPNVLN DPSLKIQFILSTTLDPTDDEGHVLLK ENLHVSKSSIEKDPSPAILVLYNVC DTSKSLVKRYRLAPTQLSAEYLVLK PDLNIDRNNSTNQIFQSRRYNLRRH SDIVLDKKVTLITSEMFDVFSFYFE DGTIESYNQNDWKLETERLISQSOL GKFKNIASPLSAGFNKGLPLPPSV EWMKVSPSMCGVIVKQYNKKWPOF YAAVQKNYADPEKDSINATALAFGYV KSLHKQISAEDLTIAAKTHILRISFLD RKRAKEFITTLKSLYSFFNISPDAPK EIMDKIITSRPLQKIMLLQLELGSCFS QENIEEMARVILYLKNVLFANVVA RNFHFAIEQISNNSNQQNPKLFQT IFSKQDLIHSLIPVAKWVFKFITLQ EILILINDPTNKEYTLVHGIFGAKMS RTLILSILNEIKKVTQIVAKFETSYP LNESSFLKLVSESPVDFEFETFL VDVNNKFIALCEQQPSQEREFSLV KAEIPPEYAKVGDFLQYANNAVISH ANAAAVYFADTSGLKISNSEFFNPEI FHLLQPLEEGLIIDTKLPIKNRTSKS FSKLLYDDVTCDKLSEISDGKLR CSRCSVTRAGNISSDKTIVPTSQT KRWPTMYTRLCICSGMLFEMDG	None	None	None	None	None	None	None		

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