	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	extrac regio
0	P19659	MED15_YEAST	Saccharomyces cerevisiae	Mediator of RNA polymerase II transcription subunit 15	24.475439	NaN	S335;S736;S752;S783;S785;S789;T793;S831;S1003;S1008;S1018;S1034	33229814	MSAAPVQDKDTLSNAERAKNVNGL LQVLMDINTLNGGSSDTADKIRIHA KNFEAALFAKSSSKKEYMDSMNEK VAVMRNTYNTRKNAVTAAAANNNI KPVEQHHINNLKNSGNSANNMNV NMNLNPQMFLNQQAQARQQVAQQ LRNQQQQQQQQQQQRRQLTPQQ QQLVNQMKVAPIPKQLLQRIPNIPPN INTWQQVTALAQQKLLTPQDMEAAK EVYKIHQQLLFKARLQQQQAQAQAQ ANNNNNGLPQNGNINNNINIPQQQ QMQPPNSSANNNPLQQQSSQNTVP NVLNQINQIFSPEEQRSLLQEAIETC KNFEKTQLGSTMTEPVKQSFIRKYIN QKALRKIQALRDVKNNNNANNNGS NLQRAQNVPMNIIQQQQQQNTNNN DTIATSATPNAAAFSQQQNASSKLYQ MQQQQQAQAQAQAQAQAQAQAQA QAQAAQAAQAQAQAQAQAQAQAQA QAQAQAQAQAQAQAQAQAHAQHQP SQQPQQAQQQPNPLHGLTPTAKDV EVIKQLSLDASKTNLRLTDVTNSLSN EEKEKIKMKLKQGQKLFVQVSNFAP QVYIITKNENFLKEVFQLRIFVKEILE KCAEGIFVVKLDTVDRLIIKYQKYWE SMRIQILRRQAILRQQQQMANNNG NPGTTSTGNNNNIATQQNMQQSLQ QMQHLQQLKMQQQQQQQQQQQQ QQQQQQQQQQQHIYPSSTPGVANY SAMANAPGNNIPYMNHKNTSSMDF LNSMENTPKVPVSAAATPSLNKTIN GKVNGRTKSNTIPVTSIPSTNKKLSIS NAASQQPTPRSASNTAKSTPNTNPS PLKTQTKNGTPNPNNMKTVQSPMG AQPSYNSAIIENAFRKEELLLKDLEIR KLEISSRFKHRQEIFKDSPMDLFMS TLGDCLGIKDEEMLTSCTIPKAVVDH INGSGKRKPTKAAQRARDQDSIDISI KDNKLVMKSKFNKSNRSYSIALSNV AAIFKGIGGNFKDLSTLVHSSSPSTS SNMDVGNPRKRKASVLEISPQDSIA SVLSPDSNIMSDSKKIKVDSPDDPF MTKSGATTSEKQEVTNEAPFLTSGT SSEQFNVWDWNNWTSAT	None	None	None	None	None	None	None	None	None