

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P53552	THO2_YEAST	Saccharomyces cerevisiae	THO complex subunit 2	9.365641	NaN	NaN	33229814	MAEQTLLSKLNALSQKVIPPASPSQA SILTEEVRNWPERSKTLCSDFTALE SNDEKEDWLRTLFIELDFINKNDE NSPLKLSDVASFTNELVNERQVSQ ASIVGKMFIAVSSTVNPINDLTTISLC KLIPSLHEELFKFSWISSKLLNKEQT TLRHLKSKYELKKYNLLVENS GYGQLVALLILAYYDPDNFSKVSAYL KEYYHIMGKYSLDSIRTLDVILNVSSQ FITEGYKFFIALLRKSDSWPSSHVAN NSNYSSLNEGGMIAANIISFNLSQ YNEEVDKENYERYMDMCCILLKNG FVNFYSIWDNVKPEMEFLQEYIQNL ETELEEESTKGVENPLAMAAALSTE NETDEDNALVVNDVNMKDKISEE TNADIESKQKQKTQODILLFGKIKLL ERLLIHGCVIPVIHVLKQYPKVLYVSE SLSRYLGRVFEYLLNPLYTSMSSGE SKDMATALMITRIDNGILAHKPRLIH KYKTHEPFESLELNSSYVFYSEWN SNLTPFASVNDLFENSHIYLSIIGPYL GRIPTLLSKISRIGVADIQKNHGSESL HVTIDKWIDYVRKFIFPATSLQNNP IATSEVYELMKFFPFKRYFIYNEM MTKLSQDILPLKVSFNKAEREAKSIL KALSIDTIAKESRRFAKLISTNPLASL VPAVKQIENYDKVSELVVYTTKYFND FAYDVLQFVLLLRLTYNRPVAVQFDG VNQAMWVQRLSIFIAGLAKNCPNM DISNIITYILKTLHNGNIIAVSILKELII TVGGIRDLEVNMMKQLLMLNSGSP LKQYARHLYDFRDDNSVISSRLTSF FTDQSAISEIILLTYTLNLKANTQNS HYKILSTRCDEMNTLLWSFIELIKHC LKGKAFEENVLPVELNRFHLSTP WTFHIWRDYLDNQLNSNENFSIDEL IEGAEFSDVDLTKISKDLFTTFWRLS LYDIHFDKSLYDERKNALSGENTGH MSNRKKHLIQNQIKDILVTGISHQRA FKKTSEFISEKSNVWNKDCGEDQIKI FLQNCVVPRVLFSPSDALFSSFFIFM AFRTENLMSILNTCITSNILKTLLFC CTSSEAGNLGLFFTDVLKLEKMRL NGDFNDQASRKLYEWHVSVITEQVID LLSEKNYMSIRNGIEFMKHVTSVFP VKAHIQLVYTTLEENLINEEREDIK LPSSALIGHLKARLKDALELDEFCTL TEEEAEQKRIREMELEEIKNYETAC QNEQKQVALRKQLELNKSQRLQND

								PPKSVASGSAGLNSKDRYTYSRNEP VIPTKPSSSQWSYSKVTRHVDDINHY LATNHLQKAISLVENDDETRNLRLK SKQNMPIFDFRNSTLEIFERYFRTL QNPQNPDFAEKIDSLKRYIKNISREP YDITSSYSEAAPEYTKRSSRYSGN AGGKDGYSNRYRGPNSDRSAPKNI KPISSYAHKRSELPTRPSKSKTYNDR SRALRPTGPDRGDGFDQRDNRLREE YKNSSQRSQRLFPEKPFQEGKDSS KANPYQASSYKRDSPSENEEKPNKR FKKDETIRNKFQTQDYRNRTRDSGAA HRANENQRYNGNRKSNTQALPQGP KGGNYVSRYQR
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