

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P53010	PAN2_YEAST	Saccharomyces cerevisiae	PAN2-PAN3 deadenylation complex catalytic subunit PAN2	24.475439	NaN	NaN	33229814	MNNWQHFFNNPVDLSEHLKPYFR FDNRDKEITAISFDEKANLIWSGDSY GCISSYDPTFQLYTRYRGHIGGNSVK DILSHRDGILSISEDLSLHFANRRGVT KLNLTSIDIAAFSELNTMCYSPHSLK NNIYCGGDNTNWGIASIDLNRGCLD SLLNYSSKVKLMCSNKNVLSIGRQT GTVDLLDPTSNRTIKSFNAHSASISA MDLRDNTLVTVGKSKRFYNYADPF VNVYDLRTMRQLPPVFSKGGTTMGS GGADFVQLHPLLPTVMIVASSGSF DFIDLSNPTLRTQYVHPCQSIKKLCL SPNGDVLGILEADNHLDTWRRSSN NMGMTNTPEMLAYPDYFNDITS GPISVDDETYPLSSVGMPYYLDKLLS AWPPVVFKEGTIPQLTGKSPLPSSG KLKSNLAVISSQNEKLSTQEFPLLR DRTKYGMRAIPDYVCLDIRKQITS GLETSDIQTYTSINKYEVPPAYSRLPL TSGRFGTDNFDFTPFNNTSEYGLDP DVDNHYTNAIQLYRFIPEMFNFVVG CLKDENFETLLDGLYFDMMERS HGKICSSSNFQASLKLSTDKRQLEN GEPQEHLEEYLESLECIRESIEDFNSS ESIKRNMPQKFNRFLLSQLIKEAQ TVNHNITLNQCFGLETEIRTECSCD HYDTTVKLLPSSLISGINKTVIKQLNK KSNQONILPYIEYAMKNVTQKNSICP TCGKTEITQECTVKNLPSVLSLELS LLDTEFNSIRSSKNWLTSEFYGSIIK NKAVLRSTASELKGTSHFYKYLNGY VAKITDNNNETRLVTVYKKNPKEN CFKWLMFNDYLVVEITEEALKMTY PWKTPEIIYCDAAELRKPFFSVDYYS INYDILFRDYFANGIRDATARREYKLL THDEAPKSGTLVAIDAEFVSLQSELC EIDHQGIRSIIRPKRTALARISIRGEE GELYGVFVDDYVNTNHIEDYLTR YSGILPGDLDPKSTKRLVRRNVVYR KVVLLMQLGCVFVGHGLNNDFKHI NINVPRNQIRDTAIYFLOGKRYLSLR YLAYVLLGMNIQEGNHDSIEDAHTA LILYKYLHLKEKAIFEKVLNSVYEE GRAHNFKVPETSKG	None	None	None	None	None	None	None		