

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P53010	PAN2_YEAST	Saccharomyces cerevisiae	PAN2-PAN3 deadenylation complex catalytic subunit PAN2	9.365641	NaN	NaN	33229814	MNNWQHFFNNPVDLSEHLKPKPYFR FDNRDKEITAI SFDEKANLIWSGDSY GCISSYDPTFQLYTRYRGHIGGNSVK DILSHRDGILSISEDLSLHFANRRGVT KLNLT SIDIAAFSELNTMCYSPHSLK NNIYCGGDNTNWGIASIDLNRGCLD SLLNYSSKVKLMCSNNKVL SIGRQT GTVDLLDPTSNRTIKSFNAHSASISA MDLRDNTLVTVGKSKRFYNLYADPF VNVYDLRTMRQLPPVSFSKGTTMGS GGADFVQLHPLLPTVMIVASSSGSF DFIDLSNPTLRTQYVHPCQSIKKLCL SPNGDVLGILEADNHLDTWRRSSN NMG MFTNTPEMLAYPDYFNDITSD GPISVDDETYPLSSVGM PYYLDKLLS AWPPVVF KSEGTIPQLTGKSP LPSSG KLKSNLAVISSQNEKLSTQEFLLRY DRTKYGM RNAIPDYVCLRDIRKQITS GLETSDIQTYTSINKYEVPPAYSRLPL TSGRFGTDNFDFTPFNNT EYSGLDP DVDNHYTNAIIQLYRFIPEMFN FVVG CLKDENFETLLTDLGYLFDMMERS HGKICSSSNFQASLKS LTKRQLEN GEPQEHLEEYLESLCIRE SIEDFNSS ESIKRNMPQKFNRFLLSQLIKEEAQ TVNHNITLNQC FGLETEIRTECSCD HYD TTVKLLPSLSISGINKTVIKQLNK KSNQNILPYIEYAMKNVTQKNSICP TCGKTETITQECTVKNLPSVLSLELS LLDTEFSNIRSSKNWLTSEFYGSIK NKAVLRSTASELKGTSHIFKYELNGY VAKITDNNNETRLVTVYVKKYNPKN CFKWLMFNDYLVEITEEEALKMTY PWKTPEIIICYDAEELRKPFVSVD TYS INYDILFRDYFANGIRDARREYKLL THDEAPKSGTLVAIDAEFVSLQSELC EIDHQGIRSIIRPKRTALARISIIRGEE GELYGVPFVDDYVVNTNHIEDYLTR YSGILPGDL DPEKSTKRLVRRNVVYR KVWLLMQLGCVFVGHGLNNDFKHI NINVPRNQIRD TAIYFLQ GKRYLSLR YLAYVLLGMNIQEGNHDSIEDAHTA LILYKKYLHLKEKAIFEKVLNSVYEE GRAHNFKVPETSKG