

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
P25339	PUF4_YEAST	Saccharomyces cerevisiae	Pumilio homology domain family member 4	24.475439	NaN	T205;T212;T252;S256	33229814	MSTKGLKEEIDDVPSVDPVVSETVN SALEQLQLDDPEENATSNAFANKVS QDSQFANGPPSQMFPHPQMMGGM GFMPYSQMMQVPHNCPFFPPPDF NDPTAPLSSSPLNAGGPPMLFKNDS LPFQMLSSGAAVATOGGQNLNPLIN DNSMKVLPASADPLWTHSNVPGSA SVAIEETTATLQESLPSKGRESNKA SSFRRQTFHALSPTDLINAANNVTLN KDFQSDMQNFSKAKKPSVGANNTA KTRTQISISFDNTPSSTSFIPTNSVSE KLSDFKIETSKEDLINKTAPAKKESP TTYGAAYPYGGPLLQPNPIMPGHPH NISSPIYGIRSPFPNSYEMGAQFQPF SPILNPTSHSLNANSPIPLTQSPIHLA PVLNPSNSVAFSDMKNDGGKPTT DNDKAGPNVRMDLINPNLGPSMQP FHILPPQNTPPPPWLYSTPPPFNA MVPPHLLAQNHMPLMNSANNKHH GRNNNSMSSHNDNDNIGNSNYNN KDTGRSNVGMKMKNMKNSYHGYYN NNNNNNNNNNNNNSNATNSNS AEKQRKIESSRFADAVLDQYIGSIH SLCKDQHGCERFLQKQLDILGSKAAD AIFEETKDYTVELMTDSFGNYLIQKL LEEVTTQQRIVLTKISSPHFVEISLNP HGTRALQKLEICKTDEEAQIVVDSL RPYTVQLSKDLNGNHVIQKCLQRLK PENFQFIFDAISDSCIDIATHRHGCC VLQRCLDHGTTEQCDNLCKLLALV DKLTLDPFGNYVVQYIITKEAEKNKY DYTHKIVHLLKPRAIELSIHKFGSNVI EKILKTAIVSEPMILEILNNGGETGIQ SLLNDSYGNVYLQTALDISHKQNDY LYKRLSEIVAPLLVGPINRTPHGKRRI GMLHLDS	None	None	None	None	None	None	None	None	

0