

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi appa
Q8TB72	PUM2_HUMAN	Homo sapiens	Pumilio homolog 2	17.150651	S85;S589;S644(Q8TB72-4);S664	S67;S82;S102;S136;S178;S182;T184;T396;S587;S592;S684;S700	30379171;31492838;33214551;34019948;35254053;35289036;30059200;22121020;28510447;38665916	MNHDFQALALESRGMGELLPTKKF WEPDDSTKDGQKIGIFLGDDEWRET AWGASHHSSQPMVQRRSGQGPFH GNSEVNAILSPRSESGGLGVSMVEY VLS SSPADKLD SFRFRKGNFGRDAE TDGPEKGDQK GKASPF EEDQNRDL KQGD DDDSKINGRGLPNGMDADCK DFNRTPGSRQASPT EVVERLGPNTN PSEGLGPLNPTANKPLVEEFSNPE TQNL DAMEQVGL ESLQFDY PGNQV PMDSSGATVGLFDYNSQQQLFORT NALTVQQLTAAQQQYALAAAQQPH IAGVFSAGLAPA AFV PNPYIISAAPPG TDPYTAAGLAAAATLAGPAVVPQYY GVPWGVYPANL FQQQAAAAANNTA SQQAASQAQPGQQVLRAGAGQRP LTPNQGGQQQAESLAAAAANPTL AFGQGLATGMPGYQLAPTAYYDQT GALVVGPGARTGLGAPVRLMAPTPV LISSAAAQAAAAAAGGTASSLTGST NGLFRPIGTOPPPQQQPPSTNLQS NSFYGSSSLTNSOSSSLFSHGPGQ PGSTSLGFGSGNSLGAAGSALSGFG SSVSSASSATRRESLSTSSDLYKR SSSSLAPIGQPFYNSLGFSSSPSIG MPLPSQTPGHSLTPPPSLSSHGSSSS LHLGGLTNGSGRYISAAPGAEAKYRS ASSTSSLFSSSQLFPPSRLRYNRSD IMPSGRSRLLEDFRNRFNQLQLRD LIGHIVEFSQDQHGSRFIQKLERAT PAERQMVFNELQAAYQLMTDVFV NYVIQKFFEFGLDQKLALATRIRGH VLPLALOMYGCRVIQKALESISSDQ VISEMVKELDGHVLCVKDQNGNH VVQKCI ECVPQSLQFIIDAFKGVF VLSTHPYGCRVIQRILEHCTAEQTLPI LEELHQHTEQLVQDQYGNVVIQHV EHGRPEDKSKIVSEIRKVLALSQHK FASNVEKCVTHASRAERALLIDEVC CQNDGPHSALYTMMDQYANYVVQ KMIDMAEPAQRKIMHKIRPHITTLR KITYGKHILAKLEKYLLKNSPDLGPI GGPPNGML	True	False	4.706	4.71	1.721	1.386	0.724