	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regi
0	Q9ULJ6	ZMIZ1_HUMAN	Homo sapiens	Zinc finger MIZ domain-containing protein 1	12.21516	NaN	NaN	35138101;34019948;33214551;38665916	MNSMDRHIQQTNDRLQCIKQHLQN PANFHNAATELLDWCGDPRAFQRP FEQSLMGCLTVVSRVAAQQGFDLDL GYRLLAVCAANRDKFTPKSAALLSS WCEELGRLLLLRHQKSRQSDPPGKL PMQPPLSSMSSMKPTLSHSDGSFPY DSVPWQQNTNQPPGSLSVVTTVWG VTNTSQSQVLGNPMANANNPMNP GGNPMASGMTTSNPGLNSPQFAGQ QQQFSAKAGPAQPYIQQSMYGRPNY PGSGGFGASYPGGPNAPAGMGIPPH TRPPADFTQPAAAAAAAAVAAAAATA TATATATVAALQETQNKDINQYGPM GPTQAYNSQFMNQPGPRGPASMGG SMNPASMAAGMTPSGMSGPPMGM NQPRPPGISPFGTHGQRMPQQTYPG PRPQSLPIQNIKRPYPGEPNYGNQQY GPNSQFPTQPGQYPAPNPPRPLTSP NYPGQRMPSQPSSGQYPPPTVNMG QYYKPEQFNGQNNTFSGSSYSNYSQ GNVNRPPRPVPVANYPHSPVPGNPT PPMTPGSSIPPYLSPSQDVKPPFPPDI KPNMSALPPPPANHNDELRLTFPVR DGVVLEPFRLEHNLAVSNHVFHLRP TVHQTLMWRSDLELQFKCYHHEDR QMNTNWPASVQVSVNATPLTIERG DNKTSHKPLHLKHVCQPGRNTIQIT VTACCCSHLFVLQLVHRPSVRSVLQ GLLKKRLLPAEHCITKIKRNFSSVAA SSGNTTLNGEDGVEQTAIKVSLKCPI TFRRIQLPARGHDCKHVQCFDLESY LQLNCERGTWRCPVCNKTALLEGLE VDQYMWGILNAIQHSEFEEVTIDPT CSWRPVPIKSDLHIKDDPDGIPSKRF KTMSPSQMIMPNVMEMIAALGPGP SPYPLPPPPGGTNSNDYSSQGNNYQ GHGNFDFPHGNPGGTSMNDFMHG PPQLSHPPDMPNNMAALEKPLSHP MQETMPHAGSSDQPHPSIQQGLHV PHPSSQSGPPLHHSGAPPPPPSQPP RQPPQAAPSSHPHSDLTFNPSSALE GQAGAQGASDMPEPSLDLLPELTNP DELLSYLDPPDLPSNSNDDLLSLFE NN	True	False	4.355	5.0	1.707	1.068	0.531	1.556	1.486