

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q8NF64	ZMIZ2_HUMAN	Homo sapiens	Zinc finger MIZ domain-containing protein 2	17.733921	S135;T145;T159;T232	NaN	38665916;34019948;40136647;38253038;35254053	MNSMNPMPKALPPAPHGDGSFAYE SVPWQQSATQPAGSLSVTTVWGVG NATQSQVLGNPMPGAGSPSGSSMM PGVAGGSSALTSPOCLGQQAFAEGG ANKGYVQQGVYSRGGYPGAPGFTTG YAGGPGGLGLPSHAARPSTDFQAA AAAAVAAAAATATATATVAALQEK QSQELSOYGAMGAGQSFNSQFLQH GGPRGSPVAGMNPMTGIGVMGSPS GLSPLAMNPTRAAGMTPLYAGQRLP QHGYPGPPQAQPLPRQGVKRTYSEV YPGQQYLQGGQYAPSTAQFAPSPGQ PPAPSPSYPGHRLPLQQGMTQSLSV PGPTGLHYKPTQFNGQGASFNGGS VSYSQPGLSGPTRSIPGYSSPLPGN PTPPMTPSSVPYMSPNQEVKSPFL PDLKPNLNSLHSSPSGSGPCDELRL TFPVRDGVVLEPFRLQHNLAVSNHV FQLRDSVYKTLIMRPDLELQFKCYH HEDRQMNTNWPASVQVSNATPLT IERGDNKTSHKPLYLKHVCQPGRNT IQITVTACCSHLFVLQLVHRPSVRS VLQGLLKRLLPAEHCITKIKRNFSS GTIPGTPGPNGEDGVEQTAIKVSLKC PITFRRIQLPARGHDCRHIQCFDLES YLQLNCERGTWRCPVCNKTALLEGL EVDQYMLGILYIQNSDYEEITIDPTC SWKVPVKPDMHIKEEPDGPALKRC RTVSPAHLMPMSVMEMIALGPGAA PFAPLQPPSVPAPSDYPGQGSSFLGP GTFPEFPPPTTPSTPLAEFTPGPPPI SYQSDIPSSLLTSEKSTACLPSQMAP AGHLDPTHNPGLHTSNLGAAPP POLHHSNPPASRQSLGQASLGFTG ELAFSPATGVMGPPSMGAGEAPEP ALDLLPELTNPDELLSYLGPDLPTN NNDDLLESLFENN	True	False	0.281	5.0	4.337	False	False	1.96