	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	go ap
0	O60293	ZC3H1_HUMAN	Homo sapiens	Zinc finger C3H1 domain-containing protein	12.331815	S736;T744;S749	S15;S28;S34;S128;S251;S352;S383;S662;S714;S717;S719;T766;S805;S809;S948;S949;S953;S998;S1046;S1301;S1303;S1304	30379171;36240223;37217939;34019948	MATADTPAPASSGLSPKEEGELEDG EISDDDNNSQIRSRSSSSSSGGGLLP YPRRRPPHSARGGGSGGGGGSSSSS SSSQQQLRNFSRSRHASERGHLRGP SSYRPKEPFRSHPPSVRMPSSSLSES SPRPSFWERSHLALDRFRFRGRPYR GGSRWSRGRGVGERGGKPGCRPPL GGGAGSGFSSSQSWREPSPPRKSSK SFGRSPSRKQNYSSKNENCVEETFE DLLLKYKQIQLELECINKDEKLALSS KEENVQEDPKTLNFEDQTSTDNVSI TKDSSKEVAPEEKTQVKTFQAFELKP LRQKLTLPGDKNRLKKVKDGAKPLS LKSDTTDSSQGLQDKEQNLTRRISTS DILSEKKLGEDEEELSELQLRLLALQ SASKKWQQKEQQVMKESKEKLTKT KTVQQKVKTSTKTHSAKKVSTTAKQ ALRKQQTKAWKKLQQQKEQERQKE EDQRKQAEEEERRKREEEIRKIRDLS NQEEQYNRFMKLVGGKRRSRSKSS DPDLRRSLDKQPTDSGGGIYQYDNY EEVAMDTDSETSSPAPSPVQPPFFS ECSLGYFSPAPSLSLPPPPQVSSLPPL SQPYVEGLCVSLEPLPPLPPLPPLPP EDPEQPPKPPFADEEEEEEMLLREE LLKSLANKRAFKPEETSSNSDPPSPP VLNNSHPVPRSNLSIVSINTVSQPRI QNPKFHRGPRLPRTVISLPKHKSVV VTLNDSDDSESDGEASKSTNSVFGG LESMIKEARRTAEQASKPKVPPKSEK ENDPLRTPEALPEEKKIEYRLLKEEI ANREKQRLIKSDQLKTSSSSPANSD VEIDGIGRIAMVTKQVTDAESKLKKH RILLMKDESVLKNLVQQEAKKKESV RNAEAKITKLTEQLQATEKILNVNR MFLKKLQEQIHRVQQRVTIKKALTL KYGEELARAKAVASKEIGKRKLEQD RFGPNKMMRLDSSPVSSPRKHSAE LIAMEKRRLQKLEYEYALKIQKLKEA RALKAKEQQNISPVVEEEPEFSLPQP SLHDLTQDKLTLDTEENDVDDEILS GSSRERRRSFLESNYFTKPNLKHTD TANKECINKLNKNTVEKPELFLGLKI GELQKLYSKADSLKQLILKTTTGITE KVLHGQEISVDVDFVTAQSKTMEVK PCPFRPYHSPLLVFKSYRFSPYYRTK EKLPLSSVSYSNMIEPDQCFCRFDLT GTCNDDDCQWQHIQDYTLSRKQLF QDILSYNLSLIGCAETSTNEEITASAE KYVEKLFGVNKDRMSMDQMAVLLV SNINESKGHTPPFTTYKDKRKWKPK FWRKPISDNSFSSDEEQSTGPIKYAF QPENQINVPALDTVVTPDDVRYFTN ETDDIANLEASVLENPSHVQLWLKL AYKYLNQNEGECSESLDSALNVLAR ALENNKDNPEIWCHYLRLFSKRGTK DEVQEMCETAVEYAPDYQSFWTFL HLESTFEEKDYVCERMLEFLMGAAK QETSNILSFQLLEALLFRVQLHIFTG RCQSALAILQNALKSANDGIVAEYLK TSDRCLAWLAYIHLIEFNILPSKFYDP SNDNPSRIVNTESFVMPWQAVQDV KTNPDMLLAVFEDAVKACTDESLAV EERIEACLPLYTNMIALHQLLERYEA AMELCKSLLESCPINCQLLEALVALY LQTNQHDKARAVWLTAFEKNPQNA EVFYHMCKFFILQNRGDNLLPFLRK FIASFFKPGFEKYNNLDLFRYLLNIP GPIDIPSRLCKGNFDDDMFNHQVPY LWLIYCLCHPLQSSIKETVEAYEAAL GVAMRCDIVQKIWMDYLVFANNRA AGSRNKVQEFKFFTDLVNRCLVTVP ARYPIPFSSADYWSNYEFHNRVIFFY LSCVPKTQHSKTLERFCSVMPANSG LALRLLQHEWEESNVQILKLQAKMF TYNIPTCLATWKIAIAAEIVLKGQREV HRLYQRALQKLPLCASLWKDQLLFE ASEGGKTDNLRKLVSKCQEIGVSLN ELLNLNSNKTESKNH	None	None	None	None	None	None	N