	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	extracellu region
0	Q9C0A1	ZFHX2_HUMAN	Homo sapiens	Zinc finger homeobox protein 2	20.509114	S1490	NaN	37217939;39894887;30379171	MATLNSASTTGTTPSPGHNAPSLPS DTFSSSTPSDPVTKDPPAASSTSEN MRSSEPGGQLLESGCGLVPPKEIGE PQEGPDCGHFPPNDPGVEKDKEQE EEEEGLPPMDLSNHLFFTAGGEAYL VAKLSLPGGSELLLPKGFPWGEAGIK EEPSLPFLAYPPPSHLTALHIQHGFD PIQGFSSSDQILSHDTSAPSPAACEE RHGAFWSYQLAPNPPGDPKDGPMG NSGGNHVAVFWLCLLCRLGFSKPQ AFMDHTQSHGVKLTPAQYQGLSGS PAVLQEGDEGCKALISFLEPKLPARP SSDIPLDNSSTVNMEANVAQTEDGP PEAEVQALILLDEEVMALSPPSPPTA TWDPSPTQAKESPVAAGEAGPDWFP EGQEEDGGLCPPLNQSSPTSKEGGT LPAPVGSPEDPSDPPQPYRLADDYTP APAAFQGLSLSSHMSLLHSRNSCKT LKCPKCNWHYKYQQTLDVHMREKH PESNSHCSYCSAGGAHPRLARGESY NCGYKPYRCDVCNYSTTTKGNLSIH MQSDKHLANLQGFQAGPGGQGSPP EASLPPSAGDKEPKTKSSWQCKVCS YETNISRNLRIHMTSEKHMQNVLM LHQGLPLGLPPGLMGPGPPPPPGAT PTSPPELFQYFGPQALGQPQTPLAGP GLRPDKPLEAQLLLNGFHHVGAPAR KFPTSAPGSLSPDAHLPPSQLLGSSS DSLPTSPPPDDSLSLKVFRCLVCQAF STDSLELLLYHCSIGRSLPEAEWKEV AGDTHRCKLCCYGTQLKANFQLHLK TDKHAQKYQLAAHLREGGGAMGTP SPASLGDGAPYGSVSPLHLRCNICDF ESNSKEKMQLHARGAAHEENSQIY KFLLDMEGAEAGAELGLYHCLLCA WETPSRLAVLQHLRTPAHRDAQAQR RLQLLQNGPTTEEGLAALQSILSFSH GQLRTPGKAPVTPLAEPPTPEKDAQ NKTEQLASEETENKTGPSRDSANQT TVYCCPYCSFLSPESSQVRAHTLSQ HAVQPKYRCPLCQEQLVGRPALHFH LSHLHNVVPECVEKLLLVATTVEMT FTTKVLSAPTLSPLDNGQEPPTHGPE PTPSRDQAAEGPNLTPEASPDPLPEP PLASVEVPDKPSGSPGQPPSPAPSPV PEPDAQAEDVAPPPTMAEEEEGTTG ELRSAEPAPADSRHPLTYRKTTNFAL DKFLDPARPYKCTVCKESFTQKNILL VHYNSVSHLHKMKKAAIDPSAPARG EAGAPPTTTAATDKPFKCTVCRVSYN QSSTLEIHMRSVLHQTRSRGTKTDS KIEGPERSQEEPKEGETEGEVGTEK KGPDTSGFISGLPFLSPPPPPLDLHR FPAPLFTPPVLPPFPLVPESLLKLQQ QQLLLPFYLHDLKVGPKLTLAGPAPV LSLPAATPPPPPQPPKAELAEREWER PPMAKEGNEAGPSSPPDPLPNEAAR TAAKALLENFGFELVIQYNEGKQAV PPPPTPPPPEALGGGDKLACGACGK LFSNMLILKTHEEHVHRRFLPFEAL SRYAAQFRKSYDSLYPPLAEPPKPPD GSLDSPVPHLGPPFLVPEPEAGGTRA PEERSRAGGHWPIEEEESSRGNLPP LVPAGRRFSRTKFTEFQTQALQSFFE TSAYPKDGEVERLASLLGLASRVVVV WFQNARQKARKNACEGGSMPTGG GTGGASGCRRCHATFSCVFELVRHL KKCYDDQTLEEEEEEAERGEEEEEV EEEEVEEEQGLEPPAGPEGPLPEPP DGEELSQAEATKAGGKEPEEKATPS PSPAHTCDQCAISFSSQDLLTSHRRL HFLPSLQPSAPPQLLDLPLLVFGERN PLVAATSPMPGPPLKRKHEDGSLSP TGSEAGGGGEGEPPRDKRLRTTILPE QLEILYRWYMQDSNPTRKMLDCISE EVGLKKRVVQVWFQNTRARERKGQ FRSTPGGVPSPAVKPPATATPASLPK FNLLLGKVDDGTGREAPKREAPAFP YPTATLASGPQPFLPPGKEATTPTPE PPLPLLPPPPPSEEEGPEEPPKASPE	True	False	False	5.0	False	False	False	False	False

									SEACSLSAGDLSDSSASSLAEPESPG AGGTSGGPGGGTGVPDGMGQRRYR TQMSSLQLKIMKACYEAYRTPTMQE CEVLGEEIGLPKRVIQVWFQNARAK EKKAKLQGTAAGSTGGSSEGLLAAQ RTDCPYCDVKYDFYVSCRGHLFSRQ HLAKLKEAVRAQLKSESKCYDLAPA PEAPPALKAPPATTPASMPLGAAPTL PRLAPVLLSGPALAQPPLGNLAPFNS GPAASSGLLGLATSVLPTTTVVQTAG PGRPLPQRPMPDQTNTSTAGTTDPV PGPPTEPLGDKVSSERKPVAGPTSSS NDALKNLKALKTTVPALLGGQFLPF PLPPAGGTAPPAVFGPQLQGAYFQQ LYGMKKGLFPMNPMIPQTLIGLLPN ALLQPPPQPPEPTATAPPKPPELPAP GEGEAGEVDELLTGSTGISTVDVTH RYLCRQCKMAFDGEAPATAHQRSF CFFGRGSGGSMPPPLRVPICTYHCL ACEVLLSGREALASHLRSSAHRRKA APPQGGPPISITNAATAASAAVAFAK EEARLPHTDSNPKTTTTSTLLAL									