

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	ex re
Q8C0C0	ZHX2_MOUSE	Mus musculus	Zinc fingers and homeoboxes protein 2	36.96114	T150;T265	S824;S826	40997131;40885482;40021952;36852467	MASKRKSTTPCMVRTSQVLEQDML EEADRAKDKGAGMPQSDVTKDSWA AEPEHSSKETEVVEVKSMDGENLSKK LQGGYECKYCPYSTQNLNEFTEHVD MQHPNVILNPLYVCAECNFTTKKYD SLSDHNSKFHPGETNFKLKLKIRNN QTVLEQSIENHVVVITASGPGSSD NDPGVSVGKTPMKTGTGKLDADAKV PKKPDEAAPENHMEGTARLVTDTAE ILARLGSVELLQDSLGHVMPVQLP PNINLVPKVPVPLNTTKYNSALDTNA TMINSFNKFPYPTQAELSWLTAAASK HPEEHIRIWFATQRLKHGISWSPEE VEEARKKMFNGTIQSVPTITVLPQA LTPTKVSPILQALPCQILGQPSLVL TQVTSGSTTVSCSPITLAVAGVTNHG QKRPLVTPQAAPEPKRPHIAQVPEPP PKVANTPLTPASDRKKTQLQIAHLKA SFLQSQFPDDAEVYRLIEVTGLARSE IKKWFSDDHRYRCQRGIVHITSESLAK DQMAITGTRHGRTYHVYPDFAPQKF KEKSQQQLKLEDSFLKSSFPQTQAE VERLRVETKLSRREIDSWFSERRKL RDSMEQAVLDSMGSGKKGSDAVAP NGALSRLDQLSGAQLAGSLPSPSSAI VQNQEQVHLLRSTFARTQWPTPQEY DQLAAKTGLVRTEIVRWFKENRCLL KTGTLVSWLEQYQRHHMSDDRRDA VSRKVAKQVAESPKNGSEAAHQYAK DPKALSEEDSEKLVPRMKVGGDPTK DCLAGKPSEATSDRSEGRDGGQSE ENEESGIVDFVEVTVGEEDAISEKW GSWSRRVAEGTVERADSDSDSTPAE AGQA	True	False	3.452	5.0	2.624	2.095	1.737	2.706	2.1