

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6NXX2	ZN532_MOUSE	Mus musculus	Zinc finger protein 532	31.870043	S433;T443;S455;T478;S481	S130;S133;S134;S306;S313;S433;S875	22645316;22517741;36852467	MTMGDMKTPDFDLDLAAFDIPDMV DPKAAIESGHHESHKQNAHVDD DSHTPSSSDVGVSVIVKNVRNIDSSE GVEKDGHNPTGNLHNGFLTASSL DSYGKDGAKSLKGDTPASEVTLKDP AFSQFSPISSAEEFEDDEKIEVDDPP DKEEARAGFRSNVLTGSAPQQDFDK LKALGGENSSKTGVSTSGHTDKNKV KREAESNSITLSVYEPFKVRKAEDKL KENSEKMLESRVLDGKPSSEKSDSG IAAAASSKTKPSSKLSSCIAAIALSA KKAASDSCKEPVANSREASPLPKEV NDSPKAADKSPESQNLIDGTTKASL KPSDSPRSVSENSSKGPSSPVGST PAIPKVRITIKTSSGEIKRTVTRVLP VDLDSGKKPSEQAASVMASVTSLLS SSASATVLSPPRAPLQTAMVTSAVS SAELTPKQVTIKPVATAFLPVSAVKT AGSQVINLKLANNNTTVKATVISAASV QSASSAIKAANAIQQQTVVVPASSLA NAKLVPKTVHLANLNLPPQGAQATS ELRQVLTQPQQIKQAIINAAASQPP KKVSRVQVSSSLQSSVVEAFNKVLS SVNPVPVYTPNLSPPANAGITLPMRG YKLECGDAFALEKLSQHYDRRSV RIEVTCHCTKNLVFYNKCSLLSHA RGHKEKGVMQCSHLILKVPADQ MIVPPSSNTAASTLQSSVGAATHVTP KVQPGIAGAVISAPASTPMSAMPPLD EDPSKLCRHSLKCLECNEVFQDEPS LATHFQHAADTSGQOMKKHPCRQC DKSFSSSHSLCRHNRIKHGIRKVVY ACSHCPDSRRTFTKRLMLERHIQLM HGKDPDVKELSDDAGDVTNDEEEE AEIKEDAKVSPKRKLEEPVLEFRPP RGAITQPLKCLKINVFVHKCAVCGF TTENLLQFHEHIPQHRSDGSSHQCR ECGLCYTSHGSLARHLFIVHKLKEP QPVSKQNGAGEDSQQENKPSPEDE AAEGAASDRKCKVCAKTFETEALN THMRTHGMFAIKSKRMSSAEK

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