

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	extracellular region
Q9Z2U2	ZN292_MOUSE	Mus musculus	Zinc finger protein 292	24.992371	T1043;S1624	S654;S1146	36852467	MADDEAEQERLSSGGCAELRRRLG ERLQELERRLCESREPAVEAAAAAYC RQLCQTLLEYAEKWKTSEDPLPLLE VYTVAIQSYVKARPLYTSECESVALVL ERLALSCVELLLCLPVELSDKQWEQ FQTLVQVAHETLMESGSCSELQFLAT LAQETGVWKNVAVLSTILSQEPLDKE KVNEFLAFEGPILLDMRIKHLIKTNQ LSQATALAKLCSHDHPEIGTKGSFKQT YLVCLCTSSPSEKLEIEISEVDCCKDAL EMICNLESEGDEKSALVLCFTAFLSR QLQQGDMYCAWELTLFWSKLQQRV EPSVQVYLERCRLSLLTKTVYHIFV LIKVINSETEGAGLATCIELCVKALRL ESTENTEVKISICKTISCLLPEDLEVK RACQLSEFLIEPTVDAYYAVEMLYNQ PDQKYDEENLPIPNLSRCELLLVKLT QWPFDPFWDWKTLLKROCLALMGE EASIVSSIDELNDSEVYEKVDYQGER GDTSVNGLSAAGLGTDSGLLMDTG DEKQKKKEIKELKDRGFISARFRNW QAYMQYCLLCDKEFLGHRIVRHAQK HYKDGISCPICAKNFNSKDSFVPHV TLHVQSSKERLAAMKPLRRLGRPP KITATHENQKTNINTVAKQQRPIKK NSLYSTDVIFVFNNDGSDDENDDKD KSYEPVIVPVQKVPVNEFNCPVTFC KKGFKYFNLIHVKGHKDESEDAKR FLEMQSKKVICQYRRHFVSVTHLN DHLQMHCGSKPYICIMCKKAGFN SYAELLAHRKEHQVFRAKCLFPKCG RIFSQAALLYDHEAQHYNTYTCKFTG CGKVYRSQSEMEXKHQDGHSHPETG LPPEDQLQPSGNDVNPDSGATAAGG RSENSIDKNLGSNRSADWEKNRAE PAVTKHGQISAAELRQANIPLSNGLE TRDNTTVLRTNEVAVSIKVSVNHGV EGDFGKQENLTMEGTGEPLITDVHK PGIGAGVQLCHPGFQEKKGHECLNE AQNLSNSELKMDLNPQSLERQ VNTLMTFSVQNEAGLEDNSQICKFE CGGDVKTSSSLYDLPLKTLSESITFVQ SQPDLSSPLGSPVPPKAPGQKFCSCQ VEGCTRTYNSSQSIGKHMKTAHPDQ YAAFQLQRKTKKGOKSNLNTPNH GKCVYFLPSQVSSNHAFFTPQTKA NGNPACSAQVQHVSPSIFPAHLASV SAPLLPSVESVLSNIPSDKHGQDQ ILCSQMENLSNAPLPAQMEDLTKTV LPLNIDSGSDPFLPLPTENSSLFSSP ADSENNSVFSQLENSTNHYPSTQTD GNINSSFLKGGSSSENGVFPQVSSA DDFSSTSAQPSTPKVKKDRGRGPN GKERKPKHNKRAKWPAILRDKGKFC SRCYRAFTNPRSLGGHLSKRSYCKP LDGAEIAQELLQTNRQPSLLASMILS TSAVNMQQPQQSNFNPETCFKDPSP FLQLLNVENRPTFLPSTFPRCDVSN FNASVSQEGSEIHKQALETAGIPSTFE SAEMLSOVVPIGSVSDAAQVSAAGM PGPPVTPLLQTVCHPNTSPSNQNT PNSKTLKECNLSLPLFTTNDLLLTIE NGLCSNFSSSSTEPQNFNTNSAHV SVISGPQNTRSSHLNKKGNSASKKR KKVAPAVSVSNTSQNVLPDLPVGL PAKNLPVPDNTNRSDMTPDCEPRAL VENLTQKLNNDHLFITDVKENCK ASLEPHTMLTPLTLKTENGDSRMM PLSSCTPVNSDLQISEDNVIQNFECT LEIIKTAMNSQILEVKSQGTGETT QNAQINYSMQLPSVNSIPDNKLPDA SQCSSFLTVMPKSEALHKEDQIQDI LEGLQNLKLENDTSAPASQSMLMN KSVALSPTTKSTPNIVQPVPEVIHV QLNDRVKNKPFVCQNGCNYSAMTK DALFKHYGKIHQYTPEMILEIKKNQL	None	None	None	None	None	None	None		

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CQLVHHFTIEEMVKKIKRPHYGRKS  
QSENLSQPNNQVKKQPSMAEETK  
TESQPAFKVPAATGDAALANATVIPE  
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TGQENISSKANQEKSKHRRTKPN  
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NDALAECTSKFVTQYPCMIKGCTSV  
VTSESNIRHYKCHKLSRAFTSQHR  
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RKGEKDEMDDELTELFITKLINEDST  
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ERQKPGNLKRVIKEKNTVQSKRRI  
DKTEPEVSLVVNNTRKEEPAVAVQ  
TTEHPASFDWSSFPMGFASFLK  
FLEESAVKQKNSDRDHSNLSGSKR  
GSHSSRRHVDKAAVAGSSHVCCK  
DSEIFVQFANPSKLQCSENVKVLDK  
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