

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
O88466	ZN106_MOUSE	Mus musculus	Zinc finger protein 106	24.992371	S1498	S657;S677;S876;S878;S881;S909;S953;T1036;S1040;S1041;S1046;S1291;S1293;S1296;S1313;S1339;S1381;T1383;S1474	36852467	MVRRERKCILCHIVYSGKKEMDEHMRSMLEHRELENLKKGRDISHECRVCRVTEVGLSAYAKHISGQLHKDNVDAQEREDDGKEEEEEEEYFDKELVLIQERKEQSRQDEPPSNQEVNSDDRQPQWRREDRIPYQDRESYSQPPRHRRGPPQRDWWKWEKDFNSTRKNSFPHSLRNSGGPRGSSVWHKATRCSSTWFLNHSNSGGGWHSSNNGMVDWNYNGTGRNSSWHSEGTGGFPSSWHMNNNSNGNWKSSVRGTNSWNYNGLGDKFQQGRNRNPNYQMEDMTKMWNKKSNNKPSKYSQERCKWQRDRDKAAKYRSPPEGYASDTFFPSEGLLEFNFEORESQTTKQDTAASKINGKNGTKARDKFRRWTPYPSQKTLDLQSALKEVIGSKSDTLEKPLFNFLITAGLRKPVDKTSNPPVIKTQKAGPPGSPSHKAISDGTAFCEVARACSITEQSEPHOKSNKIPLLLKSPLLPLPTPKSGPHKQNLKNSKKNKETSFPSPGDHSHLLNTSTLEGSHGSSYTSKSLGLCPRVLKENKTVSGTQKEPDEKLNNTSQAQDITVLQCPKTLQNPPLTPPKRMENDAKESSVEESAKDLSIESQPHSAGNSAMTSDAENHGIKSEGVASLTTEVVSCSTHTVDKEQGSQIPGTPENLSTSPRNSTVLQKEAEVQVSAATSPHSGLLLDLKTSLEDAQDNNLVKSDGPFETESFEDTSLDTELOKPDLDLNNQPPGTLPELSKLGFPASLQRDLRSRHSLSKSKTGTHLPEPNLNSARRIRNVSGHRKNETEKESGLKPTLRQILNASRRNVNWEQVIQVTKKQELGKGLPRFGIEMVPLVQNEQEVLDLDEEPLDSSLEGFQWEGVSIPSSSLARKRSLSESSVVMMDRAPVYSFFTGEGTGKENEAAQQSPSPNTALSAQSQKTAMYLEQEVAPLTPSVGTGERVGNIPQRRHSAQLPSGHIMPVMHSARDLHSQERSTPLSERHAQESTGEGNSLSSNASSGHAVSSLADAATDSSCTSGAEQTDGHSIRKRRATGDGSSPELPSLERKNNKRRKIKGKKERSQVDQLLTISLREEEELSKSLQCMDNKLQARAALQTAYVEVQRLLVLQKQITVMSALRTHRIQIQGLQETYPPEHPDQAPCSLISREQRNSRSQTSFETALLPAPFFPGFLDPPPSHASLSPGNPLQITMSTFQAHGTAPDSSVQIKQEPMSPEQEGNMNALPQGCASNVSKELLQTRRVVDDGSSVYPAIPAVIASSESTENCQEVSKDLNFSVEQGNRSRSGNSPSCQSPDLPGINRGEETAKGSSGSEACSSFLRLSFTPETPAEKETQSPADQPEQQAESTLASAETRGSKKKKLRKKKTLRATHVPENSDEQDVFTAKPARKVTKAKAAKGAQVTSQTGOEQGTARDEPDSSSLEVLEVTNPQLEVVAITDSESGDEKPDSPSKDAWIAAEQNPIETSRSGCDEVSSSELGTRYKDGVPVSVAEQTVISIKASKHSSEISSEPGDDEEPTGSEFGHQAAVNAIQIFGNFLYTCSADTTVRVYNLVSRRKCVGVFEGHTSKVNCLLVTHTSKSSVLYTGS SDHTTIRCYNIKTRCMEQLQLEDRVLCLEHNRWRTLYAGLANCTVVTFDIKNNKRQEIFECHGPRAVSCLATAQEGARKLLVVGSYDCTISVRDARNGLLRLTEGHGSKTVLCMKVVNDLVFSGSSDQSVHAHNIHTGELVRIYKGNHVAVTVNILGKVMVTAACDKFVRVYELQSHDRLOVYGGHKDMIMCMTHKSVIYTGCDYDGSIQAVRLNLMQNYRCWWYGCTLIFGVVDHLKQHLLTDHTNPNFQTLKCRWRNCDAFFTARKGSKQDVAGHIERHAEDDSKIDS	None	None	None	None	None	None	None		