

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
008808	DIAP1_MOUSE	Mus musculus	Protein diaphanous homolog 1	25.685855	NaN	S22;T751;T1104;S1237	39627609	MEPSGGGLGPGRGRTRDKKGRSPD ELPATGGDGGKHKFLERFTSMRIK KEKEKPNSAHRNSSASYGDDPTAQS LQDISDEQVLVLFQMLVDMNLNE EKQOPLREKDIVIKREMVSYLHTS KAGMNQKESRSSAMMYIQELRSGL RDMHLLSCLESRLVSLNPNVSWV QTFGAEGLASLLDILKRLHDEKEETS GNYDSRNQHEIIRCLKAFMNNKFGI KTMLETEEGILLVRAMPVAVPNM MIDAAKLLSALCILPOEDMNERVL EAMTERAEMDEVERFQPLLDGLKS GTSIALKVGCLQLINALITPAEELDFR VHIRSELMRLGLHQVLQELREIENE DMKVQLCVFDEQGDDEFFDLKGRLL DDIRMEMDDFGEVVFQIILNTVKDSK AEPHFLSILQHLLLRNDYEARPQYY KLIEECVSIQVLHKNGTDPDFKCRH LQIDIERLVDQMIDKTKVEKSEAKAT ELEKKLDSELTARHELQVEMKKME NDFEQKLQDLQGEKDALDSEKQIT AQKQDLEAEVSKLTGEVAKLSKELE DAKNEMASLSAVVAVPSVSSAAVP PAPPLPGDSGTVIPPPPPPLPGGVV PPSPPLPGGTCIPPPPLPGGACIPPP PQLPGSAAIPPPPLPGVASIPPPPL PGATAIPPPPLPGATAIPPPPLPGG TGIPPPPLPGSVGPPPPPLPGGPG LPPPPPPFGAPGIPPPPGMGVPPP PPFGFGVPAAPVLPFGLTPKKVYKPE VQLRRPNWSKFVAEDLSQDCFWTK VKEDRFENNELFAKLTAFSAQTKT SKAKDQEGGEEKSVQKKVKELK VLDSKTAQNLSIFLGSFRMPYQEIKN VILEVNEAVLTESMIQNLIKOMPEPE QLKMLSELKEEYDDLAESEQFGVV MGTVPRLRPRLNAILFKLQFSEQVE NIKPEIVSVTAACEELRSENFSLL ELTLLVGNMAGSRNAGAFGNIS FLCKLRDTSADQKMTLLHFLAELC ENDHPEVLKFPDELAHVEKASRVSA ENLQKSLDQMKKIADVERDVQNF PAATDEKDFVEKMTSFVKDAQEQY NKLRRMMHSNMETLYKELGDYFVFD PKKLSVEEFFMDLHFRNMFQAV KENQKRRETEKMRRAKLAKEKAE KERLEKQKREQLIDMNAEGDETG VMDSLLEALQSGAARRRKGRPROV NRKAGCAVTSLLASELTKDDAMAPG PVKVPKKEGVPTILEEAKELVGRAS	True	True	3.297	5.0	1.642	1.412	1.413	5.0	1.896