

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
H7BX01	H7BX01_MOUSE	Mus musculus	Dynamin-like 120 kDa protein, mitochondrial	22.380232	NaN	NaN	37507081	MWRAGRAAVACEVCQSLVKHSSGI QRNVPLQKLHLVSRSIYRSHHPALK LQRPQLRTPFQFSSLTHLSLHKLK LSPIKYGYQPRRNFWPARLAARLLKL RYIILGSAVGGGYTAKKTFDEWKDMI PDLSDYKWIVPDIWEIDEYIDLEKIR KALPSEDLASLAPDLDKITESLSLLK DFFTAGPKLVSEVLEVSEALLLGGSP GETAFRATDHGSESDKHVRKVS DKE KIDQLQEELLHTOLKYQRILERLEKE NKELRKLVLQKDDKGIHHRKLLKSL IDMYSEVLDVLSDYDASYNTQDHLP RVVVVGDQSAGKTSVLEMIQAARIFP RGS GEMMTRSPVKVTLSEGP HHVA LFK DSSREFDLTKEEDLAALRHEIEL RMRKNVKEGCTVSPETISLNVKGPC LQRMVLDLPGVINTVTSGMAPDTK ETIFSISKAYMQNPNAILLCIQDGSVD AERSIVTDLVSQMDPHGRRTIFVLT VDLAEKNVASPSRIQQIIEGKLFPMK ALGYFAVVTGKGNSSIESIAIREYEE EFFQNSKLLKTSMLKAHQVTTNRNLS LAVSDCFWKMVRESVEQQADSFKA TRFNLETEWKNNYPRLRELDNRNELF EKAKNEILDEVISLSQVTPKHWEEL QQSLWERVSTHVIENIYLPAAQTMN SGTFNTTVDIKLKQWTDKQLPNKAV EVAWETLQEEFSRFMTPEPKGEHD DIFDKLKEAVKEESIKRHKWNDFAE DSL RVIQHNALEDRSISDKQQWDA IYFMEEALQGRKDTENAIENMIGP DWKKRWMYWKNRTQEQC VHN NETK NELEKMLKVNDEHPAYLASDEITTV RKNLESRGVEVDP SLIKDTWHQVYR RHF LKTALNHCNLCRRGFYYQORHF IDSELECNVVLFWRIQRMLAITANT LRQQLTNTTEVRRLEKNVKEVLEDF A EDGEKKVLLTGKRVQLAEDLKKVR EIQEKLD AFIEALHQEK	None	None	None	None	None	None	None	None	

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