	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	E2QD73	E2QD73_DROME	Drosophila melanogaster	Phosphodiesterase	28.06426	S172	NaN	40245499	MSQESNGGPAAGGGAAAAPPPPPQY IITTPSEVDPDEVRSMADLELGSPEK QVQVQSQKFSSTSSTTKVATHSFSM SSSAGTTGQQSKQDSAQQIQQLQQL QQLQQLQQQQQQQQSQRIISSSTRS QSLQSSTIVGEATTITSGAAQILSASA AASLAQQLKAQSSTSIITSSEQRTSTS TSSSSSTRYIASGSSNLAGGNSNSAS SASSKTRFQSFLQQPEGAHGFLTAH QKHVRQFVRSTSAHSEAAAGVAGAR AEKCIRSASTQIDDASVAGVVESAGN LTDSSATGGSMQLSMSKLGLQQSSS ILISKSAETIEMKSSSAGMRTQLTLS GGFLAPPGNRKITILSPIHAPPGLHD MLKRAQGRSPLSPRISFPGSDSDLF GFDVENGQGARSPLEGGSPSAGLVL QNLPQRRESFLYRSDSDFEMSPKS MSRNSSIASERFKEQEASILVDRSH GEDLIVTPFAQILASLRSVRNNLLSL TNVPASNKSRRPNQSSSASRSGNPP GAPLSQGEEAYTRLATDTIEELDWCL DQLETIQTHRSVSDMASLKFKRMLN KELSHFSESSRSGNQISEYICSTFLD KQQEFDLPSLRVEDNPELVAANAAA GQQSAGQYARSRSPRGPPMSQISGV KRPLSHTNSFTGERLPTFGVETPRE NELGTLLGELDTWGIQIFSIGEFSVN RPLTCVAYTIFQSRELLTSLMIPPKTF LNFMSTLEDHYVKDNPFHNSLHAA DVTQSTNVLLNTPALEGVFTPLEVG GALFAACIHDVDHPGLTNQFLVNSS SELALMYNDESVLENHHLAVAFKLL QNQGCDIFCNMQKKQRQTLRKMVI DIVLSTDMSKHMSLLADLKTMVETK KVAGSGVLLLDNYTDRIQVLENLVH CADLSNPTKPLPLYKRWVALLMEEF FLQGDKERESGMDISPMCDRHNATI EKSQVGFIDYIVHPLWETWADLVHP DAQDILDTLEENRDYYQSMIPPSPPP SGVDENPQEDRIRFQVTLEESDQEN LAELEEGDESGGESTTTGTTGTTAAS ALSGAGGGGGGGGGMAPRTGGCQN QPQHGGM	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None