	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	Q2XY62	Q2XY62_DROME	Drosophila melanogaster	NaN	28.06426	T296	NaN	40245499	RLQLSRMPLDTWSKREQLILASAVS CSGDQNWITVSRTLKTICGNGSSNR PADWFSQKNCAMQYGNLLESVEAT KRKKRSSESSAGVSSPATVETPTELL LRRLTEERQAEIKAQMRQDQETYRR IQREIESLQSDAVTEQELQDMWLEIE KEQEARRIEEMKLENRMREREQRK KDLASNWRNSSLAVKRSNQAADTT SVDMDVEDINTSGNGKQQTGPSPLL TSLLKSPTGNAPATPTPGGAAPAAGG TGSVARATAPTITSLLTSGPSSAVAN QPVITMKSPTDAAFMSRPITGPPASE ALASTTPTLERSPSQAAPTLSMLLEK NKAAAKVNEGGSLVKPEGSGNQEQ GVDEDTSSAAGTADSDEADPNEAQL LEVFKNIDQIIDDIDIDMASVIDDEIL KNVDDVVASPSNDKAEVIDFEDKLD ALKREQSKSSPASDKPKSVELVIGSS DDSNDNIPLAAVASQESKDRGQSGG ESTRGTEDAESEPLTSEVAVEEIGETI TIISTSSEDTAGSPPTKMEEDGATDS LPKKEDAAEDKSEQESSSQGQNKAG GSTDEPELIDHQATREAQSKSGKDT AGKPVTMQEKTTTQSAPVPVDLHDT DEESSSLTDSSKHDDHPRSAKAKPP ALKLALDDSKSELELSGTPQPPSAPA RTPLLRKLPHRDRSESPMVDDDATT ASDHSTARSTRRRCSSTPVIDSIPNS PASSEHTDDRRETRAASKKLFLSIYA MLLDSKHAAPFKRPFHDEHAQRHA DLCLRPMDFPTIKRNIDSGFIRSLSE LHRDVLLMAHNVLVAYKPHTTQHK TARLFVQDCQAIKEFSQLPDVQAGIT ATPVSNTGSLRADKSSGSKARSGSR	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None