	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	A0A0C4DHC2	A0A0C4DHC2_DROME	Drosophila melanogaster	NaN	28.06426	S732	NaN	40245499	MPPHILGSASGSSSTAAASPPTSSAS QTPPSPAIHHSHLSQHNATSAMSKP TTPQRHGNPSLLPTLQWPWNTSLGS TSTAVPAASANKNKRTAAGSGAALG LSGEGSCLTGGGAASAKKARSDLPG SFDASKRLKVAAMEESQTKITGFFKS QMKPSPGGGKLSPQPGQQSNPANT LTMSTPATTASLNKYFNILSQLKEQK AQQQQQQQQASKTLPTPAPVAPVPS APPVAPVVVTPSPSLPVPALKKIERS NKQPAKIAQVAPNLRKTPSSGSSGSS SSSSSSSSNGSNTGKSPTKKHVAIAP RTPEMKQQQQQGKAAMVYRPPVTS TALKQKQISPPAPVAAHTPAPAPALN PTPAPANPTLYQLPVQLPNLVQLPPQ LAAAANIMQLNNVAKAAVAAAATNN AAAQAAQAAQAAAAQYFLNGTVFKL QQVTTATTTTATTATAAAAPAGNPFG LLNLATAGNPFGLPNANGQFPIEAIP GAGSYLHHQLLLARQNNMSLNETM ASNSCGNMNFVNPLNNQQVGLKD NKLYIVNSAEYLGYLMSLQIALNNQ QHQQQQIISATQPPPLAATNNNSTN TATQPPPLAATNNTNSNTANNSTAS NSINHNASNNLSNINNTAIQPQRAI VKPPLHSSTQPPPLVTISSMPACSPA AASSPAAKRSLPAAKPYQKRLTAKGR VGTAPIIATPTTPPPLVPTSATKELGQ LRKSTGTTGTPTGTPTPTPPLVSIAPS KLTPTLSVSKQGPTMKLANSAPDLF DLVKNSKLVAKVSQPLTPLPFSSPSS SSNGSHGSHGMRSSPALSTSNSCTL SAFSKIKVETTELASQTGSLTSSSIPTI SLKPQSFAGQLPKREPESETDTLKH DLLPDCTNSNSNSNSCSSSTYSHSV SSAADLSLEASTPAPSPSPSASPSGL GSPSPAASNLSASSRRAASQTDMLS ELVTSSCISSGGDDCSQATDSPPMP ALPLAKSEDATTPISTVSGGSSSGSS NYDEEDDKSVASLETHQTHKRLRDL PTPESGIGGSLSNSESSNSIADAISSK SASVGPPTTAASSASSSASDSNSLAS NAPSPASPEDCSAAPSPACSASTTGS IPPSTVVDIAMVEATSKSLPKSAISPIL SQPKTIRFPAGAGASGKGGKRHDGV CYWDKCNKKHESNSKLLDHMQTH HVNTQTGPFACLWVGCKVYNKESC SRRWLERHVLSHGGSKQFKCIVEGC GLRFGSQLALQKHVNNHFNATDNA RESTSKRTSDPPVPKQLPRMFDFFD TGIMEGLQHRLRQISTLTNGAQAIEF QGQCMMRRRNSQGGYECFVRWSP REIISDEWLPECLNRTRQHTKVLHIK QMRPAEKTRVDSLLSTAFRLRYDSH LFADDYNVNEQQVGEASGSDCDED GDGDQEEEDLEDQDEDEEEDGSSS SRSASCETVSSYQQVLSIAKLQMQQ RRKHPRKPPKVTPPAREKLVPTDALV PI	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None