	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	Q7KV90	Q7KV90_DROME	Drosophila melanogaster	non-specific serine/threonine protein kinase	28.06426	T786	NaN	40245499	MAHQQQQQLAPSVNCSLDDIDLTAL KDPAGIFELIEVVGNGTYGQVYKGR HTKTGQLAAIKVMDVTEDEEEEIKLE INVLKKYSNHRNIATYYGAFIKKSPP GKDDQLWLVMEYCGAGSVTDLVKS TKGQSLKEEWIAYICREILRGLSYLH SNKVIHRDIKGQNVLLTDNAEVKLV DFGVSAQLDRTIGRRNTFIGTPYWM APEVIACDENPDATYDNRSDLWSLG ITALEMAESQPPLCDLHPMRALFLIP RNSPPRLKSKKWSKKFHGFIDTVLV KDYHQRPYTENLLKHGFIKDQPTDR QVRIQLKDHIDRCKKRKQEKEREDY RYSGSDNDDDEPQLAGEHSSIVQAP GGDTLRRNFQQIQEGRLAAEQQQQ HHLMAQAQAQAAAAHAAAQAQAQL QQQQQQAAAAAAAAAHAAQQAQQA AQQAQAQQPQANRQPKPPSRQQVE EPGPPARPPQRLIVVPDPPHANRPLP PTPKCGEPAGQTPQQQQRNSQNNF KPSLPPRRPEPQPQTAAGGQGSQQQ AQPEAPPRNNRQSSGLSSSGGSASG GGGSSKPAAALPQQSNNHLGQPVN PLDPLDSSDSDSEPDEPNDRARNDG TLLASDPPKPLPGLGPVSEDANTTTP LSHGSGGPPNRPLPPTPDDDDQAGD RTLIMKRNRGGGGGGGGGGGGGGS SGVGADGTGLGTPGTRTSSVLPDLLS QASPATPPRHDKSSSEEYQAAISSSV HSTPSKSFIASSGSGGLGLGGGTVVG GVIISSNHSPQSTISLASSSSNSRQN SPKNSISKTRSSSSITNLLHKSASSSS ANLHHLTPCSSTSSASISNPLPPHAY ALQQKQRSFLTFGFGAGGSGPSRRE SHVNVNVTPTSHEAANDTPEIRKYK KRFNSEILCAALWGVNLLIGTENGL MLLDRSGQGKVYQLISRRRFQQME VLEGQNILVTISGKKNRVRVYYLSWL KSKILRTDGLSDQVERRNGWINVGD LQGAVHFKIVKYERIKFLVIALKDSIE IYAWAPKPYHKFMAFKNFGELEHRP LLVDLTIEDQSRLKVIYGSAEGFHAV DLDSAEVYDIYLPKHTQGAIIPHCIVA LPNSNGMQLLLCYDNEGVYVNTVG RVSKNIVLQWGEMPTSVAYIGTGQI MGWGNKAIEIRSVESGHLDGVFMH KKAQRLKFLCERNDKVFFSSAKGAS SCQIYFMTLNKPGMANW	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None