

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	extracel region		
Q7K4Q5	Y0417_DROME	Drosophila melanogaster	Probable protein phosphatase CG10417	27.146871	NaN	S289;S306;S592;S594;S599;T637;S639;S641	15952742	MGAYLSHPKTDKTSTDQFNELLAVG ASSMQGWRNSQEDAHNSILNFDN NTSFFAVYDGHGGAEVAQYCADKLP HFLKNLETYKNGQFEVALKEAFLGF DKTLDPISIVSILKILAGEHNFVDAE ADDYEEEDLAEHQEESNLPPLNEVLE KYGKLPQKDLDLKSSDHKENFKM RSPYFRGRRAAALAAEATNKAVMDP SAKPDGSSTSAAAAAALSADGVAN SRNPSNVVNPAGADSNTTTSINDL STKNAALKDDSVNDQNEGSNGTDF KHTLVSSSNKFLFATGSNDMTELN QSSKNEFTNSSTSKEFERNINSSQD DEFTDDADYEENDNVKSPDTSSAE SSDCTENDDGDEDGNEDSDEEET DEDQMANDNF CANMIEEPKDSGC TAVVCLLQGRDLYVANAGDSRCVIS RSGQAIEMSIDHKPEDDEEASRIKA GGRVTLDRVNGGLNLSRALGDHA YKTNVTLPAEEQMISALPDIKLIITP EDEFMVLACDGIWNYMSSEEVVEF VRCRLKDNKKLSTICEELFDNCLAP NTMGDGTGCDNMTAVIVQFKKKLQ ELQSTIPPNQTEDKLLKTSENVSHSL NDQSASKRCASQADADDEILEKN NSKRLKTDLEQENIKDRTPSPSNQN EDPTQKAIKEVTIIVSSS	None	None	None	None	None	None	None	None	None	None	None

0