	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	Q9W2N6	Q9W2N6_DROME	Drosophila melanogaster	U2 snRNP-associated SURP motif-containing protein	25.496522	NaN	NaN	33925313	MSKQGNKSKDSAAASATLKRISEKK LEAFTVGTFSKRQLSKKELEDQKKK EDAAAAAHAFKEFVETFQDAPTPSS KVWVKAGTYDAGSRREDKSEKGKLY KPVSKLMEKSSSDKVEDYAKTLASD LKKDSGPLKKKNQEKKKSNLELFKE ELRQIQEEREERHKYKHMASSHSAP AQQPAASAAPVPSSSVSTTSQNSSSS KESGSFDTGDPNTTNLYLGNLNPKI SEQQLMEIFGRYGPLASIKIMWPRSE EEKQRGRNCGFVAYMSRKDAERAL KTLNGRYIMGYEMRLGWGKTVPIM NTPIFTPQALLEMTLPPPPSGLPFNA QPPPSEADVLPKKNYKEFNQEDKEN MERILAKCVVKVHIPTEKAVLNVIHR MIEFVIREGPMFEALIMIREMENPLF AFLFDNESPAHIYYRWKLFSLLQGD TPNEWREEEFRMFKNGPVWKPPIA NFYTQGMPDELVVDPDAPVVHKGA LSNAQRNRLEDLIRHLTPERARIGDA MIFCIEHADAADEICECIAESLSNINT LASKKIARLYLISDILHNCTVKVANAS FFRKSVEKQLLDIFDNLHNYYLNIES RLKAEGFKSRVCNVIRTWEEWTIYP KDFMAQLTAKFLGKPYVKPVNNSPQ AEETRFDEALDEDIDGAPLSGEEKD DEDLDGVPLDGAALLKSALKRAIPDA DAGTPKRDTPKKDEYLDEIDGIPLDD DLDGVPLEKETKSTAKLPGFIPSKWE TVDPQQVEAQAITTSKWDTLDPPDP PKFFSSDDDSGEDNPQKYTDEMRQ KLRDIESKAIQYQDELESGKRRLEPG WSVPEQVEQFRKRLLKQDSRSEATV DSPLSYMHIKSKSSKVSESPALFRSS SRKRSHSPFSGGESKRRDYSPETSR SSKSSKRNRSRSPSSSPKRYTERSSR SSRYETPASGSSRSRSSPTSSRSSRR EEPSPPRHRSDKHKHKHRH	None	None	None	None	None	None	None	None	None