	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	A8WHE9	A8WHE9_DROME	Drosophila melanogaster	NaN	28.06426	S123	NaN	40245499	MTKTTKPKEKAAAGGAVIGSGSGLG SVTKAGGGSLLSNAADSKIRTAKSN NNKRQAGRAATALAATTTASALAAT TTAGATGSNAAANETEIAIETENGEA ATPTAAATAAAANLSSLESARSQALT SVVSETARQAVTTANASATSTSTVTA ATEIATATASDTAATSEAAIDDDPSAI NTNNNNNNSKAQNDASESVKTKVI SYHQSEDQQQQQQQQAHIYEQQQQ FLSQQLISHHQQEQHQQAQQQQHQ QVVQEQHQASWLAYDLTSGSAAAAA AAAAAASHPHLFGQFSYPPSHHTPT QLYEHYPSTDPIMRNNFAFYSVYTG GGGGVGVGMTSHEHLAAAAAAAAA VAQGTTPNIDEVIQDTLKDECFEDG HSTDYHVLTSVSDMHTLKDSSPYAL THEQLHQQQHHHQQQLHHHQQQQ QQLYHQQQQQQQQQQHHHHHNN STSSAGGDSPSSSHALSTLQSFTQLT SATQRDSLSPENDAYFAAAQLGSSL QNSSVYAGSLLTQTANGIQYGMQSP NQTQAHLQQQHHQQQQQQHQQH QQQQLQQQQQQHHHNQHQHHNS SSSSPGPAGLHHSSSSAATAAAVAAA TAAVNGHNSSLEDGYGSPRSSHSGG GGGGTLPAFQRIAYPNSGSVERYAPI TNYRGQNDTWFDPLSYATSSSGQAQ LGVGVGAGVVSNVIRNGRAISAANA AAAAAADGTTGRVDPGTFLSASASLS AMAAESGGDFYKPNSFNVGGGGRS KANTSGAASSYSCPGSNATSAATSA VASGTAATAATTLDEHVSRANSRRL SASKRAGLSCSNCHTTHTSLWRRNP AGEPVCNACGLYYKLHSVPRPLTMK KDTIQKRKRKPKGTKSEKSKSKSKN ALNAIMESGSLVTNCHNVGVVLDSS QMDVNDEISASSATEQQQQQRLHQ RPPVAGPTAAELHYGGSSSSSGGGS GGDQDGGLVAHLPGQLLHSVGGTY GHLNTDHGIEHIIVPESQQHY	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None