

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	extracellular region
Q9VH63	Q9VH63_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MGMVEASTSELHLEAPNSYTYLLAT VSLTLTFIVMLGACLCCCKRIPKNDL MGLEGMVKLKNPDEVIVVTNLVGN SESQAE LNASTVSNADSEKRSSTAA HRSLPDI PVAESNDNGSELYETVAD KQMVDGRNARSQSPQPSMKKQTSV SQHSSISQADDVSSPYSRVKGLPHD YAKVRGTEHPYAQLNAPSTSHAANH GSTAAAAAAAVVAGSSSEGIDPMHR SSQHSEGSREHNDSAPAQAEIPAAS AIAGMISASQDLPYMTPIANPHFSG DSQDSSKGYTSSISVREPLANILAQQP PGKPSNRNPTLSRDVND SHYATVSD DSETYAAIEDPNRRHHANAADIYT SGSETYAIQIPMQANPIVVAVEINNS SSLPTASTSAVRPTSSSGADHAMPVP PPVDSLRAQMHSRQASSSSNNSSSV CNLGS PKPEKRQANSPLPPTPKAIVG QAHHLQAGSISNLASGRNSVISVIEA GAEGHEAESGEASPQRKHSKSLSPS KEGEGNKNI EGM YAKVMKKH KFSR NSPSSQNSSPILARKQQDGLGVSP LDSAAALLVDAQGRVRSNSYSAKD HGYETIPADGQMMHG NRKSDCYAA NMLQKERQ QEGGAIAQPV TIATAIAI SASAKHYE TIINQ TQSPPPSRSNND PGYEQLQPHPVPSG EDEAKKSDYDP NYEVLK SRAHSD DGYAKVLDK KRPP VNADGSGYSTIPGADANHNYSILET KAAAETDHYARIAENAVAGTPGSSSS RLTPTSPSSTNSVSLNSSTVATSTPI SSQYESLTGTGSETDPNYESVCYLTT ASNTATTTATTNTSSGIEPGYEPLQ TDQESDTQTSSPRSGAAFTPSEQLLV DDYFHV	None	None	None	None	None	None	None		

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