

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
B7Z060	B7Z060_DROME	Drosophila melanogaster	NaN	27.156312	NaN	NaN	28604694	MNTSQTAAAGNRITFTSQPLPNGTISI AGNPGAVISTAQLPNTTITKTIQAGIG GQHQLQVHHVQQQSQSQQQQ QQQQTQSVGATQTQTLVKSNNHHA TLPAGLVSSAPAGIVTMTKTINQAGQ PLLNSMLPAGVVVGMRRQAPSQQQ QKNVPTNPLSRVVINSHMAGVRPQS PSITLSTLNTGQTPALLVKTDFNGFQL LRVGTGTPPTVTQTITNTSNNSTT STTNHPTTTQIRLQTVPAAAVSRYQQ QQHQQQSQSQSQSQQQHATST TASTSIALRKTIVRTSSTVPSSNNNN NSINATTNTTTTTTNNNNKPNATNQ QQQQKLQQQHLKAQQQKQQQ ASAAAAAAAAANVAATIKIKQNSMT NTTATSNIIVNSVASSGYANSSQPPH LTQLNAQAPQLPQITQIOTIPAQQSQ QQQVNVSSAGGTATAVSSTTAATT TQQGNTKEKCRKFLANLIELSTREP KPVEKNVRTLIQELVNANVEPEFC DRLERLLNASPOPCLIGFLKKSLLP RQALYTKELVIEGKPPPQHVLGLAG LSQQLPKIQAQIRPIGPSQTTTIGQTQ VRMITPNALGTPRPTIGHTTISKQPP NIRLPTAPRLVNTGGIRTQIPSLQVP GQANIVQIRGPHQAQLQRTGSVQIR ATTRPPNSVPTANKLTAVKVGQTQIK AITPSLHPPSLAAISGGPPPTLSVL STLNSASTTTLPIPSLPTVHLPEALR AREQMNSLNHNSNHFDKLVKIEIK APSLHPPHMERINASLTPIGAKTMA RPPPAINKAIGKKRDAMEMDAKLN TSSGGAASAANSFFQSSMSSMYG DDDINDVAAMGGVNLAEESQRILGC TENIGTQIRSKDEVFLNLPQLQARI RAITSEAGLDEPSQDVAVLISHACQE RLKNIVEKLAVIAEHRIDVIKLDPRYE PAKDVRGQIKFLEELDKAQKRHEE LEREMLLRAAKSRSRVEDPEQAKM KARAKEMQRAEMELRQRDANLTA LQAIGPRKKLKLDGETVSSGAGSSG GGVLSSSGSAPTTLRPRIKRVNLRD MLFYMEQEREFCSRSSMLFKTYLK	None	None	None	None	None	None	None		

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