

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P07663	PER_DROME	Drosophila melanogaster	Period circadian protein	34.053133	S942	NaN	30703153;2327476	MEGGESTESTHNTKVS DSAYSNSCS NSQSQRSGSSKSR LSGSHSSGSSGY GGKPSTQASSSDMI IKRNDKSRKK KKNKGAGQGAGQAQTLISASTSLEG RDEEKPRPSGTGCVEQQICRELQDQ QHGEDHSEPAIEQLQEEEEEDQS GSESEADRVEGVAKSEAAQSFPIPS LSVTIVPPSMGGCGGVGHAAGLDG LAKFDKTWEAGPGKLESM TGVGAA AAGTQGRGERVKEDSFCCVISMHD GIVLYTTPSITDVLGYP RDMWLGRSF IDFVHLKDRATFASQIT TGIPIAESRG SVPKDAKSTFCVMLRRYRGLKSGGF GVIGRPVSYEPFRLGLTFREAP EEAR PDNYMVSNGTNMLLVICATPIKSSY KVPDEILSOKSPKFAIRHTATGIISHV DSAAVSAALYLPQDLIGRSIMDFYH HEDLSVMKETYETVMKKGQTAGAS FCSKPYRFLIQNGCYV LLETWTSFV NPWSRKLEFVVGHHRVFQGPQKCN VFEEAAPTCLKISEEAQSRNTRIKEDI VKRLAETVSRPSD TVKQEVSRRCQA LASFMETLMDEVSRADL KLELPHEN ELTVSERDSVMLGEISPHHDYD SK SSTETPPSYNQLNYNELLRFFNSK PVTAPAELDPPKTEPPEPRGTCVSGA SGPMSPVHEGSGGSGSSGNFTTAS NIHMSSVINTSIAGTGGTGTGTGTG TGTGTGTGTGTGTGTGTGTGTGT GTGTGTGTGTGTGTGTGTGTGTGT SSSKGGTAAIPVTLTESLLNKHNDE MEKFMLKKHRESRGRTEKSKKSA NDTLKMLEYSGPGHG IKRGGSHSW EGEANKPKQQLTLGTDAIKGAAGSA GGAVGTGGVGGGAGVAGGGGSGT GVAGTPEGRATTTSGTGP GGAGGG GGAGAAAAAGASSVGSSTPGSSYP TCTQNINLWPPF SVGITPPVHSTHTA MAQSSFSSAGLPTFYYPASLTP TSP TRSPRMHKHPKGGTDMPTTSQQA AAAAAQAMPLOYMAGVMYPHPSLF YTHPAAAAATAMMYQPMPFPGMAN ALQIPERPLGSQ SAYNKSVYTTTPAS MTKKVPGAFHSVTTPAQVORPSSQS ASVKTEPGSSAAVSDPCKKEVPDSSP IPSVMGDYNDDPPCSSNPANNK KY TDSNGNSDDMDGSSFSFYSSFIKT TDGSESPDTEKDKPKHRK LKSMSTS ESKIMEHP EEDQTQHGDG	None	None	None	None	None	None	None		

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