	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	A0A0B4KHX4	A0A0B4KHX4_DROME	Drosophila melanogaster	NaN	28.06426	T1617	NaN	40245499	MTQKPGEWASALLARFEDQLPNRIG AYGTQARMSQDQLVACLIHISRYRFS LVISGLTKMLQRVNEAALQNRHEPE RCYFESLVIILTTLERCLTNQTKDTA RFEEAMNVKLLLREISQFVDVQSDS NPNAAQLKALASKVLFALSQNHFSA VFNRISARIQELTSCSEENPDYNDIE LIQHIDMDMIKLTKLLQETITKFRSK RAPPLILLYSLEKAIWNWIEYHPQEF QDLQRGTNRDISTCWEPLMDFVEYF KTENKKSKTLVWPLQMLLLILNPSC LEAVVNELQQSEKEKEKDKEKVASK SAQSTSRDKDFSAKQFIESIKRGLGQ HSPSKQVTESAAIACVKLCKASTYIN NTDSNNVVFKLVQFFINDLKALLFN PAKPFSRGQGYNFADIELMIDCWVS CFRINPHNIEALKVCLNLSSPQAYHF VIVCSLLRLAHIYVDFRLQNKNPFRI VNQPRLSWWPQTDVVHYRSAELRA LFTDTLNKATQGYIAHTPLRYITSLTL KSKDTQKGLTRAEEGPAHKMLLLLL VRLIHADPTLLLNTQGKVAHEVQSS TLELINGLVSLVHQTTMPDVAQEAM EALLALHAPEKIEVWNPEAPINTFW DVSSQVLFSISQKLIQHQIANYTDVL KWLREILICRNTFLQRHKDYAHVGS QIAICKQAHIKMEVVFFMYLWSVDL DAVLTSLSCFGLLCEEAEICCSSDEL TVGFIMPNYHIYQELAQLSTSATDSR ICFFDNTHGNVLSRLTLQKRIMTLLR KIEHCVHGVQPAWEETFRNWEVSS KVLQTYPKCKGEDGQAEVFHRGMG KRRASHQSSEHDLEEQINEWANMT WFLLALGGVCLHKRSSSRQMLLQQ SQNNASLGSLAQNSLYSSSTSSGHG SLHPSTVSLSTLPPAPPQDVSYCPVT QFVGQLLRLLVCSNEKIGLNIQKNV KELVGEEMSTQLYPILFDQVRAIVEK FFDQQGQVNVNVTDINTQFIEHTIYI MKSILDPKANKDPNNDQPSPSEHLG VTSIEGMMLGIVRYVRHLDMTVYAI RIKTKLCQLVEVMMKRRDDLAFRQE MKFRNKLVEYLTDWVMGTSHQIAP PSSADAAILTNTSLIFRDLDQACMEA VAALLRGLPLQPEESDRGDLMDAKS ALFLKYFTLFMNLLNDCIDSSEAEK EMNNTPLLPPRPRMAAGKLTALRN ATILAMSNLLGANIDSGLMHSIDLGY NPDLQTRAAFMEVLTQILQQGTEFD TLAETVLADRFEQLVQLVTMISDKG ELPIAMALANVVTTSQMDELARVLV TLFDAKHLLSPLLWNMFYREVEVSD CMQTLFRGNSLGSKIMAFCFKIYGA SYLQMLLEPLIRPLLDEEEETCFEVD PARLDPTEDIEQHRNNLIALTQKVFD AIINSSDRFPPQLRSMCHCLYQVLSK RFPNLLQNNIGAVGTVIFLRFINPAIV SPQELGIVDKQVHSSAKRGLMLMSK ILQNIANHVEFSKEQHMLCFNDFLR DHFEAGRRFFIQIASDCETVDQTSHS MSFISDANVLALHRLLWTHQEKIGD YLSSSRDHKAVGRRPFDKMATLLAY LGPPEHKPVDSHMMFSSYARWSSID MSSTNFEEIMVKHQMHEKEEFKTL KSMNIFYQAGTSKSGYPVFYYIARRY KIGETNGDLLIYHVILTLKPFCHSPFE VVIDFTHTCSDNRFRTEFLQKWFYV LPTVAYENVHAVYIYNCNSWVREYT KFHDRILAPLKGNRKLLFLESPNKLT DFIDAEQQKLPGATLSLDEDLKVFS NALKLSHKDTKVAIKVGPTALQITSA EKTKVLAHSVLLNDVYYASEIEEVCL VDDNQFTLSITNESGQLSFIHNDCD NIVQAIIHIRNRWELSQPDSVTVHQK IRPKDVPGTLLNMALLNLGSCDPNL RTAAYNLLCALTATFDLKIEGQLLET QGLCIPSNNTIFIKSVSEKLATNEPH	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None

									LTLEFLEESIQGFQRSTIELKHLCLEY MTPWLKNLVKFCKSNDDSKKLKVS QILDKLINLTIDQKEMYPSVQAKIWG SIGQIPELIDMVLDNFLHKSITYGLG SPQVEIMADTAVALASANVQLVSKK VITRICRVMDKSCTNPTQYLEQHMM WDDIAILGRYLLMLSFNNCLDVATS VPYLFHTITFLVCSGSLSMRASTHGL VINIIHSLCTCTNPSFSEEAQRVLRLS LDEFSLPKFYLLFGISKVKSAAVTAF RSSCRHPTDKWLGNERVTQPLPADR ERLSLPSLEVITDALLEIMEACMRDV PDCEWLNTWTSLARSFAFCYNPALQ PRALIVYGCISKSVTDHEVKQLLRILV KALESFNDLILIEALVMCLTRIQPLLR PESPIHRALFWVAISVLQLDEITLYGA GLALLEQNLHTLKSQGCFDKKETIA EVMMKTREKLEWHFKQLDHAVGLS FRSNFHFALVGHLIKGFRHPTPTTVS RTSRVLTMLLGIIAKPLHRDKFEVTP DSVAYLTALVAVSEEVRSRCHVKHA LPRWPADLSSSVENGEASGGVQAIG LPLSRRQKSWDILDQSALQFARQHK VPTLQERGSRSSVSNESNVLLDPEV LPDLSIQALVLTVLATLVKYSSDEGE TRVLYQYLAEGSVVFPKVFPVIHSLL DQKINNILSVSHDQVVLNSVQNIIQN MLASEDPSQQQLHFLQSCGFGGLW RFAGPFTKYNMMGESSELFVNCLE AMVETCLPGDESAPVPPSPRPYNLS SSLSSLTLGSPTDKAA								