

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
E1JHV7	E1JHV7_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MSGRGRKRGRPPKTPNERASGRF NYQLKKPKYLSEGKSPSTPSASRG ISPQSDGESSHNNHTNRSRGSAA KRGRGRKSAVQNTSSYSGRKGYES EYHYGSDFGDSEEDKSDNEDDMLL TPSDDESLEVANESESEFVCSFNQ NGVGRPPRPPSPEPVWLQEGROYAA LDLPDSSSEDLFIANTHVLRLSIYEV LRRFRHMVRLSPRFEDLCAALACE EQSALLTEVHIMLLKAILREDAQGT HFGPLDQKDTVNISLYLIDISITWPEV LRSYVESDKTFDRNVFHLSHTEYPY TGIDNRLEVLQFLSDQFLTNSIRDV MLOEGPIHYDDHCRVCHRIGDLLC CETCPAVYHLECVDPMPMNDVPTED WQCGLCRSHKVSQVVDVLPQEKQ GVLIRHDSLGVDRHGRKYWFIARRIF IEDQENFTCWYYSTTSKLLKLLSRLD AEELETRLHSQITERRDEIERQMKLT ETLTNEHKHTKRSVIEIEQEAKNELL EKEVLDEDEKDGDAKSESQSIEGTK KQEECKMVTQKSNQLTNGTLHFK LGMEQGFKNYVNVQYSTNPIALNKPQ RNEERDKRRRLSHKFSLLTASDFK WIGITMGTTDNMITTLRQTLINFESN IAASFLNINWVVKKIWNAAVMNAR RPSEFAVLLLLFQASLKSUVFANVW HEQLGHTTLQRITSAEREERKKLEK REKRERDDEEERNLAFNYKYTLG LKHQVWKQKGEYRVHGQGWGLW LSSRRRCGVRARRAQPLTHNRVYVH YTMGEENDVNEIILVDPRTQRFMQQ CESSNVDGQVCHYLPDQYKNVKVIE DVTEKIKGHIDVSKALNAPGRITYYSK VARKSRLDLDLDRRLKLAEEVEEQMA SKIPSDMKPLLSSQNNTANSKQTF LEKRLRLRTEVQAKGGPANVNLELV NSLAKQIQTVRLQFSQLNRFKVFPR CYTKECNTNSNAVSQITQNTCYSPL CLOKARAKKELLLLKKAHTAGNGS KETVAAILGAVKKPSILEQKLTEGKR ESTQVAVDDSEEGKPAESEAPLDDL QDWEHARAHAVPFSDSLLETCILVD QECVTNTKIQEVNASSGCNTTPDS NTQSDKIDYIESMDVCSNVEIESTE DSIVTGLNSGNAEDVDMTPGWRRK RNQSKSKSYIGTKDVLDTLDKDIPL NKQNRFPITARPVKRECVKKYERE TFENGNERVYSTSSPRGRVYLLNDA AKLYEQAVKTEDKSTITKKPSYSRYP LISNFLTHKKRSLLVLPREFELLKLA RLGGKSSSTNGFHHAAKNNTIWIYQ CSRPLFRFCWSYRTSNATSLSSLALQ LRILWSCLRWDMMIAKPPSTDGKHQ VTDTIEIVTLELLKLHSGRYGKETS YLRRKVVIPLEMPKTVREVTSSIRSGL RKRKRAESPQTEPQITEEWVDEDK LELWEIKFMGEKQEKARLSAVTRSV ASRQLEASGNSGNTSTNGALGVAG RVQLAPKLSDEVKEKMEQQLKLQRA VHQQRKLVATGEITRSVTPVKQVIG SRRVIVKPNPDGTTRIQQAQVTVSRT GGANTAAAAASPTVGGSTSTQSNPS TSTPHKVQIIRGPDGKVSVRGLNPGQ QLVQMPDQKLVLTSTSSSAGQG NKMKVPIKPASTSSSPAISQAQTTN PVTVPVQIAVKHVTKNSATQSIASSS RVALPLAQIKNLLLAQQQQSTSSS PATSSSPVQKIVSKVNTSTSGQTLQ QVQVQSGSKLVVQNAQQGQKVIIST AAQQQGTSPVQQQLVQSQPIQSP QQISMTQQQIVVGGQRILSPGQTIV TQRNVQSQALQMVQQQIQTTQQQQ QQHHVVQPQQQFVVQSNQIVQSSP SAQTKLVKQLVVQQSQQTIEEKTQI	None	None	None	None	None	None	None		

