	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	B7Z0D3	B7Z0D3_DROME	Drosophila melanogaster	Large proline-rich protein BAG6	33.312435	T719;S792	NaN	28604694;20146544	MLINLKVKTLDARIHEFSIDNELTIR QFKDQIAEKTNIAAENQRIIYQGRVL VDDKQVKEYDVDGKVLHVAERPPFS QRGANARNNDEPMRTFRNVARPPP PGMRTSPYFRALDGMLVGTMAIPVN NGPVAGTRPPPNRYPNSSSFCINRIT VALHMIDCADNIAAYLENPAVGLNN QSLDILQRGRWSMESTVVEVGVSST DLPRNNNIIDMVQDAVTAALSHTGA RNYTVVQLPTVYTNENGETSQQRTA EAITSEGAASGAASNASGETTAATVII EDVIETDDEVADGASDRSVTPTPEPE AEGAVGGQQVTAAETPTSSVGSAND AAAEGGNNSGPRRRTRPQVLAQVIQ HYRGVQARLAPFVDRYYEILQNDPT FEESDTDGRENAQRIFDRVSEAFHY LSHAQHAISDLMLDLSQPGPRVLTC RPILVEQSGYIRSNNIFTPNFLAPPSG IINEPFRNRAPGSASAAGTQTATTAG TQTPTSAGTPTAVAPSQAANLQAAT DASRSAAAMAEIASRAAGAAAEMAA GAASAAAGAANAAAAAARDLSSDQV EDPIDEPMVGPNAAGAETLAQEQQR LEMDHDHDQAQSETPERETRPVPR LRVYVPVTLPPPNPQLEMARIIQAMV NGQRPNDIHVEFNAPNVMSINLPV HVMTTVRQAPAAGSNETADQSASES SSPESAPASGNGESPNAASTSSGTR SGDQRANTLPTTATQTRSTSRPQIQI GGNNNWGGRIAPTHTAFDRFLPCN SHHIREPEQLLQNNSTSRSTSTAPA GGATVVPPTSAAVTRPVTTGRIQRGS NTIRPMWSSRRQRPASEQLSNLGPA AWAQAQTQTTHIPSAHVGAGSTNGA GVGAGRVRPVGGSTINRDRLATRTA HVGGGSTNGGLPSGRRGRLQAATSR LSRHLPTQLANFLINNMRNNAPTAS VGSAVAAVGDTAASPSGEAAPIASTT PAAPLQTVLPTSTPTPRGDSNNLRSQ LRTFLNDSLFVGVPINEQTIPGAIGR ALDWFGESLVYLPQYERPEYNSRDS VCNILRVSLRLIIELCNGAPAGVDSA QFEQSLKQICDQFRKRLYSVLFLCLG SANAELYWRQLMRLLCAPMRSNFR NEALQFLCIYIDPTIPAQTDTVDAQQ FLVLRSIQAAPPTAADEPFVAPPPFA FNPQQQQLQEQTLDTDVEMSEVTA SGSNSSPADELPAVIVGSEPWHMSF PNDWLPVITRDLQTQAEQSNRPQPP FSDAYISGMSAKRRKIIQSEKPTASV ECLIANGVQRAIQSVGLGGSNGGSA LNASISMDTVIGSIAHDSTIQAAYTD AVRNSLKERTERDVDFKTSKYPQIAK FTEQK	None	None	None	None	None	None	None	None	None