	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	B7FNR3	B7FNR3_DROME	Drosophila melanogaster	NaN	28.06426	T1047	NaN	40245499	MVNQSKVFMLHMENTGLREIEPYF LQSTGMYRLKISGNHLTEIPDDAFTG LERSLWELILPQNDLVEIPSKSLRHL QKLRHLDLGYNHITHIQHDSFRGLE DSLQTLILRENCISQLMSHSFSGLLI LETLDLSGNNLFEIDPNVFVDGMPR LTRLLLTDNILSEIPYDALGPLKSLRT LDISHNVIWSLSGNETYEIKASTKLN LDNLHLEYNHIEVLPPNSFKYFDTV NRTFFDGNPIHTLREDAFKPARIREI YMRYCGLTNISPVAFDSLVNSLQILD LSGNNLTKLHHKLFNNFDVLRVISM RDNKIKIQKPTETFNAVHYTLLKLDL SGDRNDPTNLQTLRNMTRMRNMR SLSISRLGSSSVGPEDFKDFGVELED LQITRASLSGIQSHAFKHVRGLKRLD FSENGISSIENDAFHEIGHSLISLKM SHGYSGSALPAEPLRHLTSLQELDFS NNHISSMSDTSFHFLKNLRLLELHD NRIEQVLKGTFQGDIHSKLEEISLRF NHLTSISQHTFFDLEALRKLHLDDN KIDKIERRAFMNLDELEYLSLRGNKI NNLADESFQNLPKLEILDMAFNQLP NFNFDYFDQVGTLSNLNVNVSHNQ IRQLMYNSSWSGRNEHGGMYHSNI KILDLSHNNISIIHPGYFRPAEISLTH LHLGYNSLMNTTRDVFGNMPHLQ WLDLSYNWIHELDFDAFKNTKQLQ LVFFGHNYLSDIPQDIFKPVQGLRIV DFSHNHLRGLPDNLFYNGGMEKLD VSHNMMLKIPSSSLSSLAALTLCEL HLSNNFISTIHSMDLSNKFRSLRYL DISYNYLLRIDDAVFATMPKLAVLDL SHNRDLKVMDKSFMGLENSLIKLG LENISLSTVPEIRLKYLREFRLGYNEL PSIPQELAHNMSNLRMLDLSNNDL TNVPLMTQALPHLRRLMLSGNPITS LNNNSFDGVNEDLEMLDISNFRLH YFEYGCLDSLPHLRSLKLTAYSHLEH FNIPHLLRHHYNIRQLWIEAPQPFTR IVKKGSGPTQEMQTLQLGKPTDLQR EMEGHLPSKLTNITFSGPQFTNLNE RILRGMRSPYLYMQLFNTSLQALPP NFFKYMGRVRNISLDIRYHNRNLKK IPNPNTGAVPYLPNSVFLTDLKMSH TDLNCDCDLGWVEFWQRKRRQYIC SSQTWTDTVFRTFMNSPCQVYGRH NCDEHDDDLRETRCENKGGQQLM EALKFDLECGWDNANCREAAFVVV MVCVAMVFWM	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None