

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
M9NG31	M9NG31_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MRLFSLPLLALLLVQAAGQSEVTS DDPATDAGSTTNTTDTKPRIPSQDE MDAFYMMFPAALGSLKLGWGLFPAY SILGAIPDNLQPTAAASKVVLVLADD ATAKTRVARQNPPNPLGQLMNWP ALPQDFQLPSMDLGPQVGSFLAQLP AMPTVPGLLGAAAPVPAPAPAAAAAP PPAPAPAADPPAAPVPDAPQAILGQ AALQNAFTFFNPANFDASSLLGQSV PTFAPPNLDVFAQMQRQFFPGMTP AQPAAGTDAQASDISEVRVRPEDPY SQEAQMKIKSALEMEQERQQQAQV KDQEQVPLLWFRMPTTQNDQATEE KTLEDLRVEAKLRAFERQVIAELRM LQKIELMAKQMRSSAAAQNGDSPYR ISYPLSRTPIHKITRADIEQALRDDYV RRLVNKEAQRARRRNSGINTOKANAL KRQAKSQDQTLSEKEDIVQIMAYAYR MANEQMESEKQKQKVYAAAYRTEQ NPMMMQQRQWSEEQAKIQQNQQQ IQQNPMMMQQRQWSEEQAKIQQN QQIQQNPMMMQQRQWSEEQAKI QQNQQQIQQNPMMMQQRQWSEE QAKIQQNQQIQQNPMMMVQQRQW SEEQAKIQQNQQIQQNPMMMQ ROWSEEQAKIQHDQMAQQMAQQ GLMMTEQRQRQWSEDAQIQQAQ QMAQQTMMMPQMQRQWTEDP QMVMQMQQRQWAEQDTRMQMAQ QNPMMQQRQMAENPOMMQQRQ WSEEQTKIEQAQQAQQNQMMMQ MQQRQWSEDAQIQQQQRQMM QQTMMMKERQWAEENPQSVQQQ GPMMMQQMPMSMMQREVEDEDN KAEDDLVGEAGQMPENEGTARHK VDALGVGGNKRKSKSKSAPPTVIN YAYAAPQRPVVQSYGTSYGGGGYGS NAYGVPRPVNSYQSQGYRAAVGNDE VDEMLRQHQT MARTINPKQPGEVG GSESQKSNPNPTTLTPAQEQPQE HRVHKRLAHFHRFGREAGLNATTS KGGCGRLDCLGRSCRCGRRGLE SRVSSRTSGTCQCKASHRNKRSVE YGTLETIDEGSLNELRREYKLGKEI TLPDEDPAEALMRYNAASIREALER ASMEPLEIGGDQYEEDAQQEPMEEE QLQHDPNTEPOYNHKDFVRLTTSTA SPITSTTEAATPTGSDSTSEATVTPEV TTTTSTTTTTTESTKDEGLDMQQD SQAEAESSHVTKSISKQEAIEIHLHS IVEELKNEILKLNLRCSIIISNNVAKE PVTEKNPPVVEEPSKQEDKPKVVEEK VIAEEQAPVEQEELEDEDEDSTSIST TTETSPSPGSYSTKPGLSLGSPRVDE QSGSSNKLDYDDNNWQRILANRG YDTDYLTKSHERQFAQQNLEMPK NCNYDNGSQEYGPYEPFQADEPST DTEGKAKRALSVKQQAQLLNAALND SGSDSSDGTTTTTTPSPYAMRGKRV RRRSTARRVPIPKIGKASDEVVWRSP RQAKMPQRPKKSMSPKPKQSSQVT TQATVSSTKLDLVDLKDVLRLQIQ KEKKSLLRTQSNLSTKPKSIKPV KVIKRKRLRRRHKSIIATIRSIQTK A	None	None	None	None	None	None	None		