	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	Q8MQW5	Q8MQW5_DROME	Drosophila melanogaster	NaN	28.06426	S81;T583;S588	NaN	40245499	PPPTSQESQQQQPQPPAHQVPPGAT PPPGIAMPKPHYQHDVQTPPLGRPF EPTGLMLKYGDPLAAKYGPPQDLKY PMPPVSQAGPADVKPYGGENLIKSS PYGPPPESPIDASARSTPGQDSQGSN SNSQPPSMPPQPQQFQSPHPSPHM PSPAGGGLPPGMHPQNLIHGPPPGA AGGSGPQPPPPPTSLHQPTPTSAGPP SLQHGLHPGHQHSQLSAASSIPPSSI GIPPTLSTMAPSHMHPHLHPHAHL QGLHRPHDLPPSMHPHAPMPLSLQ GHPQHGHGLPPSHTSQQQQQQQQ QQQPGGPAGTVRTPSPAQQPPRSM HDPQSSREPPTSQPSTTMAGSSGPG GPPPQQSPHAHRTSPLPGLAGSGPP PPGLIGHPMAIHPHLAHLPPGHPAH AALAHPGHHLLSHSIAGLGPGGGPIA LLAGPGGLGGIPESALSRRTPPSHLP HSHASSAPLTAHSVASMTSTSMSLT TSTVPSSAFSRASPSVQISSSGGGPS GPGSVGPGGMPNSSAAAAAAAAAH RAASPASSVSSLSRQSPLHPVPQSPL SHHPSSSALSAAAAAVAERDRHALM RQQSPHMTPPPVSNASLMASPLSK MYAPQPGQRGLGTSPPPHLRPGASP PVIRHPQMPLPLPLIAPGGGIPQIGV HPGQSPYPHPLLHPSVFYSPHHHPF NSPYGYAPYGPGFPAYMKPPPQPGQ LDPAAVMAAHHAGLQGPPPQQMRQ DEQNAAAAAAQAAAEKQHQAAAAA AAQQHKAPQQQQPGGMPPNKPPTP KTPQGPGGGMPPGMGGPGTPTGLP PGAYPGSHMPGYPQGPPHGSPFAPQ DGQPHGLKPTSHMDALRAHAHSAN SAGMGGGHHPTEPLPIDIEPDPEPEI PSPTHNIPRGPSPEAKPDDTECHRS QSAIFVRHIDRGDYNSCTRTDLIFKP VADSKLARKREERDRKLAEKERERR QQQQQQQQQQQQQQQAAAAQQAAQ QAKMKAELKPPYADTPALRQLSEYA RPHVAFSPVEQMVPYHHPMGPMYR ERELEEIKNAQAAAASQSRLDPHW MEYYRRGIHPSQFPLYANPAISQME RERLGIPPPHHVGLDPGEHMPQPPE AGFQLPPNVGQYPRPNMLIPREPHS DVLLRMSYADQLQYLQAAEFQRQSL HDQYFRQRPR	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None