	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	ext reg
0	Q9VMS2	Q9VMS2_DROME	Drosophila melanogaster	NaN	28.06426	S96;T224;T269;T1385;S1405;S1670	NaN	40245499	MPLKQLEPSLVRILVSLPWDAAQRL RQLANDGNPELRALNIQSVQFEGDS VINLKVGGQDIKITKDNVNETLEAAG SLSGNKSGLLATGFDGPSTSKAASAF MQQQQQRILQQPQNDAISLQQRRA AAAGLGLQKMPLHTTQQQQQQQQQ QQVAVTPAVFKSPNTVCPMEGKVPL LLPSPSATRDFPFESMRQARVLQGR EAGGLGPPLPPPPPPNVTLKVLKTQQ QQAPQNGELTPTTPTSAAPTPPGSKS QFIQPPPPPYPGLGAATASSVSSPIAI AGVSAKPAIVTAPSQLNHPLGHPPLP TATSTGSNNIAISSPLLVNLLQNDGN GMSNPNQLKSPQQQQQQSPLIGMS PSGAQMMNSPMRSSGPATPSDFLM GDVLSPVVPSSPTTPQAAAPPCPPPV VMRNLQQQQQQHQQAMLSPSGNA NHLYTQQQQQGPQQQQPQQQQQG PPHRFQQQHQQMSPQAVALRQQQL QRQQQMRFMQPQQQLQGQQQPQQ QFQPAPDCFNNMGMGPMQQQQIR HQLRPGGLPLAPPPPHPQQQQQQRL MGLSMPQASPQQQFMSGASPLGPA LSPASSHHSMHSPLMSNHQAQQQQ QPQMVSPAATPGPTAPSEMAGQLM PANQHVPAAPPPDYNQAAVNSRWP LAGINKPMDSATKSSFQEFTRYQMQ YNLQQQQQQVSLPGQPQQQQQQPQ QQQLPPPALPTQQQQQQQSHVQQP QVAGQTQGQVDSLISLSVLDTLTTN DLDALLPTLNCDLDSTLSLEDKNEL ESLLQDAKDLDLDLIEDNLSAVDMR DALNTLQEAPLEQQQQQQQQQAAM QPQQQTAQMMQMPYQQQPQQQQQ MQQQLMMQQQQRQQQLPQMQQR QQQHLQRQLQLQQQQQQQQQQQL QIVQRQQQQQQSQLQVQHPQQPQR PPQKQFIINPHTGDMEPMASDDSET EAEEETAQPQFRNSHSGLGSFNFNP ANDMLLPTNLFSEEDSNSAQQHPM GISSDQERSRDSLASNKSATSRARKT PVSKANHSNFSAGDNSPRSSNSSPL IGLNSPQSVNANAGGAPSKKAKPNL LRGKLQQGVKERKAKDPTAAPKQKR ERAKGNVKTKAAEEKIKLRLKLEKVE TTPAAGAGYEGQIIINQQQQQTIVVN NHMQQLPMQTQLLALPTQQQQQQ QQQQQQQQQQMQQPQMQISQQQH QQQQQTSQLQQQPAQLQQQQPQQQ QQQPIQQQQQHHPVYSNFQQQQQQ QQQQQQNVPGPNEPRVPPLQIRLRG KNHVVVKNTRKDRKKGQNQEASID ERQLKRSYSDQPPQQLLMDPAESKQ AKLTPPPHVNGLPILIQKTTATVASP QLPQQLQGGVATTTKTTTIVAGKNG LTISAIVEAHKAQAQAQAQSLNDPQL IVGSTNTGTTPTSTTLQQQQLSKIAT SLPSSITLSAINSNNNNNNVNSGNN NNRLLTMKNPIKTAATITPIVGGKPM TKNHKPPPYITAVQQLQLQKQQQQQ QQQQQQQQQQQQQQQQQRQLAAAV TMLPSATTLKRVEITKVTPQVTAEQA PSINPSPVVVAAAATATLTTSSTPTTQ TTQLICDRKLPVAMPPMTTVNNVVV ATVNSSSAALSTLSSVSTASSASIITT STSTSSPAALPSTLRNSPASNGSGTP GHGNNGGGEDSGIESMDALSEKSP HQLSSSSPIQVSKAAVTMVQQQQQQ TNAIVTITTSAAVIAPVIAPTSTQAGTT PTVAPSLQQQQKQQADDEIEKALAK MEGDFPDDLEDIVSSMIKETSECAS AAGGGFLNSLESSLPAATVATVVSTA STNTSSIGSIKLNGEHHILEHDDLIK KLTESKPPAIKVKLEEMPPLLTIKKEP VIKPMASGLKVEAKVEVKLEPKVEPK SEEKLQEKPPSESSSASLQPISIEIPA QVDGETPRIRTRASSRLESPLDAQKA SPDPTTVTGVSTTAATAANASTTVKS LSRASSTASPRVVTPTPNHNNNNNK RRRQESECGSSNDGADLGENSIKRP	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	Nor

									RISSGTTSNNNATETAAEHNAGAGL PKKVEIESSDSDEPLIEVAGKVRNSK QAQVEAADAAAVGVEKHVTRRNAQ QLQPPQNKTAGNHAPSSTPALGTPR SGKAQAPVGATSAANNNHNSSTAN NASVVPSSVTTTTMPGGAVVTTSQV NSGTSVVHATRGATAAAASTTNNNS ASSPTDEKIGTRRSVRASAAANKIIYS RSNAAAAAAAAAAAAEPKGTTGKAG AAVGANNAGNVESVAEARRKTRSAV IGESMLTEGRRRRTSRDYK								