	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	ex re
0	M9NED8	M9NED8_DROME	Drosophila melanogaster	NaN	28.06426	S821;T823;T824;T825;T826;T827;T931	NaN	40245499	MSADSPRRHPSGVVGSGIGLGSGSG TGLGSGSTGGSKSGAAVPAIVPPQTV SDRSILDSAIGFINDVTLANQPVQDP KDTITWARFETCADVSDPRFGDDWE LEGNAAPPLLLILGYGLGVQVWAIPA NGEAVEVLSWRHGVVTALRVLPTPA TAAALDENGRADEPVDSFAEKRPLV AFVDGGSAAASGLLAGSSGLGLGGG GGGVTTVGGSVGGVSGIGVSASAQF SAVNFMSLKTGVQVKTIKFKNAVLDI QANRSAVVITFHERIAVFDARTLEDR LTITTCYPSPGINPNPIALGPRWLAYA EHKLLHSKRSGGGCDGEGVPSYTAT VLNAAKSLSKGLREFGEQVAAGLTG TTAGSGASSKSSSFDSASGGPDAKQ SGVVTIIDVKHPVKDYSPTSGTPLSST AGSQGGGDPIVAHFVAHSEALVAME FDSSGMLLLTADRRGHDFHVFRVQ PHPVGPSLAAVHHLYVLHRGDTSAK VQHIAFSLDSRWAAVSTLRGTTHVF PITPYGGAMGVRTHTSLHVVNKLSR FHRSAGLGADGRSSSPISHSESTTFV QSLQPYHNPTLPPYPRPSVVQPLAQL RQPFTLGSPPGSAGLGGGVGGGGV MGSGVSAGGHGVGVGVGSNSQRQR QRLSSLSDDSGKPLSVCSIFAKSRSW LLEPPMATREQPHRVQRKAVDSLFV MAGHGALIQYDLDTKLASHVAKEKI CDDTPIELEVEARAQWNLGRRKDGS QEIAPPLGLDNWLIKDRHASLLLDSA NQFDDPDERTESWLAQVEIITHAGP HRRLWMGPQFVFKNYNTPSGSNLN HVDAEAVEIGVSKTTTTTLPSTAASS ALGLGTIIGKDRSSPLNMPLSAATSV GGAGIGSSAVTGRSGAGVPVLIESGS YSSIEQSPKLMDRFRHGHLDSDYGH GDTRLKEDLADAMRESPSTAAARRE TTGNYFTTDQLDGLALNNNNNINN NIIPTKDNASPNPNTNTNPNAIPSSN KVQKAEVTDAVDYPIRHSYGDHGEL STVVNIEVFDDQLSMSSISTNSRLSL EGPPSQSSPPLSLTNGLMDTNLMHF SESVTGTGVAQAQVHGRGANRFEV DDDDEEEEEEEELDEEAEPDDDER EDRPLGRRNL	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	No