	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	E9QAE1	E9QAE1_MOUSE	Mus musculus	NaN	25.875327	S1587;S1592	NaN	30059200	MEGKRQLDRRDFGKRLSLDSSLVEY MDSNKCIEHLLAQLREQHRSLWRE KLAVARLQREVAQRTSQGAMHEKLI HELEEERHLRLQSEERLREVTLESE RNRIQMLGLQQQFSRMEETVRSLL QSQGSPEQKKEEPAKITAYQEGLPD EERKEQAALGDLHVVAADEDSRSEH SSVEEGKEDSRLLLKRLKALEAENS ALALENENQREQYERCLDEVANQV VRALLTQKDLREECVKLKTRVFDLE QQNRALSLLFQQRVRPASDVLLQEH LQNAKSGTPALKHSGSGVVIPDHLC PRNSCSSGSELSLSSTCSEYSSGSSY TWQDRKLIGKRQSSRIWDKRLSIDS SLPSGFASPADELPPTRIKESHILEGL RKLQKRKVFLESPSLVISKWGYKDC MNSNEGIYSPGIKTSSLKECPPCQPS GMGIPCQNLHQTFIYDANSHEDAHE ELPSSALPLTSPSQGYRIQGSKLTHS VSDSLFNWEPHRKYFAEGTASVYTR EMPERLARCASHCPLERKLCPSIHL PWVQRDRDPHSQVHGHVPLNLQLS DTDDNETLNELHIESSDERSPSDLS LTGDTDKSMENLEVGFRKSEGGSR DMGEEACPTHLETSPKAFSFIKQQR VVKRTSSEECVTVIFDAENGEPVEFS SHQAGVVTVTRDEISISSALAGPSEE LTELSPQEITHFQAGATARDCPFLKR PEEETAKNIPKDENHILTAPNDFFSS QTIITQNSPRPKLAKPTQGVAWQSN SRPVGNMGVYTKQGLTKIPSRGKSS PQKSKLVESMPTMLVPSSALVTPEK TPASPMGKLSQCRKAESPGSLCDVQ PESHIPKHPAQVPHIPKISNKKNWV QCPRSQTSGIQSRHLMGRSDSSEM RSNGCVSAPQTRMRSPSPPTPPGRS VSLLVRPSYDYPPLPSPAKPETQGPG DAVSPGLKSPLLKGSSAPLVSSSPVM SEIQKKPSVASRKSAFPPALPSTQAV VHTRYPEHIASSPFAVTTSGPPKVSP KRGIPKPPPHHALGTAHMDTDLQTP KTCPSSCELLEVTSCKSLSPGRKRQL SDSTVMPHRPSFSGVNESLTPQVNS PSSSSSSCRNQGSSQGGQNSPEKGP KTRLPVGLKVLMKSPQLLKKTSLVP GKQEKDSLNEASRSCVAANTSNPED SKDPASLETSRDERTTALLGVRGPDS LARGLALDSTMPESLECSISRDDGR DGADSRAIKRSFSSSKPHLKPALGM NGAKARSQSFSIHLGEKPSTPPTEGP GKIRTQIITNTAERGNSLTRQNISME GSPSKTPSVSTSDNLPNAGRPLGNP FSRQSSLCSTGSSSSQHGSPSKLPLR MLTRPEGLLTPGVEDQRLSTQEECP ERGVHEQSSSDHHRHPSPLSDCPQ FLHVPGRMQRPSNLETSGISKLETS GRHPDASSTGTSVVSPEAPLSPSIEE KVMLCIQENVEKGQVQTKSASVEVK PRAGPSFTSWFGFRKSRLPALSSRR MEASKSKVEKKDTKLKSERKKEKKK AEVQSKIENELSRGTKKADGQSPDD GLQSSESLKVSQDIYNQMKLEPRNR PSPAVCPTKDAFMTELLNRVDKRAS VQRESGSKGVSCRSVLEGTSQGSCF ASGSVSTQGSQKKNIKTKVDMEKPR ESLGAEVNEDVQEDEEDRVADTTLQ SHMIESNCQMRTLDSGIGTFPLPDS GNRSVGRYICQQDSPEDTDALLPLQ PAASMASSARAQTLDREVPSSTDSQ SSPESMIIHSISDPSMMAREMRASQ SHLPKPTSSGKPSFEKQNGAESKAH TGPSFEYAEDAVASEPLSGWRGEDA TTETQEPKQVEETKEDPENRLSEISL ESFNKFSSNTVILLGKEKSPSEVEGQ KEEKGKKTEVSSSSSERPGVDRVES LSDSLYDSFSSCASQASNEA	None	None	None	None	None	None	None	None	None