	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	A0A178V9D9	A0A178V9D9_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MYLTLSDFISLLCGLGGMNEIRSDIL PDQNMGFSRRNQSPVSGFENGNLV DGFQPLSSRTWMRSPSSAENNPVR SRSNPNQLIHQEQTGNSSFPYAHEV AEIQEATRRKSSAVAPSDKPLGDDPI LSQHDSQRFSTSPPTSGTKSYTLSRS SDSQFPGQPSSVNSFNNARKTNSSP ATKRTRSPPVYPIEEDIPRNSFPSQD CTEGEEQARAKRLARFKGELEPIAD RPVDIQLTKSPVNKTMKPLDNKQTF NSLESSRDALKGDALPDYENSEQPS LIIGVCPDMCPESERGERERKGDLD HYERVDGDRNQTSKSLAVKKYTRTA EREAILIRPMPILQNTMEYLLSLLDR PYNENFLGMYNFLWDRMRAIRMDL RMQHIFNQEAITLLEQMIRLHIIAM HELCEYTKGEGFSEGFDAHLNIEQM NKTSVELFQMYDDHRKKGITVPTEK EFRGYYALLKLDKHPGYKVEPSELSL DLANMTPEIRQTSEVLFARNVARAC RTGNFIAFFRLARKASYLQACLMHA HFSKLRTQALASLHSGLQINQGLPV SDMSNWIGMEEEDIEALLEYHGFSI KVFEEPYMVKNDLFLHADKDYKTKC SKLVHMKKSRTIVEDVSAPTVVEDV STPFPLPSLITEATIGNQQCITAHKHE MPPARSLKKQTSMRLFDKEVADSKT SLLAEEDKPMGTFVMNPPGPFVINP VVHQEKQNDLTSAGGFHSPVKLYSP FGSPKFPQTKSSNLEKQPNDDRIGM SPGEIKFSIIGDVYTNHVPGPALQQS PKSMPMEIMPVTTIAECPTSVENKY ALEESVPEAAMICTLEKEFNDIDEED EDEDGVILNQYDEEVAKAKLKLIIRL WKRWSSRQSELRERRQLAAAAALN SLSLGTPIRFSKTDQSRACGEFNIDQ AMRRRFEEREKSWSRLNISDVIADIL VGRNPESKCISWKVVLCTQTKSVNS SSSASQVTHSAASRWLSSKLMPHAE HSSLNDDNLLFSAPGVSVWNKWVA NGSDIDFTCCLSVARDVEAENDMCE TTCGASAVLFLASGGLPLNLQREQL NLILESVPNGSVLPLLVVISSCNGEH MEPDTDIVSGLGLHDIDKSKIASFSI VSIANKSQKGQEVHFFNDSRLRDGF KWLASNSPLQPNLHHVKLRELFLTH FSFSLELLKQMPDQEVGPNICISAFN DALETSRRNITSAAEANPIGWPCPET MLLEDNRKECLMVKRYLPNLDWSS AENVELLSSVLENCKLPDFEDDLTW LTVGCASGAEIENHTQRLEGCLIEYL TQRSNLMGVSLATKETGVMLERNT RLELHNSSRYHITPRWIGIFQRIFNW RIMGLFDASSSSAYVLKSDLNMSTS SYADKFLAEDASYPSCPPNLPLLHE MIEISCSPLKSPPPYDDKAQRVVETG MLIDDHRDIEESMLEKNREACRGID LMITEDDELGERSWRSKGREAAEKK TIEKRESERLDELLEKCNMVQNSIA EKLCIYF	None	None	None	None	None	None	None	None	None