	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	A0A1I9LQ49	A0A1I9LQ49_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MDRNREARRVPMAAAGNGLSRRRH RAGSFRDSPEEEGPVELPEAARLRD RGGSNKKDRDRERDRDRERERERD RERDRLNSRSKRRRGERLMMVHGN LDDGGDDSSEESVNDDEEYDDGGV GPPSSLKMLPPTSNNISAASFSSSLS NHHNGGSGNLHHHHHSHNNNHQ RKNNFPPTKVFRSSPSPAPVSPLVST WKAADEMIGVSVPRKARSACTKRPH ESWASSTTGGGVFASGEQIHRQISST SPANRVSPASILASPSPPAPTSPSSSS ISVRKKLPSGTKQKPLPPKSSSSKLS SPVAVQDEIEIEIAEVLYGMMRMPST SKQEAAGNDLTEAAKSTVEVKSRVS SPISNPQTLPQSSITLAANSSSSNVS AIAPKRKKPRHVKYEDDNSSRVTTIK SEAEAPSKSQVPFSNQLKSSGSGEG NSSVLDSIIPLTRESNASLDSEKKEN NLSKDETILPKVESSSGFRSDGEGAK SSSPEKEKFEIDLMAPPPVRSSSERG GEMMECVAAEAKPKVTEVETQEAK PLLKEDRSDPAIHDSQEKKRPRMVA EAEHHKFERNCELKLDLDKSDHVG LVNKHHVQKPPPQQQLSVPDKTAQA SHLPLHMSMPGWPGGLPTMGYMA PTQGVVPTDTSSLSAAAMQPPPHLL FNQPRPKRCATHCYIARNIQSHQQF TKMNPFWPAAAGSAPMYGTKACNL SLMPPTELQGSVLGRSSNPVQDKNS QSTSKSSETAQRNQLMLQQALPPGA ANSILHGPTFIFPLGQQPHAAATIAA ASVRPPNSGITSSGPTATSTSMNGSA SATPAGAPTMSFSYPAMPGNETQYL AILQNNGYPFPVPAHVGAQPAYRGA PGQPMPFFNGSFYSSQMIQPPHHQ PQKQHQQQLTGQMLQSHAPNNQN GSASTGSSAAQKHLQNQQLRPPINH GNSQGFPTHKVQSQPLNFQQRQQP RENATQHSETVGEDSPSTADSRGSR SNVAYGQNYGMQMQPTNLGLMSSP APGGGVVGSSSSHGEKKSQQQVSKA GVESFQSPGYAMTFATFNGANTAPT LNMSSIAQNHAMFHSMPEAARQGY QMMAAQAAQQKMNYGASLEDGKS GSIGGAATANNTPEEQRKSGGGAIG KTSGGNGGQSIAFSNKQDLADASVS AVTSGSIVDSSSRLLNLGSALPQSSG SLPTSHHQQLLQQQQQQHMQRSQS QQPYTTMYLQKQQRYATSVAASAAR TKGPVVSNGSGFPDHNMTTSPAGTT KFANANSGFPQNLVQSSSNQVQSQ QWKNNSPRTTNTTQAQSPSMLSPS TSVAAASSLRNIPHKQQSRPQQSQIS FAANSKPMTSGSPMQQVQGGTNHQ APSPPMLVGSPSTSSVSKNASGSPRT TASASSAANKGGQASTTTHSASQPS KNLQPASAASSAGGRNNGPSVLGNP TTSSGSKSQQQQQLPKHGLQPQAQL FFSNPYMQAQHQHQQQQITISPSGG YYIQRHQQQSGSAPAVPVTGAVTATS DPAKAIAAASAANNMKGGGGMGKT QQHQLGPPGFTNVHAVSSAVQVKPV DQKQQAGE	None	None	None	None	None	None	None	None	None