	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	A0A1P8AQR8	A0A1P8AQR8_ARATH	Arabidopsis thaliana	5'-3' exoribonuclease	27.144759	NaN	NaN	28154133	MEAMANLVILDPAPATYDDVFKSMF EYIDHLFTLVRPRKILYLAIDGVAPRA KMNQQRSRRFRAAKDAAEAEAEEE RLRKDFEMEGQILSAKEKAETCDSN VITPGTPFMAILSVALQYYIQSRLNH NPGWRYVKVILSDSNVPGEGEHKIM SYIRLQRNLPGFDPNTRHCLYGLDA DLIMLSLATHEVHFSILREVITYPGQ QEKCFVCGQTGHFASDCPGKSGSN NAAADIPIHKKKYQFLNIWVLREYLQ YELAIPDPPFMINFERIIDDFVFLCFF VGNDFLPHMPTLEIREGAINLLMHV YRKEFTAMGGYLTDSGEVLLDRVEH FIQAVAVNEDKIFQKRTRIKQSMDN NEEMKQRSRRDPSEVPPEPIDDKIK LGEPGYKERYYAEKFSTTNPEETEQI KQDMVLKYVEGLCWVCRYYYQGVC SWQWFYPYHYAPFASDLKNLPDLEI TFFIGEPFKPFDQLMGTLPAASSNAL PGEYRKLMTDPSSPILKFYPADFELD MNGKRFAWQGIAKLPFIEEKLLLAA TRKLEETLTVEEQQRNSVMLDLLYV HPAHPLGQRILQYYHFYQHMPPHEC LPWMIDPNSSQGMNGFLWFSERNG FQTRVDSPVNGLPCIEQNRALNVTY LCPAKHSHISEPPRGAIIPDKILTSVD IKPFPPLWHEDNSNRRRQARDRPQV VGAIAGPSLGEAAHRLIKNTLNMKS STGAASGLIDPNGYYRNVPGNYSYG GVNRPRAPGPSPYRKAYDDDSSYYY GKYNNSTQGTFNNGPRYPYPSNGS QDYNRNYNSKIVAEQHNRGGLGAG MSGLSIEDNGRSKQLYSSYTEAANA NLNPLPSPPTQWIGTQPGGNFVGGY YRDGVGYSETNGKSVKKVIYQAKTQ PSHRGANL	None	None	None	None	None	None	None	None	None