

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A1P8B2G0	A0A1P8B2G0_ARATH	Arabidopsis thaliana	RNA polymerase II C-terminal domain phosphatase-like	27.144759	NaN	NaN	28154133	MNNISKERNKETMSRLLTLVNDHF SQFLSFNQKNEIETMNQDLRSIAIA VFAGTSSEENVNQMQTPSNGDSFL AKKLTSESTRGAAYLRSRLPMLPLL DLHKDHDADSLPSPTRETTPLSPVN GRHTMVRPGFPVGRESQTTEGAKVY SYESDARKAVSTYQQKFGFLNSVFKT DDLPSPTPSGEPNDGNGDVGGEVSS SVVKSSNPGSHLIYGQDVPLPSNFN SRSMPVANSVSTVPPHLSIHAIISA PTASDQTVKPSAKSRDPRRLRAKPA ANVTIYSYSSGDARNLSKVELSADLV NPRKQKADEFLLDGPWKRQKSDT DAPKAAGTGGWLEDTESSGLLKLES KPRLENGVTSMTSSVMPTSAVSVS QKVRTASTDTASLQSLKDIANPT MLLNLLKMGERQKVPEKAIQKPM PRRAAQLPGSSVQPGVSTPLSIPASN ALAANSLNSGVLQDSSQNAPAAESG SIRMKPRDPRRILHGSTLQRTDSSM EKQTKVNDPSTLGTLMKGKAEDLE TPPQLDPRQNISQNGTSKMKISGEL LSGKTPDFSTQFTKNLKSADMMVVV SQQLGNPPASMHSVQLKTERDVKH NPSNPNAQDEDVSVSAASVTAAGP TRSMNSWGDVEHLFEGYDDIQRVAI QRERVRRLAEQNKMFASQKLSLVD IDHTLLNSAKFNEVESRHEEILRCKE EQDREKPYRHLFRFLHMGMWTKLR PGIWNFLEKASKLYELHLYTMGNKL YATEMAKLLDPKGVLFNGRVISKGD DGDPLDGDERSPKSKDLEGVMGME SSVVIIDDSVRVWPQHKNLIAVER YLYFPCSRQFGLLGPSELELDRDEV PEEGTLASSLAVIEKIHQNFHSHTSL DEVVVRNILASEQRKILAGCRIVFSRI IPVGEAKPHLHPLWQTAEQFQAVCT TQVDEHVTHVVTNSLGTDKVNWAL TRGRFVVHPGWVEASAFLYQRANE NLYAINP	None	None	None	None	None	None	None	None	

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