

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
A0A1P8AQ08	A0A1P8AQ08_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MAAPRPTGGQDLFDTYFRADLDGD GHISGAEAVAFQGSNLPKHVLAQV WSYADSKKAGYLGRAEFYNALKLVT VAQSRRELTAIEIVKAAIYSPASANIPA PKINLAATPSPQPRGVLPAQAGVT SMPSVAAGVRGPHMGTVSTSNQQ VVPQQNQFTGIPPSQTQNFQSPG MPAGGTNAPRANQPMPSDWLSGR SVGPSGNVNSQIPSSQSTYGLTAPNS TANHITKPHITPAVTSSTTRPQESA PVHNPQESSATFGSRVSNVPSNQLV PKDPKELAASGNGFTSDSLFGDVFS VTSTQPKQHPTGSASTTGISSVTTGT VAGPEITQSVVRQSSIPOQGSLSQHA VGVQTQLTGNISGQPYTSSGAASGPP GSTVGVGISATSQLAQRPPHPSQP QPRPQGSQPPWPKMTPADVQYKT KVFVQVDTDRDGKITGNQARNLFLS WRLPRDALQVWDLSDQDNDSMLS LREFCIAVYLMERYREGRPLPVFPPS SIIHSESMFTSPGQSVAPHGNASWG HPQGFQQPHPGGLRPPAGPKGKPP RPVPLSPSDGMVQTPQPKRKMVLE KPLVDQLSKEEQDSLNTKFEEATAV DKKVELEKEIADSKQKIDFFRAKM QELVLYKSRCDNRYNEIAERVLGDK RELESLAKKYEEKYKSGNVGSKLTI EEATFRDIQEKKMELYQAIVKFEEGK LDDSVKERTHEHIQSGLEELIKNLNE RCKQYGVRGKPTSLVELPFGWQPGI QEGAADWDEDWDKLEDEGFTFVKE LTLDIQNVIAAPPKEKSSAWRKEVDVS SKEGEDVSFSDADSKTGKKQSSGEE DSEQSEGKTSVDVARDKNGSLDDS KVRKGIEADSSPRTKDTRSENGHDD GESTASAGKTVNYDSHDETDVSSV NPDNGKDKDHGKYDSGFGFGFGFD DFSIKPIKTGSTISNDFLPPKLSIFAD SVSPPANASDVSPTKPSLFADSVPS TPATNNASYPGQKSFDDSVSTPAY PGNLFAEKKSYPDDSVSTPAYSTSD FGGKPFASETPRSDNLFGRSPFPMF DSVPSTPAAHDDFSNNSFSRFDSPFN SNNDAFSLSRDSDMRSTSEPDPA SRFDSFNRYRYSFNAQSYDSSSNN NASETPKASLRFDSIGSTRDSDYSH GFGFDDHDPFGSTGPFKTTTTAET PRSSDNWNAF	None	None	None	None	None	None	None		

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