

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
A0A1P8ASK1	A0A1P8ASK1_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MDKARHQQLFOHSMPEGYRNETVP QPFMPDQTGSASANMRPPNSNGSD VKAVHNFSIQTGEEFSLFMRDRVI PQRSSNPNGAGDMNYNTGYMELRG LIGISHTGSECASDVSRFSTVENGT DIERTNSSLHEFGNKLNHVQSAPQA LLSKDSSVGNLHGYKNTSSSASGSV TAKVKILCSFGGKILPRPGDKLRYV GGETHIISIRKDISWQELRQKILEIYY QTRVVKYQLPGEDDLALVSVSSEED LQNMLEEYNEMENRGGSQKLRMF LFSISDMDDALLGVNKNDDGSEFQY VVAVNGMDIGSGKNSTLLGLDSSSA NNLAELDVRNTEGINTIAGDVVGVG ASQLMVNGFQQTSAQQSEIIPPSSS LHYSQSIPLNAAAYLQOQSVPPSSALH YPQSITPGSSLQYPQSITPGSSYQYPQ SIIPGSASSYGIYPQYGHVVQHGGER ERFPLYPDHSSNYSAGETTSSPIQG HVSQGGWAEGYPPGSTPKSTQAL AEEQKVSSDMKIREEVEPENRKT NDHQNPPIDDVEVRNHNQVREMA VATTPSQDAHLLPPSRDPRQNTTA KPATYRDAVITGQVPLSGIEDQLSTS SSTYAPVHSDSESNLIDLNYPEPEQS SRVYCSERIPREQLLELNRLSKSDN SLSSQFVTSESPANTAQQDSGKEAV GKSHDEFKTVNDDANHHHTHKDVE IFEKVGVSDETLSEPLHKIVNPPDA NKNRVVNGADTEIGVSNLSHVNA MSHVIPEEQASLQGDILINDRFP DFLSEIFSQAISEDSTVRPYPHDGA AVSMNVQNHDRKNWSYFQQLAED QFIQRDVVLDQADSRIPSDRKGGE SSRLPYVPLSRDGISTNLANPQLTL GQDYGGNFSEKGGGTGSIIPALEN EQMKVTESEFGAMVENLRTPDSE PKDEKTETRAALPPLGSEFDYSGL QIKNEDLEELRELGSFTGTVYHGK WRGSDVAIKRIKSCFAGRSSEQER LTGEFWGAEILSKLHHPNVVAFYG VVKDGGGTLATVTEYMDGSLRHV LVRKDRHLDRRRLIAMDAAFME YLHSKNTVHFDLKCNDLLVNLKDP RPICKVGDVGLSKIKRNTLVSGGVRG TLPWMAPELLNGSSSKVSEKVNIN MIRCTVNINYHFNC	None	None	None	None	None	None	None		

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