

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
A0A1P8B770	A0A1P8B770_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MDYDDSDFFQNLHLAGEANNKFP PVLQPYALPKFDDTLNTHLRFDS LGESEAFLEIEGNEEDNNWIEDFSRG SSGIVFSSGATESCAISRHNNVWSE ATSSSEVAMLLNSVGQDEVIVREDTI KKSDTSHELGCTMETVEPGQTSHER SLSKEETVNLQPNPSVDDTPGESSV VKTDGQEQVLVKDDSPAVEEASV EEKNSILTSNTATVEAIDTDLGKIGT ETDNLDDQTEEKANVESRMEDDC SDGNVQTHITCSARSILVEGHSDSH IDSASEVEKVEAENIGKTAKPDLKEI ELSDVTVLGERGQAPSTLEVGGQDV SGTECQDLLVSTVHTSVAVEASLELA GELTTITNSVIEKPELLSHQHMEVI TSEHESTFQIETETYPQIHVFETSES VYISTMDSMVEAREGGVSKKSDNEG SARTSNLEQSMELPVNANDRDQDV KNSQILSESVVSGSVGYVSGGSTSEL AESESQSDSIPTDKSETMIDSSLNLE ELQPLSQDGAPAVSLTSSIDLHMVK TSSDSDSQGSYSETKKVYGEPENGQ TVPPVDASCSSGQMDQEARAKRAEGT KQSTYSVEGCPRSEGSKDAVDADGV GQVLQQQSEELIFEENVVTEAVKAP ETLSVLDKDNKNEMPITSSLPILGSE AGKDGQEEEDNTAASGGIMAAGTPVT HPKGDIVLGDSTRASTCESSVKSIV TAIEDAATNLKTPLDSPFTVKTSELQ FNNTETNSVKKPEDQNISGFMSAGS PVLNRNETSSSEMNLTPDQLKAGKI SKAVIFSQATLVSPIVVGSPTSSSLDK TAAKSSKAKSERKPRRTSKSVGKETS RKGTSVKGATPIEQFQSGGKTNVAVN QSLASHIQITQSTEQRSLSQSPALKA FGSLSTPTASLPDLNSSLALSILRRPF TDLQQVQLRAQIFVYGALIQGTAPDE AYMISAFGGADGGKGSWEKSWRTC VVRAQKSLVATPETPLQSRPGKTETP SAGHTNSKESSTNPMIPLSSPLWS LSTSVDTLQSSSVQRGSAATHQPLLS ASHAHQTPPTQNIIVGHNTPWMSPL PFRNAWLASQQTSGFDVGSRFVYP ITDPVKLTPMKESMTLSGAKHVQS GTSNVSKVTPTEPTSTVVAPAQHS TRVSRKRKMPVSVESGPNILNSL KQTELAASPLVPFTPTPANLGYNAGT LPSVVSMTAVPMDLVSTFPGKKIKSS FPSPIFGGNLVREVKQRSVLESDTIE KLKEAKMHAEDASALATAAVSHSEY VWKQIEQQSHAGLQPETQDRLASAA VAIAAAAAVAKAAAAANVAANAAL QAKLMAEEASLPNASDQGLPKSYDS ILPGQTPASILKGEAVVNSSSVLIA AREASKRVEAATAATKRAENMDSI VKAELASEAVSQAGILVSMGHPPLL NKLVEAGPSNYWRQAQESQEVQPC KTVVLEKETVSTSEGTAFAGPKIVQTE FGGSVNTADGVSGVPATGKLGQGE GDYADLAKNNDVVFPEVGSKFSI DAQQTIKATKNEEDIKESNVVEFKE EPGLRTAWYSANVLSLEDDKAYVLF SDLVEQGTDLKKEVWALKGEGDQ APKIRPARSVTALPYEGTRKRRRAAL GDHIWKIGDRVDSWVHDSWLEGVIT EKNKEDENTVTVHFPAAAAEETLTIKA WNLRPSLVWKGDKWIECSSSGETIS SSHEGDTPEKRPRLGTPALVAEVK DTSMKIVDDPDLGKPPQTGVLNLGV SENTFNIGKSTREENKDPRLMKRT GLQKQGSKVFVGPVKGKRRKFMV SKHYVSEASTKTQERKEPVKPVRSIV PQNSGIGSWKMPKSTISIEKQTTISR PKTFKPAKPKKEKPGATARIIIPKDS RNTTASDMESESAENRGPVSGSVS FKGTVEEQTTSSSHDTGSKNSSLS	None	None	None	None	None	None	None		

