

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
A0A178UVW2	A0A178UVW2_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MSQTNWEADKMLDVYIHDYLVKRD LKATAQAFQAEQKVSDDPVAIDAPG GFLFEWVWVWFWDIFIARTNEKHSEV AASYIETQMIKAREQQLQSQHPQV SQQQQQQQQQIQMQLLQRAQQ QQQQQQQHHHHQQQQQQQQ QQQQQQQQQHQNQPPSQQQQQ STPQHQQPTPQQPQRRDGSHLA NGSANGLVGNNSEPVMRQNPQSGS SLASKAYEERVKMPTQRESLDEAAM KRFQDNVQQLDPSHASILKSAAS GQPAGQVLHSTSGGMSPOVQTRNQ QLPGSAVDIKSEINPVLTPRTAVPEG SLIGIPGSNQGSSNNLTKGWPLTGF DQLRSGLLQQKPFMQSQSFHQLN MLTPQHQQQLMLAQNLNSQSVSE ENRRKMLLNNRSMKSLGKDLGSS VGDVLPNVGSSLQPGGSLPRGDTD MLLKLKMLLQQQQHQQQGGN PPQPQPQPLNQLALTNPQSSN HSIHQEQEKLGGGSITMDGSISNSF RGNEQVLKNQSGRKRKQPVSSGPA NSSGTANTAGPSSAPSTPSTHTPG DVISMNLPHSGSSKSMFMFGTE GTGTLTSPSNQLADMDFVEDGSLD DNVESFLSQEDGDQRDAVTRCMDV SKGFTFTEVNSVRASTTKVTCCHFS SDGKMLASAGHDKKAVLWYDTMK PKTTLEEHTAMITDIRFSPQLRLAT SSFDKTVRVWDADNKGYSRLRFMG HSSMVTSLDFHPKDDLICSDNDN EIRYWSINNGSCTRVYKGGSTQIRFQ PRVGKYLAASSANLVNVLVETQAIR HSLQGHANPINSVCWDPSGDFLASV SEDMVKVWTLGTGSEGEVHELSC NGNKFQSCVFHPAYPSLLVIGCYQS LELWNMSENKMTLPAHEGLITSLA VSTATGLVASASHDKLVKLVK	None	None	None	None	None	None	None		

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