

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A1I9LRR1	A0A1I9LRR1_ARATH	Arabidopsis thaliana	ATP-dependent RNA helicase	27.144759	NaN	NaN	28154133	MMLEKSKSRKENDRKDRDRSKKEN GRRDTTEMRSRVKRCDSSEEEERIRI RRDRKSSDFEEEEYERDSKRRGEDK GRGRRERDRDRGKYLKRDRERRER EKEKGRKKQKKERSREDCNEESDD VKCGLKRRRTERSRHGDDVVEKTR DEQVEDEQKQLAEVEKRRRRVQE WQELKRNNEEAQIESKGPETGKAW TLDGESDDEVKSDSEMDVDRDTKL ENGGDAKMVAENETAVTVSENGG DRAADEDEIDPLDAFMNTMVLPEVE KLSNIVIDGILDFKMNGKETGDOAK KGFNKAALGRIIQGEDSDSYSEPKS DDDPDLDEDEEFMKRVKTKAEKL SLVDHASKIEYEPFRKNFYIEVKDISR MTQDAVNAYRKELELKVHGKDVPR PIQFWHOTGLTSKILDTLKKNYEKP MPIQAQALPIIMSGRDCIGVAKTGSG KTLGFVLPMLRHIKDQPPVEAGDGPI GLVMAPTRELVQQIYSDIRKFSKALG IICVPVYGGSGVAQQISELKRGTIIVV CTPGRMIDILCTSSGKITNLRRTYL VMDEADRMFDMGFEPQITRIVQNIR PDRQTVLFSATFPRQVETLARKVLN KPVEIQVGGRSVVNKDITQLVEIRPE SERFSRLLELLGEWYKGVVLFVVR SQEKSISDFKSDVCNLLIATSVAARG LDVKELELVNFDAPNHVEDYVHRV GRTGRAGRKGCAVTFISEDDAKYAP DLVKALELSEQVPPDDVKAVAEGFM AKVKQIEQAHTGYGGSGFKFNEE EDEVKAAKKAQKEYGFEEKSDS EDENDVVRKAGDISQQITLAQIAA IASAASKAPVTANQLLPNGGGLATEP GIPPTDAGRVAAMIAANVQQYLA KIQADAIPEHYEALEINDFPQNARW KVTHKETLGPISWGSASITTRKGFY EAGRIPGPEERKLYLVEGPTAISVKT AKAELKRVLEDITNQTFSLPGGAQS GRYSVL	None	None	None	None	None	None	None		