

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
A0A178VNI5	A0A178VNI5_ARATH	Arabidopsis thaliana	ATP-dependent RNA helicase	27.144759	NaN	NaN	28154133	MLEKSKSRKENDRKDRDRSKKENG RRDTTEMRSRVKRCDSSEEEERIRIR RDRKSSDFEEEEYERDSCRKRGEDKG RGRRRERDRDRGKYLKRDRRERERE KEKGRKKQKKERHGDDDVKKTRD EQVEDEQQLAEEVEKRRRRVQEW QELKRQNEEAQIESKGPETGKAWTL DGESDDEVKSDSEMDVDRDTKLEN GGDAKMVASENETAVTVSENGGDR AADEDEIDPLDAFMNTMVLPEVEKL SNIVIDGILDFKMNGKETGDAQKKG FNKAALGRIIQGEDSDSYSEPKSDD DPSLDEDEEFMKRVKTKAEKLSL VDHSKIEYEPFRKNFYIEVKDISRMT QDAVNAYRKELELVHVKDVPRIQ FWHQGLTSKLLDTLKKLNYEKPMP IOAQUALPIHMSGRDCIGVAKTGSGKT LGFVLPMLRHIKQPPVEAGDGPGL VMAPTRELVQQIYSDIRKFSKALGHC VPVYGGSGVAQQISELKRGTIVVCT PGRMIDILCTSSGKITNLRVTVLVM DEADRMFDMGFEPQITRIVQIRPD RQTVLFSATFPRQVETLARKVLNKPV EIQVGGRSVVNKDITQLVEIRPESER FSRLLELLGEWYKGGKLVFVRSQE KCDALYNLMTKCGYACLSLOGGK EQTDRESAISDFKSDVCNLLIATSVA ARGLDVKKLELVNFDAPNHIEDYV HRVGRTGRAGRKGCVAFTFISEDDAK YAPDLVKALELSEQVPPDDVKAFAE GFMAKVKGIEQAHGTGYGSGGFKF NEEEDVVRKAQKQAKKEYGFEEK SDESEDVVRKAGGDISQQITLA QIAAIAASAASKAPVTANQLLPNGGGL ATEPGIPPTDGAGRVAAMIAANVQ QYLAQIQADAIPEHYEAELEINDFPQ NARWKVTHKETLGPSEWGSASITT RCKFYEAGRIPCPEERKLYLFVEGPT EISVKTAKAELKRVLEDITNQTFSLP GGAQSGRYSVL	None	None	None	None	None	None	None		