

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A1I9LT31	A0A1I9LT31_ARATH	Arabidopsis thaliana	NaN	27.144759	NaN	NaN	28154133	MGEHESWAASPPSPGLHPNGLLP GKAASVTRPLDAERWAKAEDRTAKL IACIQPNPPSEDRRNASYVRRLLIM ECFPQVQIFMFGSVPLKTYLPDGDID LTAFSANQNLKDSWANLVRDMLEK EEKNENAEFHVKEVQYIQAEVKIKC LVENIVVDISFNQIGGLCTLCLFEEV DHYINQNHFKRSIILKAWCYYESRI LGAHHGLISTYALETLVLYIFLNN SFSGPLEVLYRFLFFSKFDWQNF LSLWGPVVPVSSLPDVTAEPPRRDVG ELRVSEAFYRACSRVYAVNIAPQEIQ GQPFVSKHFNVIDPLRENNNLGRSV SKGNFFRIRSAFTLGAKKLRLLLECP KENLIHEVNOFFMNTWERHGSGR PDAPGNLWLSRLGDPPEYHQAEN VSNLNNKRNQNAIRLGGVHGARS MPSQONNCGTEITSRVTYQTQKSRG NSYQPAQEVNSNQSALNDKLOQTV KPETLVNNFHGRHIFARTRSSPELT ETHGELLQSRRSRAAPDAGKRQTN STRVDSIRKKSLESETLSSGVRYSD SSSVRHTPSPQSPDSTADMSSAVNS YYDEVGSVSVNEDFSVAGEQEEQDL VNSMTSVTGQGFNGHFPFNFST GHLFPFITPSILASMGYQQRNMPGIV PSNLPFIEAPWSTNLQFPQNFVSSPF THYFPGAHPISEKPKSTGSDDMGS SEVNVDESNDLWHEPERGTHSFG LENGGYGMHQADDKHSSFAEHSF VPSRRKNRLTRGDLENLHSPVRS SQIQSEERTVGSRSVSGASSVRSRTS SESSWDGSTTRGSKPAKDRRNRKVV SGAASLYGKGSVPEHSIQIDDDNR EWIPVSSNEIDRDLGPRPTVPSFQV QRHQIHGHELAQASGSESTVSLAPFI LGHGMQQNEVDNSGYTFYPTGPPV PIVAMLPYNYQAGGNATSDALASH HSVDEGVENHEPCKSFDSSRGLDQ SEIVVSSHSTRMGSSAEQVERKNDIL NGDFISHWQNLQYGRSCONSQHP VLYPAPVVVPPAYLQGRLPWDGPR PLAYTNAVNLMTYGPRLVPVAVQ PVSTRPPNIYPRYANETPRYRSGTGT YFPNPKISPREQRPTSGMRRGNYGH DRTDHHSQDREGNWNAGSKTRGSGR NHNNRNQADNKPISRQDRSDRHW GSSYRHESSESYSAHHSQNGPIRSNT SQDASGNIAYGGEAPHLNDEDQPRF RGAASAAMSSPDDPSSPHFPRCK	None	None	None	None	None	None	None		

0