

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
Q67YI9	EPN2_ARATH	Arabidopsis thaliana	Clathrin interactor EPSIN 2	27.144759	NaN	S270;S282	28154133	MKKVFGQTVRDLKREVNKKVLKVP GVEQKVLDATSNPEPWGPHGSLADL AQASRNYHEYQLIMVVIWKRLSDTG KNWRHVYKALTVLEYMVGHGSEVI DEIRERAYQISTLSDFQYIDSGGRDQ GSNVRKKSQSLVALVNDKERIAEVR QKAAANRDKYRSSAPGGMYKPSGGY GDKYDYGSRDEERSYGREREYGYR DDDRNSRDGDRHSRDEDRYGRDG NRDDDYRGRSRSDNYGSRGRSSE REREDDGHSSSRGSGARADDNSQD GRGGLQRKFSEONIGAPPSYEEAVS DSRSPVYSERDGGETPQVTAPGAAS PPPPQVAPEAASPPTGTNTANTTAT FVNESPSQKVETFDPRSAFASAG PPAYASTDGVTPPTVTMSAPTTSN SVEMDLLGSLADVFSNALAIVPADS IYVETNGQANAGPAPSFSTSQPSTOS FDDPPFGDSPFKAFTSTDIDSTPQQN FGASFQPPPAFTSEVSHPDTAHNF FGGDSFSAVANPDPASQNVQPPSNS PGFPQEQFATSQSGIDILAGILPPSGP PVQSGPSIPTSQFPPSGNNMYEGFH SQPPVSTAPNLPGQTPFGQAVQPYN MVPHSQNMTGAMPFNSSGGFMHQP GSQTPYSTPSGPAGQFMAHQGHGM PPSHGPORTQSGPVTLQGNNNVMG DMFSQATPNLSLTSSSSHPDLTPLTG AIEIVPPQKKFEPKSSVWADTLSRG LVNFNISGSKTNPLADIGVDFEAINR REKRLEKQTNTPATSTINMGKAMGS GTGLGRSGATAMRPPPNPMTGSGM PMGGGMGVGSYGGMNQNPQPMGM GMGAGMNQNPQPMGMGMPGMN MNMNMGYGGYPMQPNPQPMV PSPNMPGNPNMMPMMGQGGYNPQQ SYGGGYR	None	None	None	None	None	None	None		

0