

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	ex re
Q05BC3	EMAL1_MOUSE	Mus musculus	Echinoderm microtubule-associated protein-like 1	47.329949	T126	NaN	22645316;40885482;36288343;35822049;40021952	MEDGFSSYSSLYDTSSLLQFCNDDSA ASAASMEVSDRIASLEQRVMQED DIQLLSALADVRRRLNITEEQQAVL NRKGPTKARPLGQTLPLRRTVNNGT VLPKPSASLPAPSGARKEVVPVTK SINRTSSSERVSPGGRRESSGDSKGS RNRTGSTSSSSGKKNSESKPKEPA FSP EEGYVKMFLRGRPVTMYMPKD QVDSYSLEAKAELPTKRLKLEWVYG YRGRDCRNNLYLLPTGETVYFIASVV VLYNVEEQQRHYAGHNDDVKCLA VHPDRITITATGQVAGTSKDGKQLPPH VRIWDSVTLNTHVIGIGFFDRAVTC IAFSKSNGGGHLCAVDDSDNDHVLVS WDWQKEERLADVKCSNEAVFAADF HPTDTNIIVTCGKSHLYFWTLEGNSL NKKQGLFEKQEKPKFVLCVTFSENG DTITGDSSGNILVWGKGTNRISYAVQ GAHEGGIFALCMLRDGTLVSGGGKD RRLISWNGNYOKLHKAEIPEQFGPIR TVAEGKGNVILGITTRNFVLOGTSLG DFTPTIQGHTDELWGLAIHASKPQFL TCGHDKHATLWDVAVGHRPWWDKII EDPAQSSGFHPSGSVAVGTLTGRW FVFDTETKDLVTVHTDGNEQLSVMR YSPDGNFLAIGSHDNCIYIGVTDNG RKYTRVFKCSGHSFITHLDWSVNS QFLVNSGDYEILYVWPSACKQVVS VETTRDIEWATYCTLGFHFVGVWP EGSDGTDINAVCRAHERKLLCTGDD FGKVHLFSYPCSQFRAPSHIYSGHSS HVTNVDFLCEDSLITGGKDTSIM QWRVI	True	False	5.0	2.996	0.687	False	1.038	1.082	0.7