

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
Q99K41	EMIL1_MOUSE	Mus musculus	EMILIN-1	25.685855	NaN	NaN	39627609	MAPRALWSCYLCCLLTATEAASYPP RGYSLYTGGTGALSPGGPQAQNSPR PASRHRNWCAYVVRTVSCVLEDGV ETIVKPDYQPCGWGPHCSRSIMYR SFRLPRYRVAYKTVTDMEWRCCQGY GGDDCGEGPASVLGPAPSTPLPRPR PVRPNLSGSSAGSHLSGLGGEGPVE SEKVQQLERQVKSLTKELQGLRGVL QGMNGRLAEDVQRAVDTVFNQRQQ PADAAARPVHETLSEIQQLQLLD NRVSTHDQELGHLNNHHNGGPGG GGRASGVPVPSGPFSELLRQLERQ LQESCSVCLTGLDGFRRQQQEDRER LRTLKLMSSMEERQQQVGPAMA RRPPQECCPELGRRVSELERRLDV VTGSLTVLSGRRGSELGGAAGQGGH PPGYTSLASRLSRLEDRFNSTLGPSE EQEKNWPGGPGRLGHWLPAPGRL EKLEGLLANVSRLEGGMDLLEEQV AGAVRTCQICSGAPGEQDSRVNEI LSALERRVLDSEGRQLQVGSGLHEA EAAGEAQAVLEGLQGLLSRLRERM DAQEETAAEILLRLNLTAALSQLEG LLQARGDEGCGACGGVQEEGLRLR DGVERCSCPLLPPRGGAGPGVGGP SRGPLDGFVFGSSGSALQALQGE LSEVILTFSSLNDSLHELQTTVEGQG ADLADLGATKDSIISEINRLQEQATE HVTSEERFRGLEEGQAQAGQCP EGRLGRLEGVCERLDTVAGGLQGLR EGLSRHVAGLWAAVRESNSTSLTQA ALLEKLLGGQAGLRRRLGALNNSLL LLEDRLQQLSLKDFTPSGKAGPPG PPGLQGPSGAPGPPGPKDGQQA IGPPGPQGEQGAEGAPAAPVPRVAFS AALS LPRSEPGTVPFDRVLLNDGGYY DPETGVFTAPLAGRYLLSAVLTGHR HEKVEAVLSRSNLGVARIDSGGYEP EGLNKPVAESQPSGALGVFSLILP LQVGD TVCIDLVMGQLAHSEPLTIF SGALLYEDTELEQV	False	True	0.422	0.946	0.469	0.549	0.719	2.634	4.645