

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
E9PYF4	E9PYF4_MOUSE	Mus musculus	NaN	25.161187	NaN	NaN	34887587	MEGMEDAEADCSVAFEAQRWVEA VTEKNFETDDFRASLENGVLLCDLI NKLKPGVVKKINRLSTPIAGLDNINV FLRACEQIGLKEAQLFHPGDLQDLS NRVTVKQEETDRRLKNVLITLYWLG RKAQSNPYNGPYLNLKAFETLLGQ ALTKALEDDSSCLKRSGRDSGYGDIW CPDRGEFPAPPGSHRREDSFESLDS LGSRLTSCSSDFTLRVKGEGCESDI DWEFTFKMQDYNKDDMSYRRISAIE PKSALPFNRFLPNKSKQPSVVPAPLR KKRPDKHEDNRRSWASPVYETDGDG TFSSTQRRTWGPKMETWHTVQETS TSSWCVEEEEEKLTRMPNIVKDDLY VRKLSVMPSPGSSFDQFLPKCWTP EDMNWKKIKRETYKPWYKEFQGFSS QFLLLQALQTYSDDILSSETSVMKLDP TSGPRLITRRKNRSPAGYRAHLEL PALDPDLENDDFFVRKTGAFHANPC VLRAFEDFRSFSEPDDAVERDIILQC REGELVLPDLEKDDMIVRRIPAQKK EVPLSGAPDRYQPVFPFWTLPEI QAKFLSVLERTRPPEQSSGCRVLVP SYRQKDDMLARKIQSWKLGTAVPP ISFKPGPCSEADLQWEAIWEASRL RHRKRLMVERLFQKIYGENGSKSM SDVSAEDVQNLRQVRYEEMQKIKSQ LKEQDQKWQDDLAKWKNRRKSYTS DLQKKEREIEIKQALEKSDRSSK TFREMLQDRESRGSTVTSWRRTYSS DDILDDGVLPTVTLSETSYQSERLE DEARAHPAEMPKESTTFANREDSV VAETQLASHSPPEEQRLAPSPSEEP QASGSSAEPHQAPLLSTEPRTPLPS EEPHQASLSSTLQRPSSLSQQSWS TRAESTRISASLPRSYQRTDARLTS VVTARPFGTSSRGISLPRSYTMDDG WKYNGDVEVVKREQLDLGRATGPK PDSSHFISGSASEKEVIATEDVASLS SPTLPSSSLSHDRAVSSKATFSSMSG LDSVSDSGEGRGSPLREVSRLDQF SDMRVSINQTPGNKPVFGFTIKWDI SGIFVASVEQGSPAEFSLQVDDDEIL AINNTKFSYKDTKKWEEVMANAQE TGNLVMDVRRYGKSGSSETKWIDTT SGIYNSDKSSSLSVTTDFSESLSQSPY TESKEVNGIHEESNPFDSKASESISL KNLKRSSQFFEQGSSDSVAPDLVP TLSAPSRWAWDQEEERRRQERWQK EQDRLLQEKYQREQEKLREEWQRA QQEAERENSKYLDEELMVLNSNSIS LTSREPVAATWEATWSEGSKLDSE GTRAGEEDRGQLEDDAVYEDQSOKL QELEQEKKRKEQEAQEEERRKQRE AQAWAEAEAKAWAEQAQAEAEAK ARAEQAQAEAEAKRAEAQKLAQAE RERETSVKIYORRRPIDSYDLPKRED DTSGMLPTDRSKSRSTELNDPLIEK NGSSKYSERIGATSFHRSSKKDQA PSEAELEERQQLQEMRKRTSLHNDN SWIRQRSASVNKEPICLPGIMRRGES LDNLDQRPGSWRQSPWHSQPAGV YASSVQDFSRPPPQLLSTSNRAYM RNPSSAVPPPAGSVKTPSGSPSPRSH SPSMSQSGQLRNRSVSGKRVCSYC NNILGKGAAMIETLGLSYHLHCFKC VSCGCDLGGSSSGAEVRIRNHQLYC NDCYLRFKSGRPTAM	True	True	1.402	5.0	1.171	1.156	0.621	5.0	1.217