	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P57678	GEMI4_HUMAN	Homo sapiens	Gem-associated protein 4	24.348488	NaN	T84;S86;S205	34725712;23301498	MDLGPLNICEEMTILHGGFLLAEQL FHPKALAELTKSDWERVGRPIVEAL REISSAAAHSQPFAWKKKALIIIWAK VLQPHPVTPSDTETRWQEDLFFSVG NMIPTINHTILFELLKSLEASGLFIQL LMALPTTICHAELERFLEHVTVDTSA EDVAFFLDVWWEVMKHKGHPQDPL LSQFSAMAHKYLPALDEFPHPPKRL RSDPDACPTMPLLAMLLRGLTQIQS RILGPGRKCCALANLADMLTVFALT EDDPQEVSATVYLDKLATVISVWNS DTQNPYHQQALAEKVKEAERDVSLT SLAKLPSETIFVGCEFLHHLLREWG EELQAVLRSSQGTSYDSYRLCDSLTS FSQNATLYLNRTSLSKEDRQVVSEL AECVRDFLRKTSTVLKNRALEDITAS IAMAVIQQKMDRHMEVCYIFASEKK WAFSDEWVACLGSNRALFRQPDLV LRLLETVIDVSTADRAIPESQIRQVIH LILECYADLSLPGKNKVLAGILRSWG RKGLSEKLLAYVEGFQEDLNTTFNQ LTQSASEQGLAKAVASVARLVIVHPE VTVKKMCSLAVVNLGTHKFLAQILT AFPALRFVEEQGPNSSATFMVSCLK ETVWMKFSTPKEEKQFLELLNCLM SPVKPQGIPVAALLEPDEVLKEFVLP FLRLDVEEVDLSLRIFIQTLEANACR EEYWLQTCSPFPLLFSLCQLLDRFSK YWQLPKEKRCLSLDRKDLAIHILELL CEIVSANAETFSPDVWIKSLSWLHR KLEQLDWTVGLRLKSFFEGHFKCEV PATLFEICKLSEDEWTSQAHPGYGA GTGLLAWMECCCVSSGISERMLSLL VVDVGNPEEVRLFSKGFLVALVQVM PWCSPQEWQRLHQLTRRLLEKQLL HVPYSLEYIQFVPLLNLKPFAQELQL SVLFLRTFQFLCSHSCRDWLPLEGW NHVVKLLCGSLTRLLDSVRAIQAAGP WVQGPEQDLTQEALFVYTQVFCHAL HIMAMLHPEVCEPLYVLALETLTCY ETLSKTNPSVSSLLQRAHEQRFLKSI AEGIGPEERRQTLLQKMSSF	True	False	5.0	5.0	0.997	0.765	0.911	1.043	4.225