	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	extracellul region
0	Q9Z0H8	CLIP2_MOUSE	Mus musculus	CAP-Gly domain-containing linker protein 2	26.300724	T355;T356	S50;S203;S208;S212;S315;S924;S974;S980	29562282	MQKPSGLKPPGRGGKHSSPVGRPSV GSASSSVVASTSGSKEGSPLHKQAS GPSSSGAATTVSEKPGPKAAEVGDD FLGDFVVGERVWVNGVKPGVVQYL GETQFAPGQWAGVVLDDPVGKNDG AVGGVRYFECPALQGIFTRPSKLTRQ PTAEGSGSDTHSVESLTAQNLSLHS GTATPPLTGRVIPLRESVLNSSVKTG NESGSNLSDSGSVKRGDKDLHLGD RVLVGGTKTGVVRYVGETDFAKGEW CGVELDEPLGKNDGAVAGTRYFQCP PKFGLFAPIHKVIRIGFPSTSPAKAKK TKRMAMGVSALTHSPSSSSISSVSSV ASSVGGRPSRSGLLTETSSRYARKIS GTTALQEALKEKQQHIEQLLAERDL ERAEVAKATSHICEVEKEIALLKAQH EQYVAEAEEKLQRARLLVENVRKEK VDLSNQLEEERRKVEDLQFRVEEES ITKGDLETQTQLEHARIGELEQSLLL EKAQAERLLRELADNRLTTVAEKSR VLQLEEELSLRRGEIEELQHCLLQSG PPPADHPEAAETLRLRERLLSASKEH QRDSTLLQDKYEHMLKTYQTEVDKL RAANEKYAQEVADLKAKVQQATTEN MGLMDNWKSKLDSLASDHQKSLED LKATLNSGPGAQQKEIGELKALVEGI KMEHQLELGNLQAKHDLETAMHG KEKEGLRQKLQEVQEELAGLQQHW REQLEEQASQHRLELQEAQDQCRD AQLRAQELEGLDVEYRGQAQAIEFL KEQISLAEKKMLDYEMLQRAEAQSR QEAERLREKLLVAENRLQAAESLCS AQHSHVIESSDLSEETIRMKETVEGL QDKLNKRDKEVTALTSQMDMLRAQ VSALENKCKSGEKKIDSLLKEKRRLE AELEAVSRKTHDASGQLVHISQELL RKERSLNELRVLLLEANRHSPGPER DLSREVHKAEWRIKEQKLKDDIRGL REKLTGLDKEKSLSEQRRYSLIDPAS PPELLKLQHQLVSTEDALRDALNQA QQVERLVEALRGCSDRTQTISNSGS ANGIHQPDKAHKQEDKH	None	None	None	None	None	None	None	None	None