

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
Q9UDT6	CLIP2_HUMAN	Homo sapiens	CAP-Gly domain-containing linker protein 2	17.826004	NaN	S49;S202;S207;S211;S314;S923;S973;S979	38665916	MQKPSGLKPPGRGGKHSSPMGRSTS TGSASSSAVAASSKEGSPLHKQSS GPSSSPAAPAAPEKPGPKAAEVGDD FLGDFVVGERVVWVNGVKPGVVQYL GETQFAPGWAGVVLDDPVGKNDG AVGGVRYFPCPALQGIPTRPSKLTRO PTAEGSGSDAHSVESLTAQNLSLHS GTATPPLTSRVIPLRESVLNNSVKTG NESGSNLSDSGVSVKRGEKDLRLGDR VLVGGTKTGVVRYVGETDFAKGEWC GVELDEPLGKNDGAVAGTRYFQCPP KFGLFAPIHKVIRIGFPSTSPAKAKKT KRMAMGVSALTHSPSSSSISSVSV ASSVGGRPSRSGLLTETSRYARKIS GTTALQEALKEKQOHIEQLLAERDL ERAEVAKATSHICEVEKEIALLKAQH EQYVAEAEKLRARLLVESVRKEK VDLSNQL EEERRKVEDLQFRVEEES ITKGDLETQTQLEHARIGELEQSLLL EKAQAERLLRELADNRLTTVAEKSR VLQLEELTLRRGEIEELQCLLHSG PPPDHPDAAEILRLRERLLSASKEH QRESGVLRDKYEKALKAYQAEVDKL RAANEKYAQEVAGLKDQVQATSSEN MGLMDNWKSKLDSLASHQKSLD LKATLNSGPGAQQKEIGELKAVMEG IKMEHQLELGNLQAKHDLETAMHV KEKEALREKLQEAQELAGLQRHW RAQLEVQASQHRLELQEAQDQRRD AELRVHELEKLDVEYRGQAQAEFL KEQISLAEKKMLDYERLQRAEAQGK QEVESLREKLLVAENRLQAVEALCS SQHTHMIESNDISEETIRTKETVEGL QDKLNKRDKEVTALTSQTEMLRAQV SALESKCKSGEKKVDALLKEKRRLE AELETVSRKTHDASGQLVLISQELLR KERSLNELRVLLLEANRHSPGPERD LSREVHKAEWRIKEQKLDKDIRGLR EKLTLGLDKEKSLSDQRRYSLIDRSSA PELLRLQHQLMSTEDALRDALDQAO QVEKLMAMRSCPDKAQTIGNSGS ANGIHQODKAQKQEDKH	True	False	2.987	4.288	0.641	0.901	1.134	0.972	0.731