	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi appara
0	Q8BRT1	CLAP2_MOUSE	Mus musculus	CLIP-associating protein 2	42.345557	T475;T476	S14;S20;S322;S333;S336;S374;S376;S413;S461;S465;S469;S484;S495;S513;S531;S535;S570;S572;S581;S614;S620;T779;S884;S944;S947;S1005;S1021	36852467;22645316;22517741;33300544;35822049	MRRLICKRICDYKSFDDEESVDGNR PSSAASAFKVPAPKTPGNPVSSARKP GSAGGPKVGGPSKEGGAGAVDEDD FIKAFTDVPSVQIYSSRELEETLNKIR EILSDDKHDWDQRANALKKIRSLLV AGAAQYDCFFQHLRLLDGALKLSAK DLRSQVVREACITVAHLSTVLGNKF DHGAEAIVPTLFNLVPNSAKVMATS GCAAIRFIIRHTHVPRLIPLITSNCTS KSVPVRRRSFEFLDLLLQEWQTHSL ERHAAVLVETIKKGIHDADAEARVEA RKTYMGLRNHFPGEAETLYNSLEPS YQKSLQTYLKSSGSVASLPQSDRSSS SSQESLNRPFSSKWSTANPSTVAGR VSVGGSKANPLPGSLQRSRSDIDVN AAAGAKAHHAAGQAVRSGRLGAGA LNPGSYASLEDTSDKMDGTASDDGR VRAKLSTPLVAVGNAKTDSRGRSRT KMVSQSQPGSRSGSPGRVLTTTALS TVSSGAQRVLVNSASAQKRSKIPRSQ GCSREASPSRLSVARSSRIPRPSVSQ GCSREASRESSRDTSPVRSFQPLGP GYGISQSSRLSSSVSAMRVLNTGSD VEEAVADALLLGDIRTKKKPARRRYE SYGMHSDDDANSDASSACSERSYSS RNGSIPTYMRQTEDVAEVLNRCASS NWSERKEGLLGLQNLLKNQRTLSRI ELKRLCEIFTRMFADPHGKVFSMFL ETLVDFIQVHKDDLQDWLFVLLTQL LKKMGADLLGSVQAKVQKALDITRE SFPNDLQFNILMRFTVDQTQTPSLK VKVAILKYIETLAKQMDPRDFTNSSE TRLAVSRVITWTTEPKSSDVRKAAQS VLISLFELNTPEFTMLLGALPKTFQD GATKLLHNHLRNTGNGTQSSMGSP LTRPTPRSPANWSSPLTSPTNTSQN TLSPSAFDYDTENMNSEDIYSSLRG VTEAIQNFSFRSQEDMSEPVRRDPK KEDGDTICSGPGMSDPRAGGDAAD GSQPALDNKASLLHSMPLHSSPRSR DYNPYNYSDSISPFNKSALKEAMFD DDADQFPDDLSLDHSDLVAELLKEL SNHNERIEERKIALYELMKLTQEESF SVWDEHFKTILLLLLETLGDKEPTIR ALALKVLKEILRHQPARFKNYAELTV MKTLEAHKDPHKEVVRSAEEAASVL ATSISPEQCIKVLCPIIQTADYPINLAA IKMQTKVIERVSKETLNMLLPEIMPG LIQGYDNSESSVRKACVFCLVAVHA VIGDELKPHLSQLTGSKMKLLNLYIK RAQTGSAGADPTADVSGQS	None	None	None	None	None	None	None