	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	Q99JD4	CLAP2_RAT	Rattus norvegicus	CLIP-associating protein 2	23.439218	NaN	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	38843836	MRRLICKRICDYKSFDDEESVDGNR PSSAASAFKVPAPKTPGNPVNSARKP GSAGGPKAGGTSKEGGAGAVDEDD FIKAFTDVPSIQIYSSRELEETLNKIR EILSDDKHDWDQRANALKKIRSLLV AGAAQYDCFFQHLRLLDGALKLSAK DLRSQVVREACITVAHLSTVLGNKF DHGAEAIVPTLFNLVPNSAKVMATS GCAAIRFIIRHTHVPRLIPLITSNCTS KSVPVRRRSFEFLDLLLQEWQTHSL ERHAAVLVETIKKGIHDADAEARVEA RKTYMGLRNHFPGEAETLYNSLEPS YQKSLQTYLKSSGSVASLPQSDRSSS SSQESLNRPFSSKWSTANPSAVAGR VSVGGSKASPLPGSLQRSRSDIDVNA AAGAKAHHAAGQAVRSGRLGAGAL NPGSYASLEDTSDKMDGTASEDGRV RAKLSTPLVAVGNAKTDSRGRSRTK MVSQSQPGSRSGSPGRVLTTTALST VSSGAQRILVNSASAQKRSKIPRSQG CSREASPSRLSVARSSRIPRPSVSQG CSREASRESSRDTSPVRSFQPLGPGY GMSQSSRLSSSVSAMRVLNTGSDVE EAVADALLLGDIRTKKKPARRRYESY GMHSDDDANSDASSACSERSYSSR NGSIPTYMRQTEDVAEVLNRCASSN WSERKEGLLGLQNLLKNQRTLSRVE LKRLCEIFTRMFADPHGKVFSMFLE TLVDFIQVHKDDLQDWLFVLLTQLL KKMGADLLGSVQAKVQKALDVTRES FPNDLQFNILMRFTVDQTQTPSLKV KVAILKYIETLAKQMDPGDFINSSET RLAVSRVITWTTEPKSSDVRKAAQSV LISLFELNTPEFTMLLGALPKTFQDG ATKLLHNHLRNTGNGTQSSMGSPL TRPTPRSPANWSSPLTSPTNTSQNT LSPSAFDYDTENMNSEDIYSSLRGV TEAIQNFSFRSQEDMSEPLKRDPKK EDGDTVCSGPGMSDPRAGGDAPDS SQPALDNKASLLHSVPLHSSPRSRD YNPYNYSDSISPFNKSALKEAMFDD DADQFPDDLSLDHSDLVAELLKELS NHNERIEERKIALYELMKLTQEESFS VWDEHFKTILLLLLETLGDKEPTIRA LALKVLKEILRHQPARFKNYAELTV MKTLEAHKDPHKEVVRSAEEAASVL ATSISPEQCIKVLCPIIQTADYPINLAA IKMQTKVIERVSKETLNLLLPEIMPG LIQGYDNSESSVRKACVFCLVAVHA VIGDELKPHLSQLTGSKMKLLNLYIK RAQTGSAGADPTTDVSGQS	False	True	2.733	2.031	1.387	0.976	4.291	5.0	0.806