

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q65CL1	CTNA3_MOUSE	Mus musculus	Catenin alpha-3	33.218136	NaN	S160;T361;S637;S647;T649	37507081;22645316	MSAETPITLNMMDTQDLQIQTFTVEK LLEPLIIQVTTLVNCPQNPSNRKKGGR SKRARVLLASVEEATWNLLDKGEMI AKEATVLKEELAAALQEVKESKALK VSAERFTDDPCYLPKREAVVQAARA LLAAVTRLLVLADMIDVIMCLLQHV SFQRTFESLKNVSNKSDLQRTYQKL GKELESLDYLAFAKROQDLKSPSQRD EIAGARATLKENSPLLHSICSACLEH SDVASLKASKDTVCEEIQNALDVISN ASQGIQNAPAPPEPQAATLGSAFDEL ENLIVLNPLTVTEEDVVRPSLEKRLEA IISGAALLADSSCTRDHLHRERIIAECN AIRQALQDLLTEYMSNTGKTERSNT LNTAIVNMSKKTRDLRRQLRKAIDH ISDSFLDITVPLLVIEAAKNGRVKEI KDYAIFHEHTGRLVEVANLACSMS TNEDGKIVRIAAHLETLCPIINAA LALASRPKSQVVKNMEMYKRTWE HYIHVLTAVDDITSIDDFLAVSESHI LEDVNCIIALRDQADNLDRAAGAI RGRAARVAHIVAGEMDSYEPGAYTE GVMRNVNFLTSTVIPEFVTQVNVAL DALSKNSLTALDDNQFVDISKIIDYD IHDIRCSVMMIRTPEELEDVSDLED DHEVRSHTSIQTEGKTDRAKMTQLP EAEKEKIAEQVADFKKVSKLDAEIE IWDTSNDIIVLAKKCMIMMEMT DFTRGKGPLKHTTDVYAAKMISESG SRMDVLRQIANQCDDPPCKQDLLA YLEQIKFYSHQLKICSQVKAIEIQLG GELIVSALDSVTSLIQAANKLMNAV QTVKMSYIASTKIIRIQSSAGPRHPVV MWRMKAPAKKPLIKREKPEETWAA VRRGSAKKIHPVQVMSEFRGRQVY	None	None	None	None	None	None	None	None	

0