	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	Q61301	CTNA2_MOUSE	Mus musculus	Catenin alpha-2	27.478975	NaN	T632;S640;S651;S901;S939	29187734;33300544	MTSATSPIILKWDPKSLEIRTLTVERL LEPLVTQVTTLVNTSNKGPSGKKKG RSKKAHVLAASVEQATQNFLEKGEQ IAKESQDLKEELVAAVEDVRKQGET MRIASSEFADDPCSSVKRGTMVRAA RALLSAVTRLLILADMADVMRLLSH LKIVEEALEAVKNATNEQDLANRFK EFGKEMVKLNYVAARRQQELKDPH CRDEMAAARGALKKNATMLYTASQ AFLRHPDVAATRANRDYVFKQVQEA IAGISSAAQATSPTDEAKGHTGIGEL AAALNEFDNKIILDPMTFSEARFRPS LEERLESIISGAALMADSSCTRDDRR ERIVAECNAVRQALQDLLSEYMNNT GRKEKGDPLNIAIDKMTKKTRDLRR QLRKAVMDHISDSFLETNVPLLVLIE AAKSGNEKEVKEYAQVFREHANKLV EVANLACSISNNEEGVKLVRMAATQ IDSLCPQVINAALTLAARPQSKVAQD NMDVFKDQWEKQVRVLTEAVDDIT SVDDFLSVSENHILEDVNKCVIALQE GDVDTLDRTAGAIRGRAARVIHIINA EMENYEAGVYTEKVLEATKLLSETV MPRFAEQVEVAIEALSANVPQPFEE NEFIDASRLVYDGVRDIRKAVLMIRT PEELEDDSDFEQEDYDVRSRTSVQT EDDQLIAGQSARAIMAQLPQEEKAKI AEQVEIFHQEKSKLDAEVAKWDDSG NDIIVLAKQMCMIMMEMTDFTRGK GPLKNTSDVINAAKKIAEAGSRMDK LARAVADQCPDSACKQDLLAYLQRI ALYCHQLNICSKVKAEVQNLGGELIV SGTGVQSTFTTFYEVDCDVIDGGRAS QLSTHLPTCAEGAPIGSGSSDSSML DSATSLIQAAKNLMNAVVLTVKASY VASTKYQKVYGTAAVNSPVVSWKMK APEKKPLVKREKPEEFQTRVRRGSQ KKHISPVQALSEFKAMDSF	None	None	None	None	None	None	None	None	None