

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P30999	CTND1_MOUSE	Mus musculus	Catenin delta-1	25.367067	S47	S4;S47;T59;T112;S125;T217;T221;S225;T228;S230;S252;T257;S268;S269;T280;S288;T291;S300;T304;S320;S346;S349;S352;S617;S713;S811;S847;S857;S859;S861;S864;T865;S868;T869;S879;S899;T904;T906;T916;S920	34418053;20665223;37453647	MDDSEVESTASILASVKEQEAQFEK LTRALEEERRHVSAQLERVRVSPQD ANSLMANGTLRRHQNGRFVGDAD LERQKFSDLKLNQPQDHNHLLYSTI PRMQEPGQIVETYTEEDPEGAMSVV SVETDDGTTTRRTETTIVKVKVTMT TRTVQPVPMGPDGLPVDASAVSNNY IOTLGRDFRKNNGGPGPYVGQAGT ATLPRNFHYPPDGYGRHYEDGYPPG SDNYGSLSRVTRIEERYRPSMEGYR APSRQDVYGPQPQVRVGGSSVDLHR FHPEPYGLEDDQRSMGYDDLIDYGM MSDYGTARRTGTSPDRRRLRSYED MIGEEVPPDQYYWAPLAQHERGSLA SLDSLKGMPPPSNWRQPELPEVIA MLGFRLDAVKSNAAYLQHLCYRN DKVKT DVRKLGIPILVGLLDHPKKE VHLGACGALKNISFGRDQDNKIAIK NCDGVPALVRLLRKARDMDLTEVIT GTLWNLSSHDSIKMEIVDHALHALT DEVIIPHSGWEREPNEDCKPRHIEW ESVLTNTAGCLRNVSSERSEARRKL RECDGLVDALIFIVQAEIGQKDSK LVENCVCLLRNLSYQVHREIPQAER YQEALPTVANSTGPHAASCFGAKKG KDEWFSRGKKPTEDPANDTVDFPKR TSPARGYELLFQPEVVRIYISLLKESK TPAILEASAGAIQNLCAGRWTYGRYI RSALRQEKALSAIAELLTSEHERVVK AASGALRNLAVDARNKELIGKHAIP NLVKNLPGGQQNSSWNFSEDTVVSI LNTINEVIAENLEAAKKLRETQGIEK LVLINKSGNRSEKEVRAAALVLQTIW GYKELRKPLEKEGWKKSDFQVNLN NASRSQSSHSYDDSTLPLIDRNQKS DKKPDREEIPMSNMGSNTKSLDNN YSTLNERGDHNRTLDRSGDLGDME PLKGAPLMQKI