

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O35927	CTND2_MOUSE	Mus musculus	Catenin delta-2	56.546553	S307;S319;S320;T329;T337;S340;S437;T447;S453	S7;S264;S273;S324;S357;S412;S458;S511;T513;S1087;S1098	22645316;22826440;22517741;36852467;33300544;19458039;16452088;35822049;26192747;34678516	MFARKQSGAAPFGAMPVVPDQPPSAS EKNSSLSPLNNTSNGDGSETETTTSAI LASVKEQELQFERLTRELEAERQIVA SQLERCKLGSETGSMSSISSAGEQF HWQTQDQKQDIEDELTTGLELVDSC IRSLQESGILDPQDYSTSERPSLLSQS ALQLNSKPEGSFQYPASYHSNQTLA LGD TAPS QLPARSTQARAAGQSFSQ GTTGRAGHLAGSEPAPPPPPPREPFA PSLGS AFHLPDAPPAAAAALYSSSTL PAPPRGGSPLTTTQGGSPTKLQRGG SAPEGAAAYAAPRGSSPKQSPSRLAKS YSTSSPINIVVSSAGLSPIRVTSPTV QSTISSSPIHQLSSTIGTYATLSPTKR LVHASEQYSKHSQELYATATLQRPG SLAAGSRASYSSQHGHLAPELRALQ SPEHHIDPIYEDRVYQKPPMRSLSQS QGDPLPPAHTGTFRTSTAPSSPGVDS VPLQRTGSQHGPQNAATAFQRASY AAGPASNYADPYRQLQYCASVDS PYSKSGPALPPEGLTARSPSIDSIQKDP REFGWRDPELPEVIQMLQHQPFSV QSNAAAALQHL CFGDNKIKAEIRRO GGIQLLVDDLDRMTEVHRSACGAL RNLVYGKANDDNKIALKNCGGIPAL VRLLRKTTDL EIRELVTGVLWNLSSC DALKMP IIQDALAVLTNAV IIPHS GWENSPLQDDRKIQLHSSQVLRNATGC LRNVSSAGEEARRRMRECDGLTDAL LYVIQSALGSSEIDSKTVENCVCILR NLSYRLAAETSQGGHMGTDDEL DGLL CGETNGKDTESGCGWKKKKKKK QDQWDG VGPLPDCAEPPKGIQMLW HPSIVKPYLTLLECSNPDTLEGAAG ALQNLAAAGSWKGAEDVAGMAYAL RSLPEGAPCLPQWSVYIRAAVRKEK GLPILVELLRIDNDRVCAVATALRN MALDVRNKELIGKYAMRDLVHRLPG GNNSNNSGSKAMSDDTVTAVCCTL HEVITKNMENAKALRDAGGIEKLVG ISKSKGDKHSPKVKAASQVLNSMW QYRDLRSLYKKGWSQYHFVASSST IERDRQRPYSSRTPSISPVVSPNN RSASAPASPREMISLKERKTDYESAG NNATYHGTKEHTSRKDTMTAONT GVSTLYRNSYGAPAEDIKQNQVSTQ PVPQEPSRKDYETYQFPNSTRNYD ESFFEDQVHHRPPASEYTMHLGLKS TGNVDFYSAARPYSELNYETSHYPA SPDSWV