

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q02248	CTNB1_MOUSE	Mus musculus	Catenin beta-1	36.579648	NaN	S23;S29;S33;S37;T41;S45;T64;T142;S191;S246;T331;T333;S552;T556;S675	33300544;29391154;34678516;34346496;22645316;11959983	MATQADLMELDMAMEPDRKAAVS HWQQQSYLDSGIHSGATTTAPSLSG KGNPEEEDVDTSQVLYEWEQGFSQ SFTQEQVADIDGQYAMTRAQVRVRAA MFPETLDEGMQIPSTQFDDAAHPTNV QRLAEPSQMLKHAVVNLINYQDDAE LATRAIPELTKLLNDEDQVVVNKAAV MVHQLSKKEASRHAIMRSPQMVSIAI VRTMQNTNDVETARCTAGTLHNLS HHREGLLAIFKSGGIPALVKMLGSPV DSVLFYAITTLHNLLHQEGAKMAV RLAGGLQKMVALLNKTNVKFLAITT DCLQILAYGNQESKLIILASGGPQAL VNIMRTYTYEKLLWTTSRVLKVLVSV SSNKPAIVEAGGMQALGLHLTDPSQ RLVQNCLWTLRNLSDAATKQEGME GLLGTLVQLLGSDDINVTCAAGILS NLTCNNYKNKMMVCQVGGIEALVR TVLRAGDREDITEPAICALRHLTSRH QEAEMAQNAVRLHYGLPVVVKLLH PPSHWPLIKATVGLIRNLALCPANHA PLREQGAIPRLVQLLVRAHQDTQRR TSMGGTQQQFVEGVRMEEIVEGCT GALHILARDVHNRIVIRGLNTIPLFV QLLYSPIENIQRVAAGVLCELAQDKE AAEAIEAEGATAPLTELHLSRNEGVA TYAAAVLFRMSEDKPQDYKKRLSVE LTSSLFRTEPMAWNETADLGLDIGA QGEALGYRQDDPSYRSFHSGGYGQ DALGMDPMEHEMGGHHPGADYP VDGLPDLGHAQDLMDGLPPGDSNQ LAWFDTDL