

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q5U302	Q5U302_RAT	Rattus norvegicus	Catenin alpha-1	23.439218	NaN	NaN	38843836	MTAVHTGNINFKWDPKSLEIRTLAV ERLLEPLVTQVTTLVNTNSKGPSNK KRGRSKKAHVLAASVEQATENFLEK GDKIAKESQFLKEELVAAVDDVRKQ GDLMKSAAGEFADDPCSSVKRGNM VRAARALLSAVTRLLILADMADVYKL LVQLKVVEDGILKLRNAGNEQDLGI QYRALKPEVDKLNIMAARKQOQLKD VGHRDQMAAARGILQKNVPILYAS QACLQHPDVAAAYKANRDLYKQLQQ AVTGISNAAQATASDDASQQQGGGG GGGELAYALNNFDKQIIVDPLSFSEE RFRPSLEERLESIISGAALMADSSCT RDDRREIVAECAVNRQALQDLLSE YMGNAGRKERSDALNSAIDKMTKK TRDLRRQLRKAVMDHVSDFLETN VPLLVLIEAAKNGNEKEVKEYAQVFR EHANKLIEVANLACSISNNEEGVKL VRMSASQLEALCPQVINAAALAAK PQSKLAQENMDLFKEQWEKQVRVL TDAVDDITSIDDFLAVSENHILEDVN KCVIALQEKDVGDLDRTAGAIRGRAA RVIHVVTSEMDNYEPGVYTEKVLEA TKLLSNTVMPRFTEQVEAAVEALSS DPAQPMDENEFIDASRLVYDGIRDIR KAVLMIRTPEELDDSDFETEDFDVR SRTSVQTEDDQLIAGQSARAIMAQLP QEQAQKIAEQVASFQEEKSKLDAEV SKWDDSGNDIIVLAKQMCMIMMEM TDFTRGKGPLKNTSDVISAAKKIAEA GSRMDKLGRTIADHCPDSACKQDLL AYLQRIALYCHQLNCSKVKAEVQNL GGELVVGVDSSAMSLIQAAKNLMN AVVQTVKASYVASTKYQKSQGMASL NLPVSWKMKAPKPLVKREKQD ETQTKIKRASQKKHVNVPVQALSEFK AMDSI	False	True	3.205	2.477	1.144	0.862	0.938	5.0	1.23