| | 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** | P26233 | CTNB1_XENLA | Xenopus laevis | Catenin beta-1 | 26.176668 | NaN | NaN | 15389870 | MATQADLMELDMAMEPDRKAAVS<br>HWQQQSYLDSGIHSGATTTAPSLSG<br>KGNPEDEDVDTNQVLYEWEQGFSQ<br>SFTQDQVADIDGQYAMTRAQRVRAA<br>MFPETLDEGMQIPSTQFDSAHPTNV<br>QRLAEPSQMLKHAVVNLINYQDDAE<br>LATRAIPELTKLLNDEDQVVVNKAAV<br>MVHQLSKKEASRHAIMRSPQMVSAI<br>VRTMQNTNDVETARCTAGTLHNLS<br>HHREGLLAIFKSGGIPALVKMLGSPV<br>DSVLFYAITTLHNLLLHQEGAKMAV<br>RLAGGLQKMVALLNKTNVKFLAITT<br>DCLQILAYGNQESKLIILASGGPQAL<br>VNIMRTYSYEKLLWTTSRVLKVLSVC<br>SSNKPAIVEAGGMQALGLHLTDSSQ<br>RLVQNCLWTLRNLSDAATKQEGME<br>GLLGTLVQLLGSDDINVVTCAAGILS<br>NLTCNNYKNKMMVCQVGGIEALVR<br>TVLRAGDREDITEPAICALRHLTSRH<br>QEAEMAQNAVRLHYGLPVVVKLLH<br>PPSHWPLIKATVGLIRNLALCPANHA<br>PLREQGAIPRLVQLLVRAHQDTQRR<br>TSIGGTQQQFVEGVRMEEIVEGCTG<br>ALHILARDIHNRIVIRGLNTIPLFVQL<br>LYSPIENIQRVAAGVLCDVAQDKEAA<br>EAIEAEGATAPLTELLHSRNEGVATY<br>AAAVLFRMSEDKPQDYKKRLSVELT<br>SSLFRTEPMPWNEAADLGLDIGAQG<br>EPLGYRQDDSSYRSFHAAGYGQDA<br>MGMDSMMDHDMGGHHPGADYPV<br>DGLPDLSHAQDLMDGLPPGDSNQL<br>AWFDTDL | None | None | None | None | None | None | None | None | None |