	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	extracellu region
0	Q8IWY9	CDAN1_HUMAN	Homo sapiens	Codanin-1	33.321647	S203	T71;S265;S285	30620550;32574038;40307207	MAAVLESLLREEVSVAAVVRWIARS TQGSEDNAGEAAALSSLRALRKEFV PFLLNFLREQSSRVLPQGPPTPAKTP GASAALPGRPGGPPRGSRGARSQLF PPTEAQSTAAEAPLARRGGRRRGPG PARERGGRGLEEGVSGESLPGAGGR RLRGSGSPSRPSLTLSDPPNLSNLEE FPPVGSVPPGPTGTKPSRRINPTPVS EERSLSKPKTCFTSPPISCVPSSQPSA LDTSPWGLGLPPGCRSLQEEREMLR KERSKQLQQSPTPTCPTPELGSPLPS RTGSLTDEPADPARVSSRQRLELVAL VYSSCIAENLVPNLFLELFFVFQLLT ARRMVTAKDSDPELSPAVLDSLESP LFQSIHDCVFFAVQVLECHFQVLSN LDKGTLKLLAENERLLCFSPALQGR LRAAYEGSVAKVSLVMPPSTQAVSF QPETDNRANFSSDRAFHTFKKQRD VFYEVLREWEDHHEEPGWDFEKGL GSRIRAMMGQLSAACSHSHFVRLF QKQLLQMCQSPGGAGGTVLGEAPD VLSMLGADKLGRLWRLQERLMAPQ SSGGPCPPPTFPGCQGFFRDFILSAS SFQFNQHLMDSLSLKIQELNGLALP QHEPNDEDGESDVDWQGERKQFAV VLLSLRLLAKFLGFVAFLPYRGPEPP PTGELQDSILALRSQVPPVLDVRTLL QRGLQARRAVLTVPWLVEFLSFADH VVPLLEYYRDIFTLLLRLHRSLVLSQ ESEGKMCFLNKLLLLAVLGWLFQIP TVPEDLFFLEEGPSYAFEVDTVAPEH GLDNAPVVDQQLLYTCCPYIGELRKL LASWVSGSSGRSGGFMRKITPTTTT SLGAQPSQTSQGLQAQLAQAFFHNQ PPSLRRTVEFVAERIGSNCVKHIKAT LVADLVRQAESLLQEQLVTQGEEGG DPAQLLEILCSQLCPHGAQALALGR EFCQRKSPGAVRALLPEETPAAVLSS AENIAVGLATEKACAWLSANITALIR REVKAAVSRTLRAQGPEPAARGERR GCSRACEHHAPLPSHLISEIKDVLSL AVGPRDPDEGVSPEHLEQLLGQLGQ TLRCRQFLCPPAEQHLAKCSVELAS LLVADQIPILGPPAQYRLERGQARRL LHMLLSLWKEDFQGPVPLQLLLSPR NVGLLADTRPREWDLLLFLLRELVE KGLMGRMEIEACLGSLHQAQWPGD FAEELATLSNLFLAEPHLPEPQLRAC ELVQPNRGTVLAQS	True	True	4.551	5.0	0.281	0.281	False	4.703	False