	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golg app
0	Q05A80	CAPR2_MOUSE	Mus musculus	Caprin-2	41.84854	T375;S378;S634;S638;S644;S797;S873;T882	S852;S853	21540332;36852467;34678516;22645316	MKSAKSQVNQDQQGENQRALSPLQ STLSSAASPSQAYETYIDNGLICLKH KIRNIEKKKLKLEDYKDRLKNGEQL NPDQLEAVEKYEEVLHNLEFAKELQ KTFSALSQDLLKAQKKAQRREHMLK LETEKKKLRTMLQIQYVLQNLTQEH VQKDFKGGLNGAMYLPSKELDYLIK FSKLTCPERNESLSVEDQMEQSSLY FWDLLEGSEKTVVGTTYKHVKDLLS KLLHSGYFESVPVLRNSKEKAEEVL MQSEMKKQLLKSESIKESESLTELV QPEIQPQEFLNRRYMTEVKFSRKQE NVEQSWEADYARKPSLLKCWNTLP EPDGQEKKKESLESWKSSLKTQEVS KPVVSLVQGKLRPTLQEEQKQQVPIT PVSQWKPESPKSKVGSPQEEQNVQE TPKPWVVQSQKEQDPKKLPPGSWA VSVQSEQSGSRSWTTPVCREQASVQ PGTPVSWENNAENQKHSLVPQSQIS LKSWGAASAGLLPNGQVLTRKLNVE PKDVPKPLPQPIDSSSALPKDPVLRK EKLQDLMSQIQGTYNFMQESVLDF DKPSSAIPSSQPPSACPVSTVSAEQN LSNQSDFLQEPSQASSPVTCSSNAC LVTTDQASSGSETEFTTSETPEMVVS PCKPKPASALASPNPPLSKSFQLPPA SGSSEAISTAPFQAMQTVFNVNAPL PPRKEQEMKEPPYSSGYNQNFTSSS TQTVSQCQLPAVHIDQTTQPPETGA GYHPDGTVQVSNGSLAFYPAPTSMF PRPAQPFISSRGTLRGCSHGGRLLM SSYQSPGGYKGFDSYRGLPSVSSGN YSQLQLQAREYSGTAYSQRDNFQQC YKRSGTSSGLQANSRAGWSDSSQVS SPERDSETFNSGDSGLGDSRSMTPV DVPVTSPAAAILPVHIYPLPQQMRVA FSAARTSNLAPGTLDQPIVFDLLLNN LGETFDLQLGRFNCPVNGTYVFIFH MLKLAVNVPLYVNLMKNEEVLVSAY ANDGAPDHETASNHAVLQLLQGDQI WLRLHRGAIYGSSWKYSTFSGYLLY QD	None	None	None	None	None	None	Non