

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	extrac region
089001	CBPD_MOUSE	Mus musculus	Carboxypeptidase D	24.474581	NaN	T264;S269;S1355;S1358;T1365;T1367	33300544	MASGRDERPPWRLGRLRLPPPLL LLLLLLRSSQAHHKKAETTIVG GSEAAEGQFDHYHHEAALGEALEAA AAAGPPGLARLFSIGSSVEGRPLWVL RLTAGLGPPTAAAGLDAAGPLLPGR PQVKLVGNMHGDETVSRQVLVYLA RELASGYRRGDPRLVRLNTTDVYLL PSLNPDGFERAREGDCGLGDSGPPG TSGRDNSRGRDLNRSFPDQFSTGEP PSLDEVPVRALIDWIRRNKFLVLSGN LHGGSVVASYPFDDSPHKTGLYS KTSDDDEVFRYLAKAYASNHPIMKTG EPHCPGDEDETFKDGITNGAHWYD VEGGMQDYNYVWANCFEITLELSC CKYPPASQLRQEWENNRESLITLIEK VHIGIKGFVKDSVTGSGLENATISVA GINHNITGRFGDFHRLVPGTYNL TALSTGYMPLTINNIMVKEGPATEM DFSLRPTVM SVMPGSTEAVTTPGTV AVPNIPPPTSSHQPIQPKDFHHHH FPDMEIFLRRFANEYPNITRYSGLK SVESRELYVMEISDNPGVHEPGEPE FKYIGNMHGNEVVGRELLNLIEYL CKNFGTDPEVTDLVRSTRIHLMPSM NPDGYEKSQEGDSISVVGRRNNSN FDLNRNFPDQFVPIPETQPETIAVM SWVKAYPFVLSANLHGGSLVNNYPY DDNEQGVATYSKSPDDAVFQIALS YSKENSQMFGQRPCKDMYLNIEYFP HGITNGASWYNVPGGMQDWNYLQT NCFEVTIELGCVKYPFENELPKYWE QNRRLIQFMKQVHQVKGFLVDA TDGRGILNATLSVAEINHPVTYKAG DYWRLLVPGTYKITASARGYNPVTK NVTVRSEGA VQVNFVLRVRSADANN ESKKGGRGHSTDDTSDPTSKEFEA LIKHLAENGLEGFMLSSSDLALYR YHSYKDLSEFLRGLVMNYPHITNLT TLGQSVYRHIWSLEISNKPNISEPE EPKIRFVAGIHGNAPVGTPELLALAE FLCLNYKRNPVVTQLVDRTRIVIVPS LNPDRERAQEKDCTSKTGHTNAH GKDLDTDFTSNASQPETKAIHENLIQ KQDFLSIALDGGSLVVTYPYDKPVQ TVENKETLKHASLYANNHPMSMHM GQSPCPNNSDENIPGGVMRGAEWH SHLGSMDKYSVTYGHCPETIVYTSC CYFPSAAQLPALWAENKKSLLSMLV EVHKGVHGLVKDKAGKPIKAVIVL NEGKIVYTKEGGYFHVLLAPGVHNI NAIADGYQQHTQVFVHDAASSVV IVFDTDNRIFGLPRELVVTVSGATMS ALILTACIIWCISIKSNRHKDGFHRL RQHHDEYEDEIRMMSTGSKKLLS HEFQDETDTEETLYSSKH	True	False	1.578	4.238	0.784	1.181	4.362	3.762	3.702