

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	extracell region
Q9JHW1	CBPD_RAT	Rattus norvegicus	Carboxypeptidase D	23.439218	NaN	T265;S270;S1356;S1359;T1366;T1368	38843836	MASGWDERPPWRLESLRLLPPPPLL LLLLLRSSAQAAHIKKA EATTTTVG GSEAAEGQFDHYHHEAALGEALEAA AAAGPPGLARLFSIGNSV EGRPLWV LRLTAGLGPPTPAAVGLDAAGPLLP GRPQVKLVGNMHGDETVSRQVLVY LARELASGYRRGD PRLVRLNTTDV YLLPSLNP DGFERAREGDCGLGDSG PPGTSGRDNSRGRDLNRSFPDQFST GEPPLDEVPEVRALIDWIRRNKFVL SGNLHGGSVVASYPFDD SPEHKTG IYKTSDDDEVFRYLAKAYASNHPIMR TGEPHCPGDEEETF KDGITNGAHWY DVEGGMQDYNVWANC FEITLELS CCKYPPASQLRQEWENNRESLITLIE KVHIGIKGFVKDSVTGSGLENATISV AGINHNITGRFGDFHRLIPGSYNL TAVSPGYMPLTINNIVVKEGPATEID FSLQPTVMSVVPDSTEAVTTPGTAVV PNIPPGT PSSHQPIQPKDFHHHFFP DMEIFLRRFANEYPNITRYSLGKSV ESRELYVMEISDNPCVHEPGEPEFK YIGNMHGNEVVGRELLNLIEYLCK NFGTDPEVTDLVRSTRIHLMPSMNP DGYEKSQEGDSISV VGRNNSNFDL NRNFPDQFVPIITDPTQPETIAMSWS VKAYPFVLSANLHGGSLV VNYPYDD NEQQVATYSKSPDDAVFQQIALSYSK ENSQMFQGRPCDMYLN EYFPHGI TNGASWYNVPGGMQDWN YLQTN C FEVTIELGCVKYPF EKELPKYWEQN RRSLIQFMKQVHQGVKGFVLDATDG RGILNATLSVAEINH PVTTYKAGDYW RLLVPGTYKITASARGYNPVTKNVTV RSEGAIQVNFTLVRSSTDANNESKK GKGHSTDDTSDPTSKEFEALIKHL SAENLEGFMLSSSDLALYRYHSY KDLSEFLRGLVMNYPHITNLTTLQ SVEYRHIWLSLEISNKPNISEPEEKIR FVAGIHGNAPVGT ELLALAEFLCLN YKKNPVVTQLVDRTRIVVPSLNP DG RERAQEKDCTSKTGHTNARGRDL D TDFTSNASQPETKAIENLIQKQDFS LSIALDGGSVLVTYPYDKPVQTVENK ETLKHLASLYANNHPSMHMGQPS C PNNSDENIPGGVMRGAEWHS HLGS MKDYSVTYGHCP EITVYTS CCYFP SA AQLPALWAENKKSLLSMLVEVHKG VHGLVKDKTGKPI SKAVIVLNEGIRV HTKEGGYFHVLLAPGVHNNIAIDG YQQQHSQVFVH HDAASSVIVFDTD NRIFGLPRELVTVSGATMSALILTA CIWCICSIKSNRHKDGFHRLRQHH DEYEDEIRMMSTGSKSLLSHEFQD ETDTEETLYSSKH	True	True	1.1	5.0	0.981	1.204	5.0	3.709	3.642