

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region
Q5NBX1	COBL_MOUSE	Mus musculus	Protein cordon-bleu	27.404006	S926	S47;S50;S212;S235;S272;S294;S346;S349;S372;T522;S649;S816;S1038;S1128;S1303	36852467;34678516	MDAPRALAAKPTGRKMKARAPPPP GKPAAQNVHSEQKLPHDATLGSQQ SLVYMKEALQNSTLDITVVLPSGLEK QSVVSGSHAMMDLLVELCLQNHNL PSHHVLEIWSSETTQPLSFKPNTLIG SLNVHTVLLKEKVPPEERVKPLTKA PEKSVRLVVNYLRTQKAVRVSPPEVP LQNILPVICAKCEVNPPEHVILLRDNV AGEELELSKSLNELGKELYAWDNR REMFKSSSLGNDETDEKKEKFLGFF KANKRSNSKAEHLGLSGADSDPA KSASGGDLNGCVTTTPNSPLHSRSL TLGPSLSLGNISGVMKSDMKKRRRA PPPPSPKLLGQDKVSEKASLSSQADL QKKRRAPAPPPPPQPPPSFVVPNR KEDKEENRKSTVGVGRQVQKPPRG TARGPPQLVLP PPPPPYPPD TDVTEP VTFPGEGAGSETSELRPKLSLPLGPG SHCSMGGVSQVPAESEETASEDTTE DSGVMSSPSDAISLDSQODSMRSKD KWSTDQEDGSDQDLAGTPELGPKQ SPSWGKSGSGSSILRTEKATMPTND DEDLFTGHQHQTAELEDELEGME ENYETDTSLLTNSVNGVSNHSLQEA IIPDSGVDDIPVTFIGEVSDEPFDSGL FSSRCNATTFTNGSIASQRSHLSPS QTEHSQPFVRTSRKEPDPSPSQDN RKRNOPTLANTS ENENPVETDPTVT SLVSKLLIDDPKAKDKGVHGSSHS EKTQAGHGINSRLRVNPRDGKDESSN SAPPWWSHHQALGGSYGLKYGLTT YKIVPPKSEMRCYDRDVSSTGAIKI DELGNLVSPHMGSRTPSPSAVVE TDTPIGKVKFWRNRSMKYLNGP AECTIKRAPSTTTATPEKPPQDNGM KAAFTVTTTQQQPASQEYGAHLEEE RSRPQSAVSCSVKVPASNPTDITFLK PQRRTSSQYVASAIKKMGPPKVHA DVVRPHKATTEQCHEEAKLARSPT RKDDAAPNLHSEARQHEHGTNQSS VCLPSNPGVQLPAGGHPKVEVNSTY GKSSQDYPAAVHRNSYFLPGRSSH RDRVSVGQSCGFNEKQTTSNQKAN STSNFSQALDKAHPPPLLLAEARDS GRILMNGSARTPGNCEPPHSPKEST LTSYIILQTEEEKPSSLSTDGQDADDT LPSSIFGPKKKFKPVIQRPLPKDVSL HSALMEAIHSSGGREKLRKTAEQTS EGRPKPSYVEAESERSALLAAIRGH SGTSLRKRVSLSAEELQSFRNAALG APGLDKPQEDLGLPPPALPPPPAP APQAPSASVTVSFRSTGTSPNSVNR QALMDAIRSGTGAARLRKVPLLV	False	False	1.472	3.35	0.992	0.555	0.518	4.282	0.83