

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	extracellular region
0 Q06890	CLUS_MOUSE	Mus musculus	Clusterin	25.849477	S329	S132;S395	36852467;37453647;36288343	MKILLLCVALLIWDNGMVLGEQEV SDNELQELSTQGSRYNKIEIQNAVQ GVKHIKTLEKTNAERKSLLSLEEA KKKKEDALEDRDSEMKLKAFPEVC NETMMALWEECKPCLKHTCMKFYA RVCRSGLVGGQLEEFNLQSSPFY FWMNGDRIDSLLES DRQQSQVLDA MQDSFARASGIIDTLFQDRFFAREL HDPHYFSPIGFPHKRPFLYPKSRLV RSLMSPSHYGPPSFHNMFQPFEMI HQAQQAMDVQLHSPAFQFPDVFDFL REGEDDRTVCKEIRRNSTGCLKMKG QCEKCEILSVDCSTNNPAQANLRQ ELNDSLQVAERL TEQYKELLQSFQS KMLNTSSLEQLNDQFNWVSQLAN LTQGEDKYYLRVSTVTHSSDSEVPS RVTEVVVKLFDSDPITVVLPEEVSKD NPKFMDTVAEKALQEYRRKSRAE	True	True	5.0	5.0	5.0	4.585	1.958	2.751	4.766