

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q91W50	CSDE1_MOUSE	Mus musculus	Cold shock domain-containing protein E1	43.758632	NaN	S123;S276;S514;S584;T761	28135057;40684658;28528544;25153642;22645316	MSFDPNLLHNNHNGYNGTSAAL RETGVIEKLLTSYGFQCSEERQARLF FHCSQYNGNLQDLKVGDDVEFEVS SDRRTGKPIAIAIKLVKIKPEIHPEERM NGQVVCAPPHNLESKSPAAPGQSPT GSVCYERNGEVFLTYTSEDVEGNV QLETGDKINFVIDNNKHTGAVSARN IMLLKKKQARCQGVVCMKEAFGFI ERGDVVKEIFFHYSEFKGDLETLPQ GDDVEFTIKDRNGKEVATDVRLLPQ GTVIFEDISIEHFEGTVTKVIPKVPSK NQNDPLPGRIKVDVFKELPFGDKD TKSKVILLEGDHVRFNISTDRRDKL ERATNIEVLSNTFQFTNEAREMGVI AAMRDGFGFIKCVDRDARMFFHFS EILDGNQLHIADEVEFTVVPDMLSA QRNHAIKIKLPGTVSFHSHSDHR FLGTVEKEATFSNPKTTSPNKGKDK EAEDGHHAYDDCGVKLTIAFQAKDVE GSTSPOIGDKVEFSISDKQRPQQIA TCVRLGRNSNSKRLGYVATLKDN FGFIETANHDKIEFFHYSEFSGDVDS LELGDMVEYSLSKGGKGNKVAEKVN KAHSVNGITEANPTIYSGKVIRPLR GVDPTQIEYQGMIEIVEEGDMKGEV YPPGIVGMANKGDCLQKGESVKFQL CVLGQNAQTMAYNITPLRRATVECV KDQGFINYEVGDSKLLFFHVKEVQ DGVELQAGDEVEFSVILNORTGKCS ACNVWRVCEGPKAVAAPRPDRLVN RLKNITLDDASAPRLMVLQRPRGPD NSMGFGAERKIRQAGVID	True	False	3.0	False	4.056	1.397	2.0	2.394