	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membr
0	Q9P2I0	CPSF2_HUMAN	Homo sapiens	Cleavage and polyadenylation specificity factor subunit 2	15.178176	S460	S419;S420;S423;S660	23301498;37217939;30379171;40596516;36240223;40914422;35008409;34019948	MTSIIKLTTLSGVQEESALCYLLQVD EFRFLLDCGWDEHFSMDIIDSLRKH VHQIDAVLLSHPDPLHLGALPYAVG KLGLNCAIYATIPVYKMGQMFMYDL YQSRHNTEDFTLFTLDDVDAAFDKI QQLKFSQIVNLKGKGHGLSITPLPAG HMIGGTIWKIVKDGEEEIVYAVDFN HKREIHLNGCSLEMLSRPSLLITDSF NATYVQPRRKQRDEQLLTNVLETLR GDGNVLIAVDTAGRVLELAQLLDQI WRTKDAGLGVYSLALLNNVSYNVVE FSKSQVEWMSDKLMRCFEDKRNNP FQFRHLSLCHGLSDLARVPSPKVVL ASQPDLECGFSRDLFIQWCQDPKNS IILTYRTTPGTLARFLIDNPSEKITEIE LRKRVKLEGKELEEYLEKEKLKKEA AKKLEQSKEADIDSSDESDIEEDIDQ PSAHKTKHDLMMKGEGSRKGSFFK QAKKSYPMFPAPEERIKWDEYGEIIK PEDFLVPELQATEEEKSKLESGLTN GDEPMDQDLSDVPTKCISTTESIEIK ARVTYIDYEGRSDGDSIKKIINQMKP RQLIIVHGPPEASQDLAECCRAFGGK DIKVYMPKLHETVDATSETHIYQVRL KDSLVSSLQFCKAKDAELAWIDGVL DMRVSKVDTGVILEEGELKDDGEDS EMQVEAPSDSSVIAQQKAMKSLFGD DEKETGEESEIIPTLEPLPPHEVPGH QSVFMNEPRLSDFKQVLLREGIQAE FVGGVLVCNNQVAVRRTETGRIGLE GCLCQDFYRIRDLLYEQYAIV	True	False	1.184	5.0	1.008	0.682	False	1.091