	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac regior
0	Q922K7	NOP2_MOUSE	Mus musculus	28S rRNA (cytosine-C(5))-methyltransferase	25.685855	NaN	S59;S68;T169;S179;S649;S713;S767;S782;S793	39627609	MGRKLDPTKKEKRGPGRKARKQKGAETELVRFLPAAGDENSKRLSSRARKRAAKRRAGSVDVPKPNKSPGIKTLPGELSKGAVQARGKKRPAPIQNSDGDEEEDSGEDDVVTQGDLWGSEDSDEDMVDDYGAASNSEDEEEKLLPIERAALKQKAQDATAGVLWNEEDTDEDEDDDGVSPESHPRKDDKAEGDLQINVEDEEAFVLPPAGETDQDGQAPDLQRVHKRIQDIVGVLRDFGAQREEGRSRAEYLSRLQKDLATYYSYGDFLLSKLMELFPLSELIEFLEANEVPRPITLRTNTLKTRRRDLAQLLINRGVNLDPLGKWSKSGLVVYDSSVPIGATPEYLAGHYMLQGASSMLPVMALAPQEHERILDMCCAPGGKTSYIAQLMKNTGVILANDANADRLKSVVGNLHRLGVTNTIISHYDGRQFPKVVGGFDRVLLDAPCSGTGVISKDPAVKTNKDEKDIQRCAHLQKELLLSAIDSVNAASKTGGYLVYCTCSITVEENEWVVDYALKKRNVRLVPTGLDFGQEGFTRFQARRFHPTLRSTRRFYPHTHNMDGFFIAKFKKFSNSIPQPHAGNSAAATPTEPDLKDQVTPKSENGSQPTKKARGAVKAKQQLLRQPHSKKPFQKLNGIAKGPGLSTEPSVPDAQVSTRPSQSAGNADVNSKRKRSEKLKQRGPKWKPSKEAAVPKPSAPSRVEDSGTPVPTPSEIRAAPRPKDCAPSLGKAKKKQKGKQQLAQQPANGAAPLKEDAVSKGPSAPFVSPHSSTRPPPAKRRKSMTKGNSQPLLS	True	False	2.33	4.726	1.471	0.715	0.344	0.742	1.067