

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
Q8R3F9	STPAP_MOUSE	Mus musculus	Speckle targeted PIP5K1A-regulated poly(A) polymerase	25.685855	NaN	S688;S744	39627609	MAAVSDVVS LPRGRFRCCLCDVTT ANRPSLDAHLKGRKHRDLVQLRATR KAQGLRSV FVSGFPRDVGSAQLSEY FQTFGPVANIVMDKDKGVFAIVEMG DISAREAVLSQPKHSLGGHGLRVRP REQKEFQSPASKSPKGV DSSSHQLV QALAEAADVGAQMVKLVRELSEEA ERQLRNLVVALMQEVFTEFFPGCVV HPFGSTVNSFDVHGCDLDFLDMG DMEETEPDPKAPKVPETSSLDSALA SSLDPQALACTPASPLDSLPTSVQE SESLDFDTPSSLAPQTPDSALGSDTV TSPQSLPPVSP LQEDRKEGKQGKEL ELAE EASKDEKEEAAVLELVGSILR GCVPGVYRVQTVPSARRPVVKFCHR PSGLHGDVSLSNRLALYNSRFLNLC SEMDGRVRPLVYTLRCWAQHNGLS GGGPLLNYYAL TLLVIYFLQTRDPV LPTVAQLTQRAGEGEQVEVDGWD C SFPKDASRLEPSTNVEPLSLLAQFF SCVSCDL DLSGSLSLREGRPLMVAE GLPSDLWEGLRLGPMNLQDPFDLS HNVAANVTGRVAKRLQSCCGAAASY CRSLQYQQRSSRGRDWGLLPLLQPS SPSSLLSAKLIPLPSAPFPQVIMALVD VLREALGCHIEQGTKRRRSEGARIKD SPLGGVNRQRRLGGQEKSFEEGKEE PQGCAGDHSENEVEEMVIEVRETPQ DWALLHSGPPEEELPLMTANCLDK AAEHNPMKPEVAGEGSQGETGKEA SHPSSVSWRCALWHQVWQGRRA RRRLQQQTKEEGRGGPTTGAEWLA MEARVTQELKGPNSEQERPPGPELL SFVASASQAEQTLTVAPLQDSQGLFP GLHHFLQGFIPQALKNLLK	True	False	3.1	4.591	1.081	0.881	0.696	0.971	0.701