	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	extracellular region
0	Q6DID3	SCAF8_MOUSE	Mus musculus	SR-related and CTD-associated factor 8	26.23264	S745;T750;S819	T6;S273;S617	36852467;34887587	MEAVKTFNSELYSLNDYKPPISKAK MTQITKAAIKAIKFYKHVVQSVEKFI QKCKPEYKVPGLYVIDSIVRQSRHQF GQEKDVFAPRFSNNIISTFQNLYRCP GDDKSKIVRVLNLWQKNNVFKSEII QPLLDMAAGIPPPVVTPVLASTTAAM SNTPGTPVTPVTPANVVQGLPDPWV SQIANTDTLAAVAQILQSPQGQQLQ QLIQTLQIQQQKPQPSILQALDAGLV VQLQALTAQLTAAAAAANTLTPLDQ GVSFNKKLMDRFDFGEDSEHSEES KKEMPTPQLSHVSESVNNSIFHQIA EQLQQQNLEQLRQQLLEQQQPQKV TPQDSQEGTFGSEHSASPSQGSSQQ HFLEPEANLDDSIDIQQQDMDIDEG QDVVEEEIFEPEAKKVAVRSRSRTHS RSRSRSPRKRRSRSRSGSRKRKHRK RSRSHSREKKRKASRSYSSERRARE REKERQKKGLPPVRSKTLSVCSTTL WVGQVDKKATQQDLTNLFEEFGQIE SINMIPPRGCAYVCMVHRQDSFRAL QKLSSGSYKIGSKVIKIAWALNKGVK TEYKQFWDVDLGVTYIPWEKVKVDD LDGFAEGGMIDQETVNAEWETVKA SEPVKEPVQTAQSPAPVEKESVVTTQ AEVFPPPVAMLQIPVAPAVPAVSLVP PAFPVSMPVPPPGFNPIPPPPFLRAS FNPSQPPPGFMPPPVPPPVVPPPAIP PVVPTSLVQPPLSMTPEAVKDVGFG SLVLPSGSVAGSLAPSTLPAGNVFNP PSKAEPEEKVPHLIEHQIPSGENTRP VIPSDIPSSAAMLAQPPGASSTSGILC VQRPNVSSNSEILGVRPANVSNSAAI MGAQPPNILNNSGILAIQPPNVSSGS GLLGVLPPNLPNNSGLVGLQPPNVT SPAGLLGTQPPIGPQNLPPLAIPAQR MPALPMLDIRPGLIAQAPGPRFPLLQ PGIPPQRGIPPPSVLDAALHPPPRGP FPPGDLFSQPERPFLAPGRPSIDNVP NPDKRIPLGNDNIQQEGDRDYRFPPI ETREGITRPPQVDVRDVVGRRLDPR EGPGRPPLDARDHFGRPPVDMREN LVRPSLDHLGRRDHFGFPPEKPWGP RDFDEREHRVLPVFGGPKGLHEERG RFRAGNYRFDPRSGPWNRGFGQEV HRDFDDRRRPWERQRDRDDRDFDF CREINGNRLGRDRIQNTWVPPPHAR VFDYFEGATSQRKGDNVPQVNGEN TERHAQPPPLPVQKDPELYEKLASS GDVDKEESGTVAGVESEAVVESTET EGT	None	None	None	None	None	None	None	None	None