	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endopl reticul
0	Q9UPN6	SCAF8_HUMAN	Homo sapiens	SR-related and CTD-associated factor 8	23.545102	S89;T95;S605;T611;T614;T615;S617;T619;S686;S779;T807;S821;S822;S824;S861	T6;S273;T615;S617;S779	28657654;31492838;23301498;37340703;34019948;35254053;28510447;29351928;20068230;35138101;30620550;30059200;30379171;22661428;35132862;33214551;35289036;34725712;27114449;38665916	MEAVKTFNSELYSLNDYKPPISKAK MTQITKAAIKAIKFYKHVVQSVEKFI QKCKPEYKVPGLYVIDSIVRQSRHQF GQEKDVFAPRFSNNIISTFQNLYRCP GDDKSKIVRVLNLWQKNNVFKSEII QPLLDMAAGIPPPVVTPVLASTTTAM SNTPGTPVTPVTPANVVQGLPDPWV SQITNTDTLAAVAQILQSPQGQQLQQ LIQTLQIQQQKPQPSILQALDAGLVV QLQALTAQLTAAAAAANTLTPLEQG VSFNKKLMDRFDFGEDSEHSEEPK KEIPASQLSHVSESVNNSIFHQIAEQ LQQQNLEHLRQQLLEQQQPQKATP QDSQEGTFGSEHSASPSQGSSQQHF LEPEVNLDDSIDIQQQDMDIDEGQD GVEEEVFEQEAKKVAVRSRSRTHSR SRSRSPRKRRSRSRSGSRKRKHRKR SRSRSRERKRKSSRSYSSERRARER EKERQKKGLPPIRSKTLSVCSTTLWV GQVDKKATQQDLTNLFEEFGQIESI NMIPPRGCAYVCMVHRQDAFRALQ KLSSGSYKIGSKVIKIAWALNKGVKT EYKQFWDVDLGVTYIPWEKVKVDDL EGFAEGGMIDQETVNTEWETVKSSE PVKETVQTTQSPTPVEKETVVTTQAE VFPPPVAMLQIPVAPAVPTVSLVPPA FPVSMPVPPPGFSPIPPPPFLRASFN PSQPPPGFMPPPVPPPVVPPPTIPPV VPTSLVQPSLSMTPETVKDVGFGSL VIPGGSVASNLATSALPAGNVFNAPT KQAEPEEKVPHLIDHQISSGENTRSV IPNDISSNAAILGGQPPNVTSNSGIL GVQRPNVSSNSEILGVRPSNVSSSS GIIAAQPPNILNNSGILGIQPPSVSNS SGLLGVLPPNIPNNSGLVGVQPPNV PNTPGLLGTQPPAGPQNLPPLSIPNQ RMPTMPMLDIRPGLIPQAPGPRFPLI QPGIPPQRGIPPPSVLDSALHPPPRG PFPPGDIFSQPERPFLAPGRQSVDNV TNPEKRIPLGNDNIQQEGDRDYRFP PIETRESISRPPPVDVRDVVGRPIDPR EGPGRPPLDGRDHFGRPPVDIRENL VRPGIDHLGRRDHFGFNPEKPWGH RGDFDEREHRVLPVYGGPKGLHEER GRFRSGNYRFDPRSGPWNRGFGQE VHRDFDDRRRPWERQRDRDDRDFD FCREMNGNRLGRDRIQNTWVPPPH ARVFDYFEGATSQRKGDNVPQVNG ENTERHAQPPPIPVQNDPELYEKLTS SNEINKEKSDTVADIESEPVVESTET EGT	None	None	None	None	None	None