

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9UPN6	SCAF8_HUMAN	Homo sapiens	SR-related and CTD-associated factor 8	32.23989	S89;T95;S605;T611;T614;T615;S686;S821;S822;S824;S861	T6;S273;T615;S617;S779	3626323;30379171;33214551;20068230;27114449;31492838;30059200;22661428;28657654;29351928;34019948;30620550;23301498;28510447	MEAVKTFNSELYSLNDYKPPISKAK MTQITKAAIKAIKFKYKHVVQSVKEFI QKCKPEYKVPGLYVIDSIVRQSRHQF GQEKDVFAPRFSNNIISTFQNLRYRCP GDDKSKIVRVLNLWQKNNVFKSEII QPLLDMAAGIPPPVTPVLASTTTAM SNTPGTPVTPVTPANVVQGLPDPWV SQITNTDTLAAVAQILQSPQGQQLQQ LIQTLQIQQQKQPQPSILQALDAGLVV QLQALTAQLTAAAAAANTLTPLEQG VSFNKKLMDRFDGDESEHSEEPK KEIPASQLSHVSESVNNSIFHQIAEQ LQQQNLHLRQQLLEQQQPQKATP QDSQEGTFGSEHSASPSQSSQOHF LEPEVNLDDSIDIQQDMIDIDEGQD GVVEEVFEQEAKKVAVRSRSTRHSR SRSRSPRKRSSRSRSGSRKRKRHRKR SRSRERKRKSSRSYSERRARER EKERQKKGLPPIRSKTLSVCSTTLWV GQVDKKATQDDLNLFEFQIESI NMIPPRGCAYVCMVHRQDAFRALQ KLSSGSYKIGSKVIAWALNKGVKT EYKQFWDVDLGVYIPWEKVKVDDL EGFAEGGMIDQETVNTWETVKSSE PVKETVQTTQSPTPVEKETVVTQAE VFPPPVAMLQIPVAPAVPTVSLVPPA FPVSMVPPPGFSPIPPPFLRASFN PSQPPPGFMPPVPPVPPPTIPPV VPTSLVQPSLSMTPETVKDVGFGSL VIPGGSVASNLATSALPAGNVFNAPT KQAEPEEKVPHLIDHQISSGENTRSV IPNDISSNAAILGGQPPNVTNSNGIL GVQRPNVSSNSEILGVRPSNVSSSS GIIAQPPNILNNSGILGIQPPSVSNS SGLLGVLPPNIPNNSGLVGVQPPNV PNTPGLLGTQPPAGPQNLPLSIPNQ RMPTMPMLDIRPGLIPQAPGRFPLI QPGIPPQRGIPPPSVLDSALHPPPRG PFPFGDIFSQPERPFLAPGRQSVDNV TNPEKRIPLGNDNIQEGDRDYRFP PIETRESISRPPVDVRDVVGRPIDPR EGPGRPPLDGRDHFGRPPVDIRENL VRPGIDHLGRRDHFGFNPEKPWGH RGDFDEREHRVLPVYGGPKGLHEER GRFRSGNYRFDPRSGPWNRGFGQE VHRDFDDRRRPWERQRDRDRDFD FCREMNGNRLGRDRIQNTWVPPPH ARVFDYFEGATSQRKGDNVQVNG ENTERHAQPPPIPVQNDPELYEKLTS SNEINKEKSDTVADIESEPVESTET EGT