

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	25.321177	S89;T95;S605;T611;T614;T615;S617;S686;S779;T807;S821;S822;S824;S861	T6;S273;T615;S617;S779	23301498;30379171;31492838;35254053;22661428;29351928;27114449;28657654;28510447;30059200;33214551;35132862;20068230;34725712;35289036;35138101;34019948;30620550	MEAVKTFNSELYSLNDYKPPISKAK MTQITKAAIKAIKFYKHHVQSVKEFI QKCKPEYKVPGLYVIDSIVRQSRHQF GQEKDVFAPRFSNNIISTFQNLRYRCP GDDKSKIVRVLNLWQKNNVFKSEII QPLDMAAGIPPPVTPVPLASTTTAM SNTPGTPVTPVTPANVVQGLPDPWV SQITNTDTLAAVAQILQSPQGGQLQO LIQTLQIQQQKQPQPSILQALDAGLVV QLQALTAQLTAAAAAANTLTPLEQG VSFNKKLMDFDFGEDSEHSEEPK KEIPASQLSHVSESVNNSIFHQIAEQ LQQNLEHLRQQLLEQQQPQKATP QDSQEGTFGSEHSASPSQGSSQOHF LEPEVNLDDSIDIQQDMIDIDEGQD GVVEEVFEQEAKKVAVRSRSTRHSR SRSRSPRKRSSRSRSGSRKRKRHRKR SRSRERKRKSSRSYSERRARER EKERQKKGLPPIRSKTLCVSTTLWV GQVDDKATQDDLTLNLFEEFGQIESI NMIPPRGCAYVCMVHRQDAFRALQ KLSSGSYKIGSKVIAWALNKGVKT EYKQFWDVDLGVTYIPWEKVKVDDL EGFAEGGMIDQETVNTWETVKSSE PVKETVQTTQSPTPVEKETVVTQAE VFPPPVAMLQIPVAPAVPTVSLVPPA FPVSMVPPPGFSPIPPPFLRASFN PSQPPPGFMPPVPPVPPPTIPPV VPTSLVQPSLSMTPETVKDVGFGSL VIPGGSVASNLATSALPAGNVFNAPT KQAEPEEKVPHLIDHQISSGENTRSV IPNDISSNAAILGGQPPNVTNSNGIL GVQRPNVSSNSEILGVRPSNVSSSS GIIAQPPNILNNSGILGIQPPSVSNS SGLLGVLPPNIPNNSGLVGVQPPNV PNTPGLLGTQPPAGPQNLPLSIPNQ RMPTMPMLDIRPGLIPQAPGRFPLI QPGIPPQRGIPPPSVLDSALHPPPRG PFPFGDIFSQPERPFLAPGRQSVDNV TNPEKRIPLGNDNIQEGDRDRYRFP PIETRESISRPPVDVRDVVGRPIDPR EGPGRPPLDGRDHFGRRPPVDIRENL VRPGIDHLGRRDHFGFNPEKPGWH RGDFDEREHRVLPVYGGPKGLHEER GRFRSGNYRFDPRSGPWNRGFGQE VHRDFDDRRRPWERQRDRDRDFD FCREMNGNRLGRDRIQNTWVPPPH ARVFDYFEGATSQRKGDNVQVNG ENTERHAQPPPIPVQNDPELYEKLTS SNEINKEKSDTVADIESEPVESTET EGT