

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96T23	RSF1_HUMAN	Homo sapiens	Remodeling and spacing factor 1	3.245925	NaN	S227;S392;S397;S429;S473;S524;S570;S604;S622;T628;S629;S748;S882;S1096;S1098;S1105;S1221;S1223;S1226;S1258;S1277;T1278;T1305;S1325;S1336;S1345;S1359;S1375	30379171;34019948	MATAAAAAVMAPPGCPGSCPNFAV VCSFLERYGPLLDLPELPPPELRLV QAPPPDVNGEVPKELVELHLKLMR KIGKSVTADRWEEKYLKICQEFNSTW AWEMEKKGYLEMSVECKLALLKYL CECQFDDNLKFKNIINEEDADTMRL QPIGRDKDGLMYWYQLDQDHNVR MYIEEQDDQDGSSWKCIVRNRNEL AETLALLKAQIDPVLLKNSSQODNSS RESPSLEDEETKKEEETPKQEEQKE SEKMKSEEQPMDLENRSTANVLEE TTVKKEKEDEKELVKLPVIVKLEKPL PENEEKKIIKEESDSFKENVKPIKVE VKECRADPKDTKSSMEKPVAQEPER IEFGNIKSSHEITEKSTEETEKLKN DQQAQIPLKKREIKLSDDFDSPVKGP LCKSVTPTKEFLKDEIKQEEETCKRI STITALGHEGKQLVNGEVSDERVAP NFKTEPIETKFYETKEESYSPSKDRN IITEGNGTESLNSVITSMKTGELEKE TAPLRKDADSSISVLEIHSQKAQIEEP DPPEMETSLSSEMAKDLSSKTALS STESCTMKGEEKSPKTKKDKRPPILE CLEKLEKSKKTFLDKDAQRLSPIPEE VPKSTLESEKPGSPEAAETSPPSNIID HCEKLASEKEVVECQSTSTVGGQSV KKVDLETLKEDSEFTKVEMDNLDN AQTSGIEEPSETKGSMQKSKFKYKL VPEEETTASENTEITSERQKEGIKLT RISSRKKKPDSPPKVLEPENKQEKTE KEEKTNVGRTLRRSPRISRPTAKVA EIRDQKADKKRGEDEVEEESTAL QKTDKKEILKSEKDTNSKVSKVKP KGKVRWTGSRTGRWKYSSNDESE GSGSEKSSAASEEEEEKESEEAILAD DDEPCKKCGLPNHPELILLCDSCDS GYHTACLRPPLMIIPDGEWFCPPCQ HKLLCEKLEEQLQDLVALKKKERA ERRKERLVYVGIENIIPPQEPDFSE DQEEKKKDSKSKANLLERRSTRTR KCISYRFDEFDEAIDEAIEDDIKEADG GGVGRGKDITITGHRGKDITILDE ERKENKRPQAAAARRKRRRLNDL DSDSNLDEEESDEFKISDGSQDEF VSDENPDESEEDPPSNDDSDTDFC SRRLRRHPSRPMRQSRRLRRKTPKK KYSDDDEESEESENSRDESDFSDD FSDDFVETRRRRSRRNQKRQINYKE DSESDGSQKSLRRGKEIRRVHKRRL

