

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8TAQ2	SMRC2_HUMAN	Homo sapiens	SWI/SNF complex subunit SMARCC2	23.524013	NaN	S283;S286;S302;S304;S306;S347;S387;T548;S813	34105348;22121020;21740066;33214551;34019948;27655845;34725712;32119511;35138101	MAVRKKDGGPNVKYEEAADTVTQF DNVRLWLWGKNYKKYQAEPTNKSL SSLVVQLLQFQEEVFGKHVSNAPLT KLPIKCFLDKAGGSLCHILAAAYKF KSDQGWRRYDFQNPSPMDRNVEM FMTIEKSLVQNNCLSRPNIFLCPEIE PKLLGKLDIIKRHQGTVTEDKNNNA SHVVYPVPGNLEEEEWVRPVMKRD KQVLLHWGYYPDSYDTWIPASEIEAS VEDAPTPEKPRKVHAKWILDTDTFN EWMNEEDYEVNDDKNPVSRRKKIS AKTLTDEVNSPDSDRRDKKGGNYKK RKRSPSPSPTPEAKKNAKKGPSTPY TKSKRGHREEEQEDLTKDMDEPSPV PNVEEVTLPKTVNTKKDSEAPVKG GTMTDLDEQEDESMTTGKDEDEN STGNKGEQTKNPD LHEDNVTEQTH HIIIPSYAAWFDYNSVHAIERRALPEF FNGKNKSKTPEIYLAYRNF MIDTYRL NPQEYLTSTACRRNLAGDVCAIMRV HAFLEQWGLINYQVDAESRPTMGP PPTSHFHVLA DT P SGLVPLQPKTPQ QTSASQQMLNFPDKGKEKPTDMQN FGLRTDMYTKKNVPSKSKAAASATR EWTEQETLLLLLEALEMYKDDWNKV SEHVGSR TQDECILHFLRLPIEDPYL EDSEASLGPLAYQPIPF S QSGNPVM STVAFLASVVDPRVASAAKSALEEF SKMKEEVPTALVEAHVRKVEEAAKV TGKADPAFGLESSGIAGTTSDEPERI EESGNDEARVEGQATDEKKEPKEPR EGGGAIEEEAKEKTSEAPKKDEEK KEGDSEKESKSDGDPVDPEKEKE PKEGQEEVLKEVVESEGERKTKVER DIGEGNLSTAAAAALAAA AVKAKHL AAVEERKIKSLVALLVETQMKKLEIK LRHFEELETIMDREREAL EYQROQL LADRQAFHMEQLKYAEMRARQQHF QQMHQQQQQPPALPPGSQIPPTG AAGPPAVHGLAVAPASVVPAPAGSG APPGSLGPSEQIQAGSTAGPQQQQ PAGAPQPGAVPPGVPPPGPHGSPFP NQQTPPSMMPGAVPGSGHPGVAGN APLGLPFGMPPPPPPAPSIIPFGSLA DSISINLPAPPNLHGHHHHLPFAPG TLPPP NLPVSMANPLHPNLPATTTM PSSLPLGPGLSAAAQSPAIVAAVQG NLLPSASPLPDPGTPLPPDPTAPSPG TVTPVPPPQ