

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q92922	SMRC1_HUMAN	Homo sapiens	SWI/SNF complex subunit SMARCC1	16.264	T603	S310;S328;S330;T335;S350;S357;T398;S573;S822;S825	33214551;32119511;33465208;20305658;21740066;34019948;23301498;28510447	MAAAAGGGPGTAVGATGSGIAAAA AGLAVYRRKDGGPATKFWESPETVS QLDSVRVWLKGHYKYYHADAPT KTLAGLVVQLLQFQEDAFGKHVTNP AFTKLPACFMDFKAGGALCHILGA AYKYKNEQGWRRFDLQNP SRMDRN VEMFMNIEKTLVQNNCLTRPNYILIP DIDLKLANCLKDIIKRHQGTFTDEKS KASHHIYPYSSSQDDEEWLRPVMRK EKQVLVHWGFYPSYDTWVHSNDV DAEIEDPPIPEKPWKVHVKWILDTDI FNEWMNEEDYEVDENRKPVSFRQR ISTKNEEPVRSERRDRKASANARK RKHSPSPPPPTPTESRKKSGKKGQA SLYGKRRSQKEEDEQEDLTKDMED PTPVPNIEEVVLPKNVNLKKDSENTP VKGGTVADLDEQDEETVTAGGKEDE DPAKGDQSRVLDGEDNVTEQTNHI IIPSYASWFDYNCIHVIERRALPEFFN GKNKSKTPEIYLAYRNF MIDTYRLNP QEYLTSTACRRNLTGDVCAVMRVHA FLEQWGLVNYQVDPE SRPMAMGPP PTPHFNVLADTPSGLVPLHLRSPQV PAAQQMLNFPEKNKEKPVLDLQNF LRTDIYSKKTAKSKGASAGREWTE QETLLLLLEALEMYKDDWNKVSEHV GSRTQDECILHFLRLPIEDPYLENSD ASLGPLAYQVPFVSQSGNPVMSTVA FLASVVDPRVASAAAKAAL EEF SRVR EEVPLELVEAHVKKVQEAARASGKV DPTYGLESSCIAGTGPDEPEKLEGAE EEKMEADPDGQQPEKAENKVENET DEGDKAQDGENEKNSEKEQDSEVS EDTKSEEKETEENKELDTCKERES DTGKKKVEHEISEGNVATAAAAAALA SAATKAKHLAAVEERKIKSLVALLVE TQMKKLEIKLRHFEELETIMDREKE ALEQQRQQLLTERQNFHMEQLKYA ELRARQQMEQQQHGNPQQAHQH SGGPGLAPLGAAGHPGMMPHQQPP PYPLMHHQMPPPHPPQPGQIPGPGS MMPGQHMPGRMIPTVAANIHPSGS GPTPPGMPPMPGNILGPRVPLTAPN GMYP PPPQPPPPPPADGVPPPPA PGPPASAAP