

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P97496	SMRC1_MOUSE	Mus musculus	SWI/SNF complex subunit SMARCC1	25.685855	NaN	S309;S327;S329;T334;S349;S356;T397;S572;S775;S821;S824	39627609	MAATAGGGPGAAGAVGAGGAAAA SGLAVYRRKDGGPASKFWESPDTVS QLDSVRVWLKHKYKYYHADAPT KTLAGLVVQLLQFQEDAFGKHVTNP AFTKLPKACFMDFKAGGTLCILGA AYKYKNEQGWRRFDLQNP SRMDRN VEMFMNIEKTLVQNNCLTRPNILIP DIDLKLANLKDIIKRHQGTFTDEKS KASHHIPPSSQEEDEEWRPVMRR DKQVLVHWGFYPSYDTWVHSNDV DAEIEDAPIPEKPKVHVWILDTDV FNEWMNEEDYEV DENRKPVSFRQR ISTKNEEPVRSERRDRKASANSRK RKPSPSPPTATESRKKSGKKGQAS LYGKRRSQKEEDEQEDLT KDMEDPT PVPNIEEVLKPNVNPKKDSENTPV KGGTVADLDEQDEAVTTGGKEDED PSKGDPSRSVDPGEDNVTEQTNHIII PSYASWFDYNCIHVIERRALPEFFNG KNKSKTPEIYLAYRNF MIDTYRLNPQ EYLTSTACRRNL TGDVCAVMRVHAF LEQWGLVNYQVDPESRPMAMGPPP TPHFNVLADTPSGLVPLHLRSPQVP AAQQLNFPKKNKEKPIDLQNFGLR TDIYSKKT LAKSKGASAGREWTEQE TLLLEALEMYKDDWNKVSEHVG RTQDECILHFLRLPIEDPYLENSDAS LGPLAYQVPFQSQGNPVMSTVAFL ASVVDPRVASAAAKALEEFSRVREE VPLELVEAHVKVQEAARASGKVDV TYGLESSCIAGTGPDEPEKLEGESEE KMETDPDGGQPEKAENKVENESDE GDKIQDRENEKTEKEQSDVSESDV KPEEKENEENKELTDTCKERESDAG KKKVEHEISEGNVATAAAAALASAAT KAKHLAAVEERKIKSLVALLVETQM KKLEIKLRHFEELETIMDREKEALEQ QRQQLTERQNFHMEQLKYAELRA RQQMEQQQHGQTPQQAHOHTGG PGMAPLGATGHPGMMPHQPPYP LMHHQMPPPHPPQPGQIPGPGSMM PGQMPGRMIPAVAANIHTGSGPT PPGMPPMPGNILGPRVPLTAPNGMY PPPPQQQPPPPADGVPPPPAPGPPA SATP	True	False	2.215	5.0	1.394	0.814	False	1.216	1.253