

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
Q6PDG5	SMRC2_MOUSE	Mus musculus	SWI/SNF complex subunit SMARCC2	25.685855	NaN	S283;S286;S302;S304;S306;S347;S387;S813	39627609	MAVRKKDGGPNVYKYYEAADTVTQF DNVRLWLQGNKYKQIQAEPPTNKSL SSLVVQLLQFQEEVFGKHVSNAPLT KLPIKCFLDKAGGSLCHILAAAYKF KSDQGWRRYDFQNPSPRMDRNVEM FMTIEKSLVQNNCLSRPNIFLCPDIE PKLLGKLDIVKRHOGTISEDKSNAS HVVYPVPGNLEEEWVRPVMKRDK QVLLHWGYPPDSYDTWIPASEIEASV EDAPTPEKPRKVHAKWILDTDTFNE WMNEEDYEVSDDKSPVSRKKISAK TLTDEVNSPDSRRRDKKGGNYKRRK RSPSPSPTPEAKKNAKKGSTPYTK SKRGHREEEQEDLTKDMDEPSVPP NVEEVTLPKTVNTKDSSESAPVKGK TMTDLDEQDDESMETTGKDEDENS TGNKGEQTKNPDLHEDNVTEQTHH IIIPSYAAWFDYNSVHAIERRALPEFF NGKNKSKTPEIYLAYRNFIMIDTYRL NPQEYLTSTACRRNLAGDVCAIMRV HAFLEQWGLINYQVDAESRPTPMGP PPTSHFHVLAADTPSGLVLPQKPPQ QSSASQQMLNPFPEKGEKPADMQN FGLRTDMYTKKNVPSKSKAAASATR EWTEQETLLLLALEMYKDDWNKV SEHVGSRTQDECILHFLRLPIEDPYL EDSEASLGPLAYQIPFSQSGNPVM STVAFSLVVDPRVASAAKSALEEF SKMKKEVPTALVEAHVRKVEEAQV TGKADPAFGLLESSGIAGTASDEPERI EESGTEEARPEGQAADKKEKPEPR EGGGAVEEEAKEISEVPPKDEEKG KEGDSEKESEKSDGPIVDPEKDKE PTEGQEEVLKEVAEPEGERKTKVER DIGEGLNSTAAAAALAAAVKAKHL AAVEERKIKSLVALLVETQMKKLEIK LRHFEELETTIMDREREALYQRQQL LADRQAFHMEQLKYAEMRARQQHF QQMHQQQQQPPTLPPGSQPIPTG AAGPPTVHGLAVPPAAVASAPPGSGA PPGSLGFSEIQGAGTTAGPQQPQQ AGAPQPGAVPPGVPPGPHGPSFPF NQPTPPSMMPGAVPGSGHPGVAGN APLGLPFGMPPPPAAPSVPFGSLA DSISINLPPPNLHGHHHHPFAPG TIPPPNLVPSMANPLHHPNLPTTTM PSSLPLPGGLGSAQAQSPAIVAQVQ NLLPSASPLPDGPTLPPDPTAPSPG TVTPVPPQ	True	False	2.397	5.0	1.265	0.884	False	1.123	1.102