

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	es
Q91ZW3	SMCA5_MOUSE	Mus musculus	SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily A member 5	23.620502	NaN	T55;S65;T112;S115;S136;S170;S754	34887587;37507081	MSSAVEPPPPPPESAPSKPSAAGAG GSSSGNKGPEGGAAPAAPCAAGSG PADTEMEEVFDHGS PGKQKEIQEPD PTYEEKMQTDRANRFEYLLKQTELF AHFIQPAAQKTPTSPLKMKPGRPRV KKDEKQNLSSVGDYRHRRTQEED EELLTESSKATNVCTRFEDSPSYVK WGKLRDYQVRGLNWLISLYENGIN ILADEMGLGKTLQTISSLGYMKHYR NIPGPHMVLVPSKSTLHNWMSEFKK WVPTLRVSVCLIGDKEQRAAFVRDVL LPGEDWVCVTSYEMLIKEKSVFKKF NWRYLVIDEAHRKNEKSKLSEIVRE FKTTNRLLLTGTPLQNNLHELWSLL NFLLPDVFNASADDFDSWFDTNACL GDQKLVERLHMVLRPFLRLRIKADV EKSLPPKKEVKIYVGLSKMQREWYT RILMKDIDILNSAGKMDKMRLLNIL MQLRKCCNHPYLFDAEPGPPYTTD MHLVTNSGKMVLDKLLPKLKEQG SRVLFSQMRVLDILEDYCMWRNY EYCRLDGQTPHDERQDSINAYNEPN STKFFVMLSTRAGGLGINLATADWVI LYDSDWNPQVDLQAMDRHRIGQT KTVRVFRFITDNTVEERIVERAEMKL RLDSIVIQQGRLVDQNLNKGKIDEM LQMIRHGATHVFASKSEITDEDIDG ILERGAKTAEMNEKSKMGESSLR NFTMDTESSVNFEGEDYREKQKIA FTEWIEPPKRERKANYAVDAYFREA LRVSEPKAPKAPRPPKQPNVQDFQF FPPRLFELLEKEILYRKTIGYKVP PDL PNAQAQKEEQKIDAEPLND EELEEKELLTQGFNWNKRDFNQ FIKANEKWGRDDIENIAREVEGKTP EEVIEYSAVFWERCNELQDIEKIMA QIERGEARIQRRISIKKALDTKIGRYK APFHQLRISYGTNKGKNYTEEDRF LICMLHKLGFDKENVYDELRCIRN SPQFRFDWFLKSRTAMELQRCNT LITLIERENMELEKEKAEKKRGPK PSTQKRKMDGAPDGRGRKKLKL	True	False	2.421	5.0	1.963	0.995	0.713	1.717	1.