

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
Q920F6	SMC1B_MOUSE	Mus musculus	Structural maintenance of chromosomes protein 1B	26.222628	NaN	NaN	24788674	MGHLELLLVENFKSWRGRQVIGPFK RFTCIIGPNSGSKSNVMDALSFVMG EKTTNLRVKNIQELIHGAHTGKPVSS SASVTIYIEDSGEEKTFTRIIRGGCSE YHFGDKPVSRSVYVAQLENIGIIVKA QNCLVFQGTVESISMKKPKERTOFF EEISTSGEFIGEYEAKKKKLQKAEED AQPHFNVKKNVAERKHAKIEKEEA EHYQNLLEELKINKIQLMLFQLYYN EEKINVLNTELEQMDGNLSVVKDTL SHHENIFKAKKKDYGMLTRQLQOTA KELKSVEAILNQKRPQYIKAKENTSH HLKKLDLTKKLTIDNEKQCSKQEDG IRALVAELADLDRAWKSFQKMEEK ILQKGRDIELENSQLDRYKLLKEQVR RKVGIMTQOLEKLQWEQKAEKERL AFEKRRHGDGTQGNLKQIKEQIEEHK KRIEKLEEYTKTCMDCLEDDKQQUEE ALKKEIENTKSRMSEVNEELSLIRNE LQNAIDNHEGKRQKRAEVLEHL KRLYPDSVFGRLDLCHPIHKYQLA VTKLFGRYMVAIVVASEKIAKDCIRF LKAERAEPETFLALDYLDIKPINERL REIKGCKMMIDVIKTQFPOLKKVIQF VCGNGLVCETVEEARHIAFGGPERR KAVALDGTFLKSGVISGSSDLKHK ALCWDEKELHNLDRKRSQLVQELK ELMKTLRKETDLKQIOTLVQGTNTR LKYSQNELEMIKKKHLATFYREQSQ LQSELLNIDSQCTMLSEGINKQQQKI EEFQDKIDEVEDDIFQDFCEEIGVEN IREFENKHVKQQQENDQKRFEFK QKTRLNIQLEYSRNQLKKLNNIDT LKTTIQKGKEDIDNLKKTETEECLKIV EELMVKQEIQEVLATQSSNIEKIHI QIEEERKKVLAVDREVGKLQKEVVII QGSLEQKLEKHNLLEDCKVQDIDIS LVLGSLEDIIEMELTETESTQATADIY EKEASIQIDYSPLREDLQALQSDKEV EAHLTLLQQVASQENTLLKTTAPN LRAQENLKTVRDKFQESADVFEASR KEARICRQEFQVRRRYDAFSQCF EHISVSIDQYKCLKRNSAQAFLLSP ENPEPYLDGISYNCVAPGKRFMPM DNLSGGEKCVAAALLLFAVHSFRPA PFFVLDEVDAALDNTNIGKVSSYIKE QSQEQFMIIISLKEEFYSKADALIG VYPEHNECMFSHVLTLDLSKYPDTE DQEGSRSHRKPRVPRVMSPKSPQS R	None	None	None	None	None	None	None		