

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
Q9CW03	SMC3_MOUSE	Mus musculus	Structural maintenance of chromosomes protein 3	27.796441	NaN	T783;S787;S886;S1013;S1065;S1067;S1074;S1083	39627609;34887587	MYIKQVVIQGFRRSYRDQTVDPFSSK HNIVVGRNGSGKSNFFYAIQFVLSD EFSHLRPEQRLALLHEGTGPRVISAF VEIIFDNDNRLPIDKEEVSLRRVIGA KKDQYFLDKKVMVTKNDVMNLLESA GFSRSNPYYIVKQGKINQMATAAPDS ORLKLLEVVAGTRVYDERKEESISL MKETEGKREKINELLKYIEERLHTLE EEKEELAQYQKWDKMRRALEYTIYN QELNETRAKLDELAKRETSGEKSR QLRDAQDARDKMEDIERQVRELK TKISAMKEEKEQLSAERQEQIKQRT KLELKAKDLQDELAGNSEQRKRLK ERQKLEKIEEKQKELAETEPKFNSV KEKEERGIAARLAQATQERTDLYAKO GRGSQFTSKEERDKWIKKELKSLDQ AINDKKRQIAAIHKDLEDTEANKEK NLEQYNKLDQDLNEVKARVEELDR KYYEVKNKKDELQSERNYLWREEN AEQQALAAKREDLEKKQQLLRAATG KAILNGIDSINKVLEHFRKGINQHV QNGYHGIVMNNFECEPAFYTCVEVT AGNRLFYHIVDSDEVSTKILMEFNK MNLPGEVTFPLNKLDRDTAYPET NDAIPMISKLRYNPRFDKAFKHVFG KTLICRSMVSTQLARAFTMDCITLE GDQVSHRGALTGGYYDTRKSRLELQ KDVRKAEFEELGELEAKLNENLRNI ERINNEIDQLMNMQIQIETQQRKF KASRDSILSEMMLKEKRQSEKTF MPKQRSLQSLEASLHAMESTRESLK AELGTDLLSLSLEDQKRVDALENDE IRQLQENRQLLNERIKLEGITRVE TYLNENLRKRLDQVEQELNELRETE GGTVLTATTSELEINKRVKDTMAR SEDLDNSIDKTEAGIKELQKSMERW KNMEKEHMDAINHDTKELEKMTN RQGMMLKKKEECMKKIRELGSQPQE AFEKYQTLCLKQLFRKLEQCNTLKL KYSHVNKKALDOFVNFSEQKEKLIK RQEELDRGYKSIMELMNVLELRKYE AIQLTFKQVSKNFSEVFQKLVPGGK ATLVMKKGDVESQSQDEGEGSGE SERGSGSQSSVPSVDQFTGVGIRVSF TGKQGEMREMQLSGGQKSLVALA LIFAIQKCDPAPFYLFDEIDQALDAQ HRKAVSDMIMELAVHAQFITTTFRP ELLESADKFGYKFRNKVSHIDVITA EMAKDFVEDDITTHG	True	False	2.663	5.0	2.269	1.406	1.116	1.715