

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9CW03	SMC3_MOUSE	Mus musculus	Structural maintenance of chromosomes protein 3	9.340548	NaN	T783;S787;S886;S1013;S1065;S1067;S1074;S1083	34887587	MYIKQVIIQGFRSYRDQTIIVDPFSSK HNVIVGRNGSGKSNFFYAIQFVLSDFSHLRPEQRLALLHEGTGPRVISAF VEIIFDNSDNRLPIDKEEVSRRVIGAKKDQYFLDKKMVTKNDVMNLLESA GFSRSNPYYIVKQGKINQMATAPDS QRLKLLREVAGTRVYDERKEESISL MKETEGKREKINELLKYIEERLHTLEEKEELAQYQKWDKMRRALEYTIYN QELNETRAKLDDELSAKRETSGEKSR QLRDAQDARDKEDIQVRELKTKISAMKEEKEQLSAERQEQIKQRT KLELKAKDLQDELQAGNSEQRKRLKERQKLEKIEEKQKELAETEPKFNVS VKEKEERGIARLAQATQERTDLYAKQGRGSQFTSKEERDKWIKKELSLDQ AINDKKRQIAAIHKDLEDTEANKEKNLEQYNKLDQDLNEVKARVEELDR KYYEVKNKKDELQSERNYLWREENAEQQALAAKREDLEKKQQLLRAATG KAILNGIDSINKVLEHFRRKGINQHVQNGYHGIVMNNFECEPAFYTCVEVT AGNRLFYHIVDSDEVSTKILMEFNKMNLPGEVTFPLNKLVDVDRDTAYPET NDAIPMISKLRYNPRFDKAFKHVFGKTLICRSMEVSTQLARAFTMDCITILE GDQVSHRGALTGGYYDTRKSRLELQKDVRAEEELGELEAKLNENLRRNI ERINNEIDQLMNQMQQIETQQRKFKASRDSILSEMKMLKEKRQQSEKTF MPKQORSLQSLEASLHAMESTRESLKAELGTDLLSQLSLEDQKRVDALNDE IRQLQQENRQLLNERIKLEGIITRVE TYLNENLRKRLDQVEQELNELRETEGGTVLTATTSELEAINKRVKDTMAR SEDLDNSIDKTEAGIKELQKSMERWKNMEKEHMDAINHDTKELEKMTN RQGMLLKKKEECMKKIRELGSPLQEA FEKYQTL SLKQLFRKLEQCNT TELK KYSHVNKKALDQFVNFSEQKEKLIK RQEELDRGYKSIMELMNVLELRKYE AIQLTFKQVSKNFSEVFKLVPGGK ATLVMKKGDVESQSQDEGEGSGE SERGSGSQSSVPSVDQFTGVGIRVSF TGKQGEMREMQQLSGGQKSLVALA LIFAIQKCDPAPFYLFDEIDQALDAQ HRKAVSDMIMELAVHAQFITTTFRP ELLESADKFIGVVKFRNKVSHIDVITA EMAKDFVEDDTTHG

