

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	extrac regio
E9Q2X6	E9Q2X6_MOUSE	Mus musculus	Structural maintenance of chromosomes protein	32.227397	NaN	NaN	34887587;25153642;28528544	MRRKGTKPSTACHQEEGPPPSQDG AHSDEEMEQPAGEAESAAPATMAS EAGAPRLMITHIVNQNFKSYAGEKV LGPFHKRFSCHGPNNGSGKSNVIDSM LFFVGYRAQKIRSKKLSVLHNSDEH KDIQSCTVEVHFQKIIDKEGDDYEVL PNSNFYVSRDAYRSTSVYHISGKKK TFKDVGNLLRSHGIDLDHNRFLILQ GEVEQIAMMKPKGQTEHDEGMLEY LEDHGGRLNEPIKVLRRVEILNE HRGEKLNVRKMVEKEKDALEGEKN IAIEFLTENEMFKKKNHICQYYIYD LQNRIAEITQKEKIHEDTKEITEKS NVLSNEMKAKNSAVKDVEKKLNKV TKFIEQNKEKFTQLDLEDVQVREKL KHATSKAKKLEKQLQKDKKVEELK SVPKSKTVINETTRNNSLEKEREK EEKLKEVMDSLKQETQGLQKEKEI QEKELMGFNKSNEARSKMEVAQS ELDIYLSRHNTAVSLSKAKEALITA SETLKERKAAIKDINTKLPQTQOQLK EKEKELQKLTQOEINLKSIVHDLFQ KVEEAKSSLAMNRSRGKVLDAHQE KKSGRIPGIYGRGLGDLGAIDEKYDIAI SSCCHALDIYVDSIDTAQECVNFLLK KHNIGIATFIGLDKMTVWAKKMSKI QTPENTPRLFDLVKVKNEIRQAFYF ALRDTLVANNLDQATRVAYQRDRR WRVVTLQGOIIEQSGTMSGGGSKV MRGRMGSSVIDEISVEEVNKMESQL ERHSKQAMQIQEQKVQHEEAVVKL RHSERDMRNTLEKFAASIQGLSEQE EYLCVQIKELEANVLTTPDRKQOQL LEENVSVFKKEYDAVAEKAGKVEAEI KRLHNTIIDINNRKLAQQNKLDITIN KQLDECASAITKAQVAIKTADRNLKK AQDSVCRTEKEIKDTEKEINDLKTTEL KNIEDKAEVINNTKTAETSLPEIQK EHRNLLQELKVIQENEHALQKDALS IKLKLQIDGHISEHNSKIKYWQKEI SKIKLHPVEDNPVETVAVLSQEELEA IKNPESITNEIALLEAQCREMKPNLG AIAEYKKKEDLYLQ RVAELDKITSER DNFRQAYEDLRKQRLNEFMAGFYVI TNKLENYQMLTLGGDAELELVDSL DPFSEGIMFSVRPPKKSWKKIFNLS GGEKTLSSLALVFALHHYKPTPLYF MDEIDAALDFKNVSIYAFYIYEQTKN AQFIIISLRNNMFEISDRLLGIYKTYN STKSVAVNPKQIASKGLC	None	None	None	None	None	None	None		