

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
Q8NDV3	SMC1B_HUMAN	Homo sapiens	Structural maintenance of chromosomes protein 1B	18.379595	T815	NaN	30379171	MAHLELLLVENFKSWRGQVIGPFR RFTCIIGPNGSGKSNVMDALSFVMG EKIANLRVKNIQELIHGAHIGKPISS ASVKIYVEESGEEKTFARIIRGGCSE FRFNDNLVSRVYIAELEKIGIIVKAQ NCLVFQGTVESISVKKPKERTQFFEE ISTSGELIGEYEEKRKLQKAEEDAQ FNFNKKKNIAAERRQAKLEKEEAER YOSLLEELKMNKIQLQLFQLYHNEK KIHLNNTKLEHVNRDLSVKRESLSH HENIVKARKEHGMLTRQLQOQTEK ELKSVETLLNQKRPQYIKAKENTSH HLKKLDVAKKSIKSEKQCSKQEDD IKALETELADLDAAWRSFEKQIEEEI LHKKRDIELEASQLDRYKELKEQVR KKVATMTQOLEKLQWEQKTDEERL AFEKRRHGEVQGNLQIKEQIEDHK KRIEKLEEYTKTCMDCLKEKKQEE TLVDEIEKTKSRMSEFNELNLRSE LQNAGIDTHEGKRQKRAEVLEHLK RLYPDSVFGRLFDLCHPIHKYQLAV TKVFGRFITAIIVASEKVAKDCIRFLK EERAEPETFLALDYLDIKPINERLRE LKGCKMVIDVIKTQFPQLKKVIQFVC GNGLVCETMEEARHIALSGPERQKT VALDGTFLKSGVISGGSSDLKYKAR CWDEKELKNLRDRRSQKIQLKGL MKTLRKETDLKQIQTLIQGTQTRLKY SQNELEMIKKHLVAFYQEQSOLQS ELLNIESOCIMLSEGIKERQRRIKEF QEKIDKVEDDIFQHFCEEIGVENIRE FENKHVKRQEQEIDQKRYFYKMLTR LNVQLEYSRSHLKKLNKINTLKETI QKGSSEDIDHLKKAENCLQTVNEL MAKQQQLKDIRVTQNSSAEKVOTQI EEERKKFLAVDREVGLQKEVVSQ TSLEQKRLEKHNLLEDCKVQDIEIIL LSGSLDDHIEVEMGTEAESTQATIDIY EKEEAFEIDYSSLKEDLKALQSDQEI EAHLRLLQQVASQEDILLKTAAPNL RALENLKTVRDKFQESTDAFEASRK EARLCRQEFQVKKRRYDLFTQCFE HVSISIDQIYKKLCRNNSAQAFLSPE NPEEPYLEGISYNCVAPGKRFMPMD NLGGGEKCAALALLFAVHSFRPAP FFVLDEVDAAALDNTNIGKVSSYIKEQ TQDQFMIVISLKEEFYSRADALIGIY PEYDDCMFSRVLTLDLSQYPDTEGQ ESSKRHGESR	None	None	None	None	None	None	None		