

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	exti reg
Q96T76	MMS19_HUMAN	Homo sapiens	MMS19 nucleotide excision repair protein homolog	12.749961	NaN	S1027	34019948;23301498;28510447;38665916	MAAAAAVEAAAPMGALWGLVHDFV VGQEGPADQVAADVKSNGYTVLQ VVEALGSSLENPEPRTRARAIQLLSQ VLLHCHTLLLEKEVVHLLIFYENRLK DHHLVIPSVLQGLKALSLCVALPPGL AVSVLKAIFQEVHVQSLPQVDRHTV YNIITNFMRTREEELKSLGADFTFGF IQVM DGEKDP RNLLVAFRIVHDLISR DYSLGPFEVLEFVETSCYFPIDFTPPP NDPHGIQREDLILSLRAVLASTPRFA EFLPLLLIEKVDSEVLSAKLDSLQTL NACCAVYGQKELKDFLPSLWASIRR EVFQTASERVEAEGLAALHSLTACLS RSVLRADAEDLLDSFLSNILQDCRH HLCEPDMKLVWPSAKLLQAAAGAS ARACDSVTSNVLPILLEQFHKHSQS SQRRITILEMLLGLKLGKQKWSYEDK DQRPLNGFKDQLCSLVMALTD PST QLQLVGIRTLTVLGAQPDLLSYEDLE LAVGHLYRLSFLKEDSQSCRVAALE ASGTLAALYPVAFSSHLVPKLAEELR VGESNLTNGDEPTQCSRHLCCQLQAL SAVSTHPSIVKETLPLLLQHLWQVN RGNMVAQSSDVIACQSLRQMAEK CQQDPESCWYFHTAIPCLLALAVQ ASMPEKEPSVLRKVLLEDEVLAAAMV SVIGTATTHLSPELAAQSVTHIVPLFL DGNVSFLPENSFPRFQPFQDGSSG QRRLIALLMAFVCSLPRNVEIPQLNQ LMRELLELSCCHSCPFSSTAAAKCF AGLLNKHYPAGQQLDEFQLAVDKVE AGLGSGPCRSQAFTLLLVVTKALVL RYHPLSSCLTARLMGLLSDPELGPA AADGFSLMSDCTDVLTRAGHAEVR IMFRORFFTDNVPALVQGFHAAPOD VKPNYLKGLSHVLRNLPKPVLLPELP TLLSLLLEALSCPDCVVQLSTLSCLQ PLLEAPQVMSLHVDTLVTKFLNLS SSPSMAVRIAALQCMHALTRLPTPV LLPYKQVIRALAKPLDDKKRLVRKE AVSARGEWFLLGSPGS	None	None	None	None	None	None	None	None	

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