

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracell region
Q9D071	MMS19_MOUSE	Mus musculus	MMS19 nucleotide excision repair protein homolog	27.735065	NaN	S1028	28135057;28528544;33300544	MAAATGLEEAVAPMGALCGLVQDF VMGQQEGPADQVAADVKSGGYTVL QVVEALGSSLENAEPRTRARGAQLL SQVLLQCHSLLSEKEVVHLILFYENR LKDHHLVVPVSLQGLRALSMSVALP PGLAVSVLKAIFQEVHVQSLQVDR HTVFSIITNFMRSSREEELKGLGADFT FGFIQVMDGEKDPNLLLAFRIVHD LISKDYSLGPFVEELFEVTSCYFPIDF TPPPNDPYGIQREDLILSLRAVLASTP RFAEFLPLLEIKVDSEILSAKLDLQ TLNACCAVYGGKELKDFLPSLWASI RREVFQTASERVEAEGLAALHSLTA CLSCSVLRADAEDLLGSFLSNLQDC RHHLCPEPMKLVWPSAKLLQAAAG ASARACEHLTSNVLPILLEQFHKHS QSNQRRTILEMILGFLKLQKWSYE DRDERPLSSFQDQLCSLVFMALDTP STQLQLVGIRTLTVLGAQPGLLSAED LELAVGHLYRLTFLEEDSQSCRVAAL EASGTLATLYPGAFSRHLLPKLAEEL HKGESDVARADGPTKCSRHFRCQLQA LSAVSTHPSIVKETLPLLQHLQCAN KGNMVTESSEVVAVCQSLQQVAEK CQQDPESYWYFHKTAVPCLFALAVQ ASMPEKESVLRKVLLEDEVALAALA SVIGTATTHLSPELAAQSVTCIVPLFL DGNTSFLPENSFPDQFPFQDSSG QRRVLALLTAFVCSLPRNVEIQNLR LMRELLKQSCGHSCPFSSAATKCF AGLLNKQPPGQLEEFQLAVGTVE AGLASESSRDQAFTLLLWVTKALVL RYHPLSACLTRLMGLSDPELGCA AADGFSLMSDCTDVLTRAGHADVR IMFRORFFTDNVPALVQGFHAAPOD VKPNYLKGLSHVNLRLPKPVLLPELP TLLSLLLEALSCPDSVVQLSTLSCLQ PLLEAPQIMSLHVDTLVTKFLNLSS SYMAVRIAALQCMHALTRLPTSVL LPYKSQVIRALAKPLDDKKRLVRKEA VSARGEWFLGSPGS	None	None	None	None	None	None	None	None	

0