

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
Q9R1X5	MRP5_MOUSE	Mus musculus	ATP-binding cassette sub-family C member 5	25.875327	S1213;T1215	S14;S19;S43;S60;S505;S509;T513	30059200	MKDIDMGKEYIIPSPGYRSDRDRSAV PCQHRDPEEPFRFRTRSLQDALE TAARVEGLSLDISVHSHLQILDEEHS KGKYHHGLSVLKPFRITTKHQHPVD NAGLFSYMTFWSLSPLARVVHKKGE LLMEDVWPLSKYESSDVNSRRRLERL WQEELNEVGPDAASLRVVWFICRT RLILSIVCLMITQLAGFSGPAFVVKH LLEYTOATESNLQYSLLLVLGILLTE VVRSWSLALTWALNYRTGVRLRGAI LTMAFKKILKLNIEKSLGELINICS NDGQRMFEAAAVGSLLAGGPVVAIL GMIYNVILGPTGFLGSAVILFYPAM MFSVRLTAYFRKCVAAATDDRVQK MNEVLTYIKFIKMYAWVKAFSQCVO KIREEERRILEKAGYFQSITVGVAPIV VVIASVVTFSVHMTLGFHMTAAQAFI VTVFNSMTFALKVTPFVSKLSEAS VAVDRFKSLFLMEEVHMINKNPASP HIKIEMKNATLAWDSSHSSIQNSPK LTPKMKDKRATRGKKEKSRQLQH TEHQAVLAEQKGHLLDSDRPSPE EEEGKQIHTGSLRLQRTLYNIDLEIE EGKLVGICGSVGSKTSLSAILGQM TLLEGSIAVSGTFAYVAQQAWILNAT LRDNILFGKEFDEERYNSVLNSCCL RPDLAILPNSDLTEIGERGANLSGGQ RQRISLARALYSRDSRIYLDPLSALD AHVGNHIFNSAIRKRLKSKTVLFVTH QLQYLVDCEVIFMKEGCITERGTH EELMNLNGDYATIFNLLGETPPV EINSKKEATGSQKSQDKGPKPGSVK KEKAVKSEEGQLVQVEEKGGQSV WSVYVVYQAAGGPLAFLVIMVLFM LNVGSTAFSTWWSLYWIKQSGNST VYQGNRSFVSDSMKDNPFMQYYASI YALSMAVMLILKAIKRGVVFVKGTLRA SSRLHDELFRRLRSPMKFFDITPTG RILNRFKDMDEVDVRLPFQAEMFI QNVILVFCVGMIAGVFPWFLVAVG PLLILFSLHIVSRVIRELKRDLNIT QSPFLSHITSSIOGLATIHAYNKQE FLHRYQELDDNQAPFFLFTCAMR WLAVRLDLISIALITTTGLMIVLMHG QIPSAYAGLAISYAVQLTGLFOFTVRL ASETEARFTSVERINHXYIKTSLLEAPA RIKNAKPPHDWPOEGEVTFFENAEM RYRENPLVLKKSFTIKPKKIGIVG RTGSGKSSLGMAFLRVELSGGCIKI DGIRISDIGLADLRSLAIPQEPVLF GTVRSNLDPFNQYTEDQIWDALERT HMKECIAQLPLESEVMENGDNF SVGERQLLCIARALLRHCKILILDEAT AAMDTEDDLQETIREAFADCTMLT IAHRLHTVLGSDRIMVLAQQQVVEF DTPSVLLSNDSSRFYAMFAAANKV AVKG	False	True	1.825	1.973	1.922	1.553	3.504	5.0	1.986