

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P08183	MDR1_HUMAN	Homo sapiens	ATP-dependent translocase ABCB1	8.275682	S419	S660	34019948;33214551;29351928	MDLEGDRNGGAKKKNFFKLNKSE KDKKEKKPTVSVFMSFRYSNWLDKL YMVVGTAAIIHGAGLPLMMLVFGE MTDIFANAGNLEDLMSNITNRSIN DTGFFMNLEEDMTRYAYYYSGIGAG VLVAAYIQVSFWCLAAGRQIHKIRKQ FFHAIMRQEIGWFDVHDVVGELNTRL TDDVSKINEGIGDKIGMFFQSMATF FTGFIVGFTRGWKLTLVILAI SPVLGL SAAVWAKILSSFTDKELLAYAKAGAV AEEVLA AIRTVIAFGGQKELERYNK NLEEAKRIGIKKAITANISIGAAFLLIY ASYALAFWYGTTLVLSGEYSIGQVLT VFFSVLIGAFSVGQASPSIEAFANAR GAAYEIFKIIDNKPSIDSYKSGHKPD NIKGNLEFRNVHFSYPSRKEVKILKG LNLKVQSGQTV ALVGN SGC GKSTTV QLMQRLYDPT EGMVSVDGQDIRTIN VRFLREIIGVVSQEPVLFATTIAENIR YGRENVTMDEIEKAVKEANAYDFIM KLP HKFDTLVGERGAQLSGGQKQRI AIARALVRNPKILLLDEATSALDTESE AVVQVALDKARKGRTTIVIAHRLSTV RNADVIAGFDDGVIVEKGNHDELMK EKG IYFKLV TMQTAGNEVELENAAD ESKSEIDALEMSSNDRSSLIRKRST RRSVRGSQAQDRKLSTKEALDESIPP VSFWRIMKLNLT EWPFVVGVFCAII NGGLQPAFAIIFSKIIGVFTRIDDPET KRQNSNLFSLFLALGIISFITFFLQG FTFGKAGEILTKRLRYMVFRSMLRQ DVSWFDDPKNTTGALTTRLANDAA QVKG AIGSRLAVITQNIANLGTGIIIS FIYGWQLTLLLLAIVPIAIAAGVVEMK MLSGQALKDKKELEGSGK IATEAIE NFRTVVSLTQE QKFEHMYA QSLQVP YRNSLRKAHIFGITFSFTQAMMYFSY AGCFRFGAYLV AHKLMSFEDVLLVF SAVVF GAMAVGQVSSFAPDYAKAKI SAAHIIMIEKTPLIDSYSTEGLMPNT LEGNVTFGEVVFNYPTRPDIPVLQGL SLEVKKGQTLALVGSSGCGKSTVVQ LLERFYDPLAGKVLLDGKEIKRLNVQ WLR AHLGIVSQEPILFDCSIAENIAY GDNSRVVSQEEIVRAAKEANIHFIE SLPNKYSTKVGDKGTQLSGGQKQRI AIARALVRQPHILLLDEATSALDTESE EKVVQEALDKAREGRTCIVIAHRLST IQNADLIVVFQNGRVKEHGTHQQLL

