

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
P98198	AT8B2_HUMAN	Homo sapiens	Phospholipid-transporting ATPase ID	22.903221	S1143;S1151;S1162	S1175	38253038	MTVPKEMPEKWARAQAPPSWSRKK PSWGTTEERRARANDREYNEKFOY ASNCIKTSKYNLTFLPVLNLFQEQE VANTYFLFLILQLIPQISSLSWFTTIV PLVLVLTITAVKDATDDYFRHKSDNQ VNNRQSQVLINGILOQEQWMNVCV GDIIKLENNQFVAADLLLLSSSEPHG LCYIETAELDGETNMKVRQAIPVTSE LGDISKLAKFDGEVICEPPNKLKDF SGTLYWKENKFPPLSNQNMLLRGCV LRNTEWCFGLVIFAGPDTKLMQNSG RTKFKRTSIDRLMNTLVWIFGFVLC MGVILAIIGNAIWEHEVGMRFQVYLP WDEAVDSAFFSGFLSFWSYIILNTV VPISLYVSEVIRLGHSYFINWDKMM FCMKKRTPAEARTTTLNEELGQVEY IFSDKTGTLTQNIMVFNKCSINGHSY GDVFDVLGHKAEGERPEPVDFFSN PLADKKFLFWDPSLLEAVKIGDPHT HEFFRLLSLCHTVMSEEKNEGELYY KAQSPDEGALVTAARNFGFVFRSRT PKTITVHEMGTAITYQLLAILDFNNIR KRMSVIVRNPEGKIRLYCKGADTILL DRLHHSTQELLNTTMDHLNEYAGE GLRTLVLAYKDLDEEYEEWAERRL QASLAQDSREDRLASIEEVENNM MLLGATAIEDKLOQGVPEIALLTLA NIKIWVLTGDKQETAVERNIGYSCKML TDDMTEVFIVTGHTVLEVREELRKA REKMMDSRSVNGFTYQDKLSSS KLTSVLEAVAGEYALVINGHSLAHL EADMELEFLETACACKAVICCRVTP QKAQVVELVKYKKAVTLAIGDGAN DVSMIKTAHIGVIGSQEGIQAVLAS DYSFSQFKFLQRLLLVHGRWSYLRM CKFLCYFFYKNFAFTMVHFWFGFFC GFSAQTVYDQYFITYNIVYTSPLVLA MGVFDQDVPEQRSMEYPKLYEPGQ LNLLFNKREFFICIAQGIYTSVLMFFI PYGVFADATRDDGTQLADYQSFVIT VATSLVIVSVQIGLDTGYWTAINHF FIWGLAVYFAILFAMHSNGLFDMF PNQFRFVGNAAQNTLAQPTVWLTIVL TTVVCIMPVVAFRFLRLNPKDLSDT VRYTQLVRKKQKAQHRCMRRVGRV GSRRSYAFSHQEGFCELIMSGKN MRLSSLALSSFTTRSSSWIESLRRK KSDSASSPSGGADKPLKG	None	None	None	None	None	None	None		