	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasm memb
0	Q6P542	ABCF1_MOUSE	Mus musculus	ATP-binding cassette sub-family F member 1	26.69368	NaN	S22;S90;S103;S107;S138;S194;S225;S587	37453647;39627609;34887587	MPKGPKQQPPEPEWIGDGEGTSPAD KVVKKGKKDKKTKKTFFEELAVEDK QAGEEEKLQKEKEQQQQQQQQKKK RDTRKGRRKKDVDDDSDERVLMER LKQLSVPASDEEDEVPAPIPRGRKKA KGGNVFEALIQDDSEEEEEEEENRV LKPAKPEKNRINKAVAEEPPGLRSKK GKEEKSKGKAKSKPAAADSEGEEEE EDTAKEKEPPQQGKDRDKKEAEQGS GEEKEEKEGDLKANDPYANLSKKEK KKLKKQMDYERQVESLKAANAAEN DFSVSQAEVSSRQAMLENASDIKLE KFSISAHGKELFVNADLYIVAGRRYG LVGPNGKGKTTLLKHIANRALSIPPN IDVLLCEQEVVADETPAVQAVLRADT KRLRLLEEERRLQGQLEQGDDTAAE KLEKVYEELRATGAAAAEAKARRILA GLGFDPEMQNRPTQKFSGGWRMR VSLARALFMEPTLLMLDEPTNHLDL NAVIWLNNYLQGWRKTLLIVSHDQG FLDDVCTDIIHLDTQRLHYYRGNYM TFKKMYQQKQKELLKQYEKQEKKLK ELKAGGKSTKQAEKQTKEVLTRKQQ KCRRKNQDEESQEPPELLKRPKEYT VRFTFPDPPPLSPPVLGLHGVTFGYE GQKPLFKNLDFGIDMDSRICIVGPN GVGKSTLLLLLTGKLTPTNGEMRKN HRLKIGFFNQQYAEQLHMEETPTEY LQRSFNLPYQDARKCLGRFGLESHA HTIQICKLSGGQKARVVFAELACREP DVLILDEPTNNLDIESIDALGEAINDY KGAVIVVSHDARLITETNCQLWVVE EQGVSQIDGDFDDYKREVLEALGEV MVNRPRD	True	False	3.546	4.586	1.849	1.276	1.105	1.387