

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	extracel region
P70168	IMB1_MOUSE	Mus musculus	Importin subunit beta-1	28.265827	NaN	S12	33300544;34887587;37453647;36288343;39627609	MELITILEKTVSPDRLELEAAQKFLE RAAVENLPTFLVELSRVLANPGNSQ VARVAAGLQIKNSLTSKDPDIKAQYQ QRWLAIIDANARREVKNYVLTGTE TYRPSSASQCVAGIACAEIPVSWPE LIPQLVANVTNPNSTEHMKESTLEAI GYICQDIDPEQLQDKSNEILTAHQG MRKEEPSNNVKLAATNALLNSLEFT KANFDKESERHFIMQVCEATQCPD TRVRVAALQNLVKIMSLYYQYMETY MGPALFAITIEAMKSDIDEVALQIE FWSNVCDDEMDLAIIEASEAAEQGRP PEHTSKFYAKGALQYLVPILTQTLTK QDENDDDDWNPCKAAGVCLMLLS TCCEDDIVPHVLPFIKEHIKNDWRY RDAAVMAFGSILEGPEPNQLKPLVIQ AMPTLIELMKDPSVVRRDTTAWTVG RICELLPEAAINDVYLAPLLQCLIEGL SAEPRVASNVCFWAFSSLAEEAAYEAA DVADDQEEPATYCLSSSFELIVQKLL ETDRPDGHQNNLRSSAYESLMEIV KNSAKDCYPVQKTTLVIMERLQOV QMESHQSTSDRIQFNDLQSLCA TLQNVLRKVQHQDALQISDVVMASL LRMFQSTAGSGGVQEDALMAVSTL VEVLGGEFLKYMEAFKPFLLGIGLKN YAEYQVCLAAVGLVGDLCRALQSNIL PFCDEVMLLENLGNENVHRSVK PQILSVFGDIALAIGGEFKKYLEVVLN TLQQASQAQVDKSDFDMDVYLNEL RESCLEAYTGIVQGLKGDQENVHPD VMLVQPRVEFILSFIDHIAGDEDHTD GVVACAAGLIGDLCTAFGKDVCLKVE ARPMIHELLTEGRRSKTNKAKTLAT WATKELRKLKNQA	True	False	4.72	4.174	1.817	2.882	1.317	1.506	1.579