

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	extracellular region
035604	NPC1_MOUSE	Mus musculus	NPC intracellular cholesterol transporter 1	29.185005	NaN	NaN	22645316	MGAHHPALGLLLLLCPAQVFSQSC VWYGECEGIATGDKRYNCKYSGPPKP LPKDDGYDLVQELCPGLFFDNVSLCC DIQQLQTLKSNLQLPLQFLSRCPSCF YNLMTLFCELTCSPHQSOFLNVTAT EDYFDPKTOENKTNVKELEYFVGQS FANAMYNACRDVEAPSSNEKALGLL CGRDARACNATNWIEYMFNKDNGQ APFTIHPVFDLSILGMEPMRNATKG CNESVDEVTGPCSCQDCSIVCGPKP QPPPPMPWRIWGLDAMYVIMWVT YVAFLFVFFGALLAVWCHRRRYFVS EYTPIDSNIASFVNSSDKGEASCCDP LGAAFDCLRRMFTKWGAFVCRNP TCIHFFSLAFITVCSGLVVFVQVTINP VELWSAPHSQARLEKEYFDKHFQPF FRTEQLIIQAPNTSVHIYEPYPAGADV PFGPPLNKEILHQVLDLQIAIESITAS YNNETVTLQDICVAPLSPYNKNCI MSVLNYFQNSHAVLDSQVGDDFYIY ADYHHTFLYCVRAPASLNDTSLHGG PCLGTGGFPVFPWLVLGGYDDQNY NNATALVITFPVNNYNDTERLQRA WAWKEFISFVKNYKPNLTISFTA ERSIEDELNRESNSDVFTVHSYVVM FLYISLALGHIQSCSRLLVDSKISLGI AGILVLSVACSLGIFSYMGMPLTLI VIEVIPFLVLAVGVDNIFILVQTYQRD ERLQEETLDQQLGRILGEVAPTMFL SSFSETSAFFFGALSSMPAVHTFSLF AGMAVLIDFLQITCFVSLGLDIKR QEKNHLDLCCVRGADDGQGHAS ESYLFRFFKNYFAPLLLKDWRPIVV AVFVGVLSFSVAVVNKVDIGLDQSL MPNDSYVIDYFKSLAQYLHSGPPVYF VLEEGYNYSSRKQNMVCGMGCD NDSLVOQIFNAEELDTYTRVGFAPSS WIDDYFDWVSPQSSCRLYNVTHQF CNASVMDPTCVRCRPLTEGKORPQ GKEFMKFLPMFLSDNPNPKCGKGG HAAYGSVNVGD DTYIGATYFMTY HTILKTSADYTDAMKKARLIASNITE TMRSGSDYRVFPYSVFYVFEQYLT IIDDTIFNLSVSLGSIPLVTLVVLGCE LWSAVIMCITIAMILVNMFGVMWL WGISLNAVSLVNLVMSCGISVEFCS HITRAFTMSTKGRSVRAEEALAHM GSSVFSGITLTKFGGIVVLAFAKSQIF EIFYFRMYLAMVLLGATHGLIFLPVL LSYIGPSVNAKRRHTTYERYRGTRE RLLNF	True	True	2.383	5.0	2.346	4.513	5.0	5.0	5.0