

Product datasheet for SC209222

OriGene Technologies, Inc.

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Niemann Pick C1 (NPC1) (NM_000271) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Niemann Pick C1 (NPC1) (NM 000271) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: NPC1

Synonyms: NPC; POGZ; SLC65A1

ACCN: NM_000271

Insert Size: 790 bp

Insert Sequence: >SC209222 3'UTR clone of NM_000271

The sequence shown below is from the reference sequence of NM_000271. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AGAGACAAATACAAAAGCTAAATGGAAGAGA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).





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Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 000271.5</u>

Summary: This gene encodes a large protein that resides in the limiting membrane of endosomes and

lysosomes and mediates intracellular cholesterol trafficking via binding of cholesterol to its N-

terminal domain. It is predicted to have a cytoplasmic C-terminus, 13 transmembrane domains, and 3 large loops in the lumen of the endosome - the last loop being at the N-terminus. This protein transports low-density lipoproteins to late endosomal/lysosomal compartments where they are hydrolized and released as free cholesterol. Defects in this gene cause Niemann-Pick type C disease, a rare autosomal recessive neurodegenerative disorder characterized by over accumulation of cholesterol and glycosphingolipids in late

endosomal/lysosomal compartments.[provided by RefSeq, Aug 2009]

Locus ID: 4864

MW: 30