

Product datasheet for SC206401

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

Nephrocystin 4 (NPHP4) (NM_015102) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Nephrocystin 4 (NPHP4) (NM_015102) Human 3' UTR Clone

Symbol: Nephrocystin 4
Synonyms: POC10; SLSN4

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_015102

Insert Size: 484 bp

Insert Sequence: >SC206401 3'UTR clone of NM_015102

The sequence shown below is from the reference sequence of NM_015102. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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).

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.





Nephrocystin 4 (NPHP4) (NM_015102) Human 3' UTR Clone - SC206401

RefSeq: <u>NM 015102.5</u>

Summary: This gene encodes a protein involved in renal tubular development and function. This protein

interacts with nephrocystin, and belongs to a multifunctional complex that is localized to actin- and microtubule-based structures. Mutations in this gene are associated with nephronophthisis type 4, a renal disease, and with Senior-Loken syndrome type 4, a combination of nephronophthisis and retinitis pigmentosa. Alternative splicing results in

multiple transcript variants. [provided by RefSeq, Apr 2014]

Locus ID: 261734 **MW:** 17.8