

## **Product datasheet for SC206384**

## SNX14 (NM 153816) Human 3' UTR Clone

## **Product data:**

**Product Type:** 3' UTR Clones

**Product Name:** SNX14 (NM\_153816) Human 3' UTR Clone

Symbol: SNX14

Synonyms: RGS-PX2; SCAR20

**Mammalian Cell** 

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_153816

**Insert Size:** 501 bp

Insert Sequence: >SC206384 3'UTR clone of NM\_153816

The sequence shown below is from the reference sequence of NM\_153816. 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).



**OriGene Technologies, Inc.** 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



MW:

## SNX14 (NM\_153816) Human 3' UTR Clone - SC206384

**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.

19.8

**RefSeq:** <u>NM 153816.6</u>

**Summary:** This gene encodes a member of the sorting nexin family. Members of this family have a phox

(PX) phosphoinositide binding domain and are involved in intracellular trafficking. The

encoded protein also contains a regulator of G protein signaling (RGS) domain. Regulator of G protein signaling family members are regulatory molecules that act as GTPase activating proteins for G alpha subunits of heterotrimeric G proteins. Alternate splicing results in

transcript variants encoding distinct isoforms. [provided by RefSeq, Jul 2014]

**Locus ID:** 57231