

## **Product datasheet for SC206850**

## NCOA62 (SNW1) (NM 012245) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: NCOA62 (SNW1) (NM 012245) Human 3' UTR Clone

Symbol: NCOA62

Synonyms: Bx42; FUN20; NCOA-62; Prp45; PRPF45; SKIIP; SKIP; SKIP1

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_012245

**Insert Size:** 519 bp

The sequence shown below is from the reference sequence of NM\_012245. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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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## NCOA62 (SNW1) (NM\_012245) Human 3' UTR Clone - SC206850

**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 012245.3</u>

**Summary:** This gene, a member of the SNW gene family, encodes a coactivator that enhances

transcription from some Pol II promoters. This coactivator can bind to the ligand-binding domain of the vitamin D receptor and to retinoid receptors to enhance vitamin D-, retinoic acid-, estrogen-, and glucocorticoid-mediated gene expression. It can also function as a splicing factor by interacting with poly(A)-binding protein 2 to directly control the expression of muscle-specific genes at the transcriptional level. Finally, the protein may be involved in oncogenesis since it interacts with a region of SKI oncoproteins that is required for

transforming activity. Alternative splicing results in multiple transcript variants. [provided by

RefSeq, Jan 2016]

**Locus ID:** 22938 **MW:** 19.4