

## **Product datasheet for SC204332**

## ITGB3BP (NM 014288) Human 3' UTR Clone

## **Product data:**

**Product Type:** 3' UTR Clones

Product Name: ITGB3BP (NM\_014288) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: ITGB3BP

Synonyms: CENP-R; CENPR; HSU37139; NRIF3; TAP20

**ACCN:** NM\_014288

**Insert Size:** 365 bp

Insert Sequence: >SC204332 3'UTR clone of NM\_014288

The sequence shown below is from the reference sequence of NM\_014288. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGCATGTATAAAAGTGGCA

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.

**RefSeg:** NM 014288.5



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## ITGB3BP (NM\_014288) Human 3' UTR Clone - SC204332

Summary: This gene encodes a transcriptional coregulator that binds to and enhances the activity of

members of the nuclear receptor families, thyroid hormone receptors and retinoid X receptors. This protein also acts as a corepressor of NF-kappaB-dependent signaling. This protein induces apoptosis in breast cancer cells through a caspase 2-mediated signaling pathway. This protein is also a component of the centromere-specific histone H3 variant nucleosome associated complex (CENP-NAC) and may be involved in mitotic progression by recruiting the histone H3 variant CENP-A to the centromere. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Sep 2011]

**Locus ID:** 23421 **MW:** 14.5