

## **Product datasheet for SC209452**

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## Annexin VI (ANXA6) (NM\_001155) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: Annexin VI (ANXA6) (NM 001155) Human 3' UTR Clone

Symbol: Annexin VI

Synonyms: ANX6; CBP68; CPB-II; p68; p70

**Mammalian Cell** 

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_001155

**Insert Size:** 772 bp

Insert Sequence: >SC209452 3'UTR clone of NM\_001155

The sequence shown below is from the reference sequence of NM\_001155. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

 ${\sf TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC}$ 

ACACATAAACACA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul





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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.

**RefSeq:** <u>NM 001155.5</u>

Summary: Annexin VI belongs to a family of calcium-dependent membrane and phospholipid binding

proteins. Several members of the annexin family have been implicated in membrane-related events along exocytotic and endocytotic pathways. The annexin VI gene is approximately 60 kbp long and contains 26 exons. It encodes a protein of about 68 kDa that consists of eight 68-amino acid repeats separated by linking sequences of variable lengths. It is highly similar to human annexins I and II sequences, each of which contain four such repeats. Annexin VI has been implicated in mediating the endosome aggregation and vesicle fusion in secreting epithelia during exocytosis. Alternatively spliced transcript variants have been described.

[provided by RefSeq, Aug 2010]

Locus ID: 309 MW: 28