

Product datasheet for SC308104

OriGene Technologies, Inc.

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RNF90 (TRIM7) (NM 203293) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: RNF90 (TRIM7) (NM 203293) Human Untagged Clone

Tag: Tag Free RNF90 Symbol:

Synonyms: GNIP; RNF90 **Mammalian Cell** Neomycin

Selection:

Vector: pCMV6-Entry (PS100001)

NM 203293

E. coli Selection: Kanamycin (25 ug/mL)

Restriction Sites: Sgfl-Mlul ACCN:

Insert Size: 1536 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

> point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

> into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: NM 203293.2





RNF90 (TRIM7) (NM_203293) Human Untagged Clone - SC308104

 RefSeq Size:
 2978 bp

 RefSeq ORF:
 1536 bp

 Locus ID:
 81786

 UniProt ID:
 Q9C029

 Cytogenetics:
 5q35.3

Protein Families: Druggable Genome

MW: 56.6 kDa

Gene Summary: The protein encoded by this gene is a member of the tripartite motif (TRIM) family. The TRIM

motif includes three zinc-binding domains, a RING, a B-box type 1, a B-box type 2, and a coiled-coil region. The protein localizes to both the nucleus and the cytoplasm, and may represent a participant in the initiation of glycogen synthesis. Alternative splicing results in

multiple transcript variants. [provided by RefSeq, Jul 2013]

Transcript Variant: This variant (1, also known as GNIP1) encodes the longest isoform (1).