

Product datasheet for SC330947

OriGene Technologies, Inc.

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zinc finger protein 138 (ZNF138) (NM_001271637) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: zinc finger protein 138 (ZNF138) (NM_001271637) Human Untagged Clone

Tag:Tag FreeSymbol:ZNF138Synonyms:pHZ-32

Vector: pCMV6-Entry (PS100001)

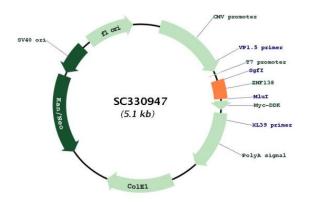
Fully Sequenced ORF: >SC330947 representing NM_001271637.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATTACCTGTCTGGAACAAGGAAAAGAGCCCTGGAATATGAAGAGACA<mark>TGA</mark>

Restriction Sites: Sgfl-Mlul

Plasmid Map:



ACCN: NM_001271637

Insert Size: 189 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).

Components:

Cytogenetics:

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: <u>NM 001271637.1</u>

 RefSeq Size:
 2624 bp

 RefSeq ORF:
 189 bp

 Locus ID:
 7697

 UniProt ID:
 P52744

Protein Families: Transcription Factors

7q11.21

MW: 7.4 kDa

Gene Summary: May be involved in transcriptional regulation as a repressor.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3) includes an alternate exon, which results in a frameshift, compared to variant 1. The encoded isoform (3) is shorter and has a distinct C-terminus, compared to isoform 1. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly.

The genomic coordinates used for the transcript record were based on alignments.