

Product datasheet for **SC337130**

ZNF135 (NM_001289401) Human Untagged Clone

Product data:

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|---------------------------|--|
| Product Type: | Expression Plasmids |
| Product Name: | ZNF135 (NM_001289401) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | ZNF135 |
| Synonyms: | pHZ-17; pT3; ZNF61; ZNF78L1 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-Entry (PS100001) |
| E. coli Selection: | Kanamycin (25 ug/mL) |

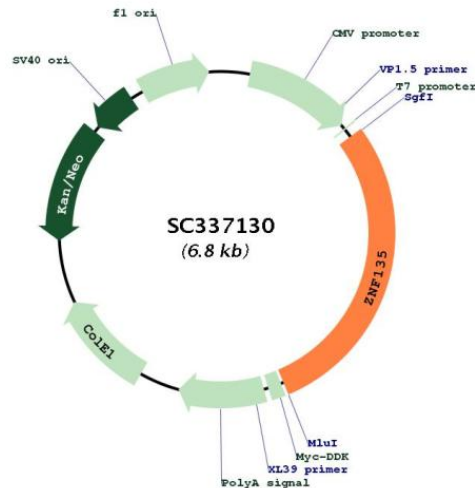


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Fully Sequenced ORF: >SC337130 representing NM_001289401.
Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGGGAATTCGTGACTG
GATCCGGTACCGAGGAGATCTGCCGCCCGCATCGCC
ATGACCCTGGGTGCGCTCTCCACAGACCCGGAGCAAGTGACGTTTGGAGACGTGGTAGTGGCTTC
AGCCAGGAGGAGTGGGGCAGCTGAAGCCTGCCAGAGGACCCTGTACCGTGATGTAATGCTGGACACC
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TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
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Restriction Sites: SgfI-MluI

Plasmid Map:


ACCN: NM_001289401

Insert Size: 1977 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: 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_001289401.1](#)

RefSeq Size: 3377 bp

RefSeq ORF: 1977 bp

Locus ID: 7694

UniProt ID: [P52742](#)

Cytogenetics: 19q13.43

Protein Families: Transcription Factors

MW: 75.3 kDa

Gene Summary:

Plays a role in the regulation of cell morphology and cytoskeletal organization. May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (6) uses an alternate in-frame splice site in the 3' coding region, and includes an alternate exon that results in a distinct 5' UTR and uses of an alternate start codon, compared to variant 1. The encoded isoform (6) is shorter and has a distinct N-terminus, compared to isoform 1.