

Product datasheet for **SC331885**

DEF8 (NM_001242822) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: DEF8 (NM_001242822) Human Untagged Clone
Tag: Tag Free
Symbol: DEF8
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC331885 representing NM_001242822.
Blue=Insert sequence Red=Cloning site Green=Tag(s)

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ATGGAATATGATGAGAAGCTGGCCGTTTCCGGCAGGCCACCTCAACCCCTTCAACAAGCAGTCTGGG
CCGAGACAGCATGAGCAGGGCCCTGGGGAGGAGGTCCCGGACGTCACTCCTGAAGAGGCCCTGCCTGAG
CTGCCCTGGGGAGCCGGAATCCGCTGCCCTGAACGCGTGATGGATCTCGGCTGTCTGAGGACCAC
TTCTCCCGCCCTGTGGGTCTGTTCTCGGCTCTGACGTCCAGCAGCTGCGGCAGGCGATCGAGGAGTGC
AAGCAGGTGATTCTGGAGCTGCCGAGCAGTCGGAGAAGCAGAAGGATGCCGTGGTGCGACTCATCCAC
CTCCGGCTGAAGCTCCAGGAGCTGAAGGACCCCAATGAGGATGAGCCAAACATCCGAGTGCCTTGAG
CACCGCTTTTACAAGGAGAAGAGCAAGAGCGTCAAGCAGACCTGTGACAAGTGTAAACACCATCATCTGG
GGGCTCATTACAGACCTGGTACACCTGCACAGGTGGGCCGAGACCCAGACGAGGAGTGAGGAATGAGAGA
GACCAAAGTTCCTGCCTCCGCTGGGCTCACATTCAGATGTGA
```

Restriction Sites: SgfI-RsrII
ACCN: NM_001242822
Insert Size: 594 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).



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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_001242822.1](#)

RefSeq Size: 929 bp

RefSeq ORF: 594 bp

Locus ID: 54849

UniProt ID: [Q6ZN54](#)

Cytogenetics: 16q24.3

MW: 23 kDa

Gene Summary: Positively regulates lysosome peripheral distribution and ruffled border formation in osteoclasts. Involved in bone resorption.[UniProtKB/Swiss-Prot Function]
Transcript Variant: This variant (9) has multiple differences in the 5' and 3' UTRs, compared to variant 1. These differences cause translation initiation at an in-frame downstream start codon, compared to variant 1. The encoded protein (isoform 2) has shorter N- and C-termini, compared to isoform 1. Variants 2, 8, and 9 encode the same protein.