

Product datasheet for SC327034

MIER1 (NM 001146112) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: MIER1 (NM_001146112) Human Untagged Clone

Tag: Tag Free Symbol: MIER1

Synonyms: ER1; MI-ER1

Mammalian Cell

Selection:

None

Vector: pCMV6-XL5

E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM_001146112, the custom clone sequence may differ by one or

more nucleotides

ATGCTGGTTCATGATTTTGATGATGAACGAACATTAGAAGAGGAAGAAATGATGGAAGGA GAAACAAACTTCAGCTCTGAAATAGAAGATCTTGCAAGGGAAGGCGACATGCCAATTCAT GAACTTCTCAGCCTTTATGGTTATGGTAGTACTGTTCGACTACCTGAAGAAGATGAGGAA GAGGAAGAAGAAGAAGAAGATGAAGATGATGAAGATGCTGATAATGATGACAACAGT GGCTGTAGTGGGGAAAATAAAGAGGAGAATATAAAGGATTCATCAGGTCAGGAGGATGAA ACTCAGTCTTCCAATGATGATCCATCACAATCTGTTGCTTCTCAAGATGCCCAGGAAATA ATCCGCCCACGTCGATGTAAATATTTTGATACAAATAGTGAAGTAGAAGAAGAATCTGAA GAAGATGAAGATTATATTCCATCAGAAGACTGGAAAAAGGAGATTATGGTGGGCTCCATG TTTCAAGCAGAAATTCCAGTTGGCATTTGTAGATACAAAGAAAATGAAAAAGTATATGAA AATGATGATCAGCTCCTGTGGGACCCTGAGTACTTACCAGAAGATAAAGTGATTATATTT CTTAAAGATGCATCTAGAAGAACAGGTGATGAGAAGGGTGTAGAAGCAATTCCTGAAGGA TCTCACATAAAAGACAATGAACAGGCTTTATATGAATTGGTTAAATGCAATTTTGATACA GAAGAAGCATTGAGAAGATTAAGATTTAATGTAAAAGCAGCTAGAGAGGAATTATCTGTT TGGACAGAGGAAGAGTGTAGAAAATTTTGAACAAGGGCTGAAGGCCTATGGAAAGGATTTT CATTTGATTCAGGCTAATAAAGTCCGAACAAGGTCAGTTGGTGAATGTGTAGCATTCTAT TACATGTGGAAAAATCTGAACGTTATGATTTCTTTGCTCAGCAAACACGATTTGGAAAG AAGAAATATAATCTTCATCCTGGTGTAACGGATTACATGGATCGTCTTCTAGACGAAAGT GAAAGTGCTGCATCTAGTCGAGCACCATCCCCTCCCCCAACTGCATCAAACAGTAGTAAC AGCCAGTCTGAGAAAGAAGATGGCACTGTAAGCACTGCTAATCAAAATGGAGTGTCATCT AATGGACCAGGTGAAATATTAAACAAAGAGGAAGTAAAAGTTGAAGGGTTACACATTAAT GGACCAACAGGTGGAAATAAGAAACCACTTCATGCAGATATGGATACTAATGGTTATGAA ACAGATAACCTTACCACTGACCCAAAACTTGCCCATATGACTGCAAGAAATGAAAATGAT TTTGATGAAAAAGTGAGAGCCTGCCAAAAGGCGAAGGGTAAACAGCAATGGAAAAGAA AGTCCAGGTTCTTCTGAATTTTTCCAAGAAGCAGTCTCACATGGGAAATTTGAAGAACTT GAAAACACAGATGAC

OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com

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Restriction Sites: Please inquire ACCN: NM_001146112

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: <u>NM 001146112.1</u>, <u>NP 001139584.1</u>

 RefSeq Size:
 5604 bp

 RefSeq ORF:
 1458 bp

 Locus ID:
 57708

 UniProt ID:
 Q8N108

 Cytogenetics:
 1p31.3

Gene Summary: This gene encodes a protein that was first identified in Xenopus laevis by its role in a

mesoderm induction early response (MIER). The encoded protein functions as a

transcriptional regulator. Alternatively spliced transcript variants encode multiple isoforms, some of which lack a C-terminal nuclear localization signal. [provided by RefSeq, May 2013] Transcript Variant: This variant (8) lacks a portion of the 5' coding region and initiates translation at a downstream in-frame start codon, compared to variant 1. The resulting isoform (h) has a shorter N-terminus, compared to isoform a. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the

transcript record were based on transcript alignments.