

## Product datasheet for **SC335828**

### MDMX (MDM4) (NM\_001278517) Human Untagged Clone

#### Product data:

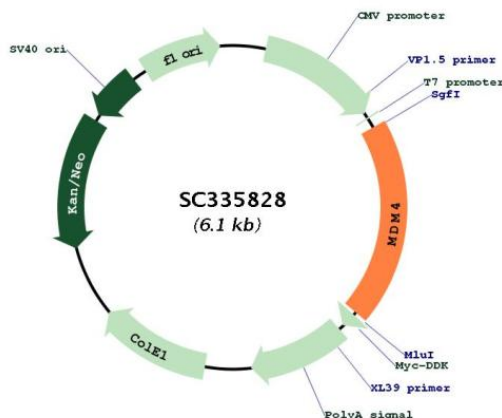
Product Type:	Expression Plasmids
Product Name:	MDMX (MDM4) (NM_001278517) Human Untagged Clone
Tag:	Tag Free
Symbol:	MDM4
Synonyms:	BMFS6; HDMX; MDMX; MRP1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC335828 representing NM_001278517. Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTGTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGACATCATTTTCCACCTCTGCTCAGTGTCAACATCTGACAGTGCTTGCAGGATCTCTCTGGACAA
ATCAATCAGGATCACAGTATGGATATCCAAGTCAAGACCAACTGAAGCAAAGTGCAGAGGAAAGTTCC
ACTTCCAGAAAAAGAACTACAGAAGACGATATCCCACACTGCCTACCTCAGAGCATAAATGCATACAT
TCTAGAGAAGATGAAGACTTAATTGAAAATTTAGCCCAAGATGAAACATCTAGGCTGGACCTTGATTT
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GACTTGTGGTTTTTGAATGAGTCAGTATCAGAGCAGTTAGGTGTTGGAATAAAAGTTGAAGCTGCTGAT
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GATTGGTATTCAGATTGTTCAAAGTTAACCATTCTCTCCACGTCTGATACTGCCATACCTGAA
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GATGCGTATATAAAGAAAGAAAACCTCAAACCTTTTGTCCCTGCAACTCAGTGGAAATCTTGATTTG
GCTCACAGTCTGAAAGCCAAGAGACCATCTCAAGCATGGGAGAACAGTTAGATAACCTTTCTGAACAG
AGAACAGATACAGAAAACATGGAGGATTGCCAGAATCTCTTGAAGCCATGTAGCTTATGTGAGAAAAGA
CCACGAGACGGGAACATTATTATGGAAGGACGGCCATCTTGTCACTTGTCTTCACTGTGCCAGAAGA
CTAAAGAAGGCTGGGCTTCATGCCCTATTTGCAAGAAAGAGATTGAGCTGGTTATTAAGTTTTTATA
GCATAA
ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
```

Restriction Sites: Sgfl-MluI



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**Plasmid Map:**


**ACCN:** NM\_001278517

**Insert Size:** 1179 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\\_001278517.1](#)

**RefSeq Size:** 9796 bp

**RefSeq ORF:** 1179 bp

**Locus ID:** 4194

**UniProt ID:** [Q15151](#)

**Cytogenetics:** 1q32.1

**Protein Families:** Druggable Genome, Transcription Factors

**Protein Pathways:** p53 signaling pathway

**MW:** 44 kDa

**Gene Summary:** This gene encodes a nuclear protein that contains a p53 binding domain at the N-terminus and a RING finger domain at the C-terminus, and shows structural similarity to p53-binding protein MDM2. Both proteins bind the p53 tumor suppressor protein and inhibit its activity, and have been shown to be overexpressed in a variety of human cancers. However, unlike MDM2 which degrades p53, this protein inhibits p53 by binding its transcriptional activation domain. This protein also interacts with MDM2 protein via the RING finger domain, and inhibits the latter's degradation. So this protein can reverse MDM2-targeted degradation of p53, while maintaining suppression of p53 transactivation and apoptotic functions. Alternatively spliced transcript variants encoding different isoforms have been noted for this gene. [provided by RefSeq, Feb 2011]

**Transcript Variant:** This variant (5, also known as MDM4-G or HDMX-G) lacks three consecutive exons and part of the fourth compared to variant 1. The resulting isoform (5) has the same N- and C-termini but lacks a large internal segment compared to isoform 1.

**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.