

## Product datasheet for RC208795L4

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OriGene Technologies, Inc.

### MTA3 (NM\_020744) Human Tagged Lenti ORF Clone

**Product data:** 

**Product Type:** Expression Plasmids

**Product Name:** MTA3 (NM\_020744) Human Tagged Lenti ORF Clone

Tag: mGFP Symbol: MTA3

Mammalian Cell Puromycin

Selection:

Vector:

pLenti-C-mGFP-P2A-Puro (PS100093)

E. coli Selection: Chloramphenicol (34 ug/mL)

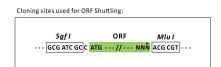
ORF Nucleotide The ORF insert of this

Sequence:

The ORF insert of this clone is exactly the same as(RC208795).

Restriction Sites: Sgfl-Mlul

**Cloning Scheme:** 





**ACCN:** NM\_020744

ORF Size: 1545 bp



### MTA3 (NM\_020744) Human Tagged Lenti ORF Clone - RC208795L4

**OTI Disclaimer:** The molecular sequence of this clone aligns with the gene accession number as a point of

reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**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 020744.2</u>

 RefSeq Size:
 2218 bp

 RefSeq ORF:
 1548 bp

 Locus ID:
 57504

 UniProt ID:
 Q9BTC8

Cytogenetics: 2p21

**Protein Families:** Transcription Factors

**MW:** 58.8 kDa

**Gene Summary:** Plays a role in maintenance of the normal epithelial architecture through the repression of

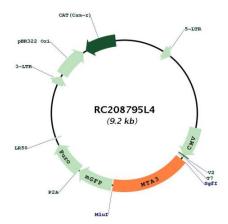
SNAI1 transcription in a histone deacetylase-dependent manner, and thus the regulation of E-

cadherin levels. Contributes to transcriptional repression by BCL6.[UniProtKB/Swiss-Prot

Function]



# **Product images:**



Circular map for RC208795L4