

Product datasheet for RN205303

Morf4l2 (NM_001007714) Rat Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Morf4l2 (NM 001007714) Rat Untagged Clone

Tag: Tag Free
Symbol: Morf4l2

Synonyms: LRRG00119
Mammalian Cell Neomycin

Selection:

E. coli Selection:

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >RN205303 representing NM_001007714

Kanamycin (25 ug/mL)

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM 001007714

Insert Size: 867 bp



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Morf4l2 (NM_001007714) Rat Untagged Clone - RN205303

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: <u>NM 001007714.1</u>, <u>NP 001007715.1</u>

RefSeq Size: 1846 bp
RefSeq ORF: 867 bp
Locus ID: 317413
UniProt ID: Q6Q189
Cytogenetics: Xq32

Gene Summary: Component of the NuA4 histone acetyltransferase complex which is involved in

transcriptional activation of select genes principally by acetylation of nucleosomal histone H4

and H2A. This modification may both alter nucleosome - DNA interactions and promote

interaction of the modified histones with other proteins which positively regulate

transcription. This complex may be required for the activation of transcriptional programs associated with oncogene and proto-oncogene mediated growth induction, tumor suppressor mediated growth arrest and replicative senescence, apoptosis, and DNA repair. The NuA4 complex ATPase and helicase activities seem to be, at least in part, contributed by the association of RUVBL1 and RUVBL2 with EP400. NuA4 may also play a direct role in DNA repair when directly recruited to sites of DNA damage. Also component of the MSIN3A complex which acts to repress transcription by deacetylation of nucleosomal histones (By

similarity).[UniProtKB/Swiss-Prot Function]