

## **Product datasheet for SC202470**

## OriGene Technologies, Inc.

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## C20orf132 (MROH8) (NM\_213632) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: C20orf132 (MROH8) (NM 213632) Human 3' UTR Clone

Symbol: C20orf132

Synonyms: C20orf131; C20orf132

Mammalian Cell

Selection:

Neomycin

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_213632

**Insert Size:** 229 bp

Insert Sequence: >SC202470 3'UTR clone of NM\_213632

The sequence shown below is from the reference sequence of NM\_213632. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GCATATGTGGCTGTGGCTGTAAGCCCTTGAGCTTTGCCCCTCTAATGTCAGCCCCTCCCAGGATGACTG ACCCCACACCCCAGTGGAGAGTGTTCCTTTTGGTCCCTCTCAAAGCCTGCATCTGTCATCCATTTATAA GTCTACAGTGTTTCTGCCAAGGGATTGACATGGAGGGAGCTGGGGAGAGTGAAGGAGTCAGAATGCCTT

AATACAAACTCGTCCAACCTGC

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**RefSeq:** <u>NM 213632.3</u>





## C20orf132 (MROH8) (NM\_213632) Human 3' UTR Clone - SC202470

Summary: The protein encoded by this gene belongs to the maestro heat-like repeat family. The exact

function of this gene is not known, however, in a genome-wide association study using hippocampal atrophy as a quantitative trait, this gene has been associated with Alzheimer's disease (PMID:19668339). Alternatively spliced transcript variants encoding different isoforms

have been found for this gene. [provided by RefSeq, Sep 2013]

**Locus ID:** 140699

**MW:** 8.1