

## **Product datasheet for SC208208**

c Maf (MAF) (NM 005360) Human 3' UTR Clone

## 9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com

OriGene Technologies, Inc.

techsupport@origene.com
EU: info-de@origene.com
CN: techsupport@origene.cn

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**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** c Maf (MAF) (NM\_005360) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: MAF

**Synonyms:** AYGRP; c-MAF; CCA4; CTRCT21

**ACCN:** NM\_005360

**Insert Size:** 652 bp

The sequence shown below is from the reference sequence of NM\_005360. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GTGTGGCAAAATACAAAGTTAGTTAAATACA

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 005360.5</u>

**Summary:** The protein encoded by this gene is a DNA-binding, leucine zipper-containing transcription

factor that acts as a homodimer or as a heterodimer. Depending on the binding site and binding partner, the encoded protein can be a transcriptional activator or repressor. This protein plays a role in the regulation of several cellular processes, including embryonic lens fiber cell development, increased T-cell susceptibility to apoptosis, and chondrocyte terminal differentiation. Defects in this gene are a cause of juvenile-onset pulverulent cataract as well as congenital cerulean cataract 4 (CCA4). Two transcript variants encoding different isoforms

have been found for this gene. [provided by RefSeq, Jan 2010]

**Locus ID:** 4094 **MW:** 24.9