

## **Product datasheet for SC206505**

## MAX (NM 145114) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: MAX (NM\_145114) Human 3' UTR Clone

Symbol: MAX

Synonyms: bHLHd4

Mammalian Cell Neomycin

Selection:

**Vector:** pMirTarget (PS100062)

**ACCN:** NM\_145114

**Insert Size:** 466 bp

The sequence shown below is from the reference sequence of NM\_145114. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TTCATTTTCTCTAAAATAACTAGTACAAAATAAAACAGATGTCAGTTGTTGA

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.



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**RefSeq:** <u>NM 145114.3</u>

Summary: The protein encoded by this gene is a member of the basic helix-loop-helix leucine zipper

(bHLHZ) family of transcription factors. It is able to form homodimers and heterodimers with other family members, which include Mad, Mxi1 and Myc. Myc is an oncoprotein implicated in cell proliferation, differentiation and apoptosis. The homodimers and heterodimers compete for a common DNA target site (the E box) and rearrangement among these dimer forms provides a complex system of transcriptional regulation. Mutations of this gene have been reported to be associated with hereditary pheochromocytoma. A pseudogene of this gene is located on the long arm of chromosome 7. Alternative splicing results in multiple transcript

variants. [provided by RefSeq, Aug 2012]

**Locus ID:** 4149 **MW:** 17.9