

## Product datasheet for SC207888

### MAGEA9 (NM\_005365) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	MAGEA9 (NM_005365) Human 3' UTR Clone
Symbol:	MAGEA9
Synonyms:	CT1.9; MAGE9
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_005365
Insert Size:	609 bp
Insert Sequence:	<p>&gt;SC207888 3'UTR clone of NM_005365</p> <p>The sequence shown below is from the reference sequence of NM_005365. The complete sequence of this clone may contain minor differences, such as SNPs.</p> <p>Blue=Stop Codon Red=Cloning site</p>

```

GGCAAGTTGGACGCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAACGATCGCC
GTTTTGGGAGAGGAGCAAGAGGGAGTCTAGCACCAGCCGCGAGCCGGGGCCAAAGTTTGTGGGGTCAGG
GCCCCATCCAGCAGCTGCCCTGCCCCATGTGACATGAGGCCCATTTCTCGCTCTGTGTTGAAGAGAGC
AATCAGTGTTCTCAGTGGCAGTGGGTGGAAGTGAACACACTGTATGTCATCTCTGGGTTCTTGTCTAT
TGGGTGATTTGGAGATTTATCCTTGCTCCCTTTTGAATTGTTCAAATGTTCTTTAATGGTCAGTTTA
ATGAACCTCACCATCGAAGTTAATGAATGACAGTAGTCACACATATTGCTGTTTATGTTATTTAGGAGT
AAGATTCTTGCTTTGAGTCACATGGGGAAATCCCTGTTATTTTGTGAATTGGGACAAGATAACATAGC
AGAGGAATTAATAATTTTTTTGAACTTGAACCTAGCAGCAAAATAGAGCTCATAAAGAAATAGTGAAA
TGAAAATGTAGTTAATCTTGCTTATACCTCTTCTCTCTCTGTAATAAATACATATACATGTAT
ACCTGGATTTGCTTGCTTCTTTGAGCATGTAAGAGAAATAAATTTGAAAGAATAA
ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
  
```

Restriction Sites: SgfI-MluI

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).


[View online »](#)

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><a href="#">NM_005365.5</a></u>
Summary:	This gene is a member of the MAGEA gene family. The members of this family encode proteins with 50 to 80% sequence identity to each other. The promoters and first exons of the MAGEA genes show considerable variability, suggesting that the existence of this gene family enables the same function to be expressed under different transcriptional controls. The MAGEA genes are clustered at chromosomal location Xq28. They have been implicated in some hereditary disorders, such as dyskeratosis congenita. [provided by RefSeq, Jul 2008]
Locus ID:	4108
MW:	23.2