

## Product datasheet for **SC205649**

### ZNF98 (NM\_001098626) Human 3' UTR Clone

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** ZNF98 (NM\_001098626) Human 3' UTR Clone  
**Vector:** pMirTarget (PS100062)  
**Symbol:** ZNF98  
**Synonyms:** F7175; ZNF739  
**ACCN:** NM\_001098626  
**Insert Size:** 527 bp  
**Insert Sequence:** >SC205649 3'UTR clone of NM\_001098626  
The sequence shown below is from the reference sequence of NM\_001098626. The complete sequence of this clone may contain minor differences, such as SNPs.  
**Blue**=Stop Codon **Red**=Cloning site

```
GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
TATAAAAGAAATTGTGCTGGTGAGAAAATAGAAATATGAATGTGACAAAGCCTTTAAATGGTTATCA
CACTGGATTGTAGGTAAGGTAATTCATTCTGGAGAAAATTCTACAAGTGTGAACAGTGTGGCAAACT
TACACAATGCTCAAACCTTATTGCACAGGAAAGCCTTTATACTTGATGAGAACAATGTACAAATATAA
AGAAAGTAAAAAGTGATTAATATCTGCACACATCTTATTTCAGCATCAGAGAATTTATACTTAATAGCA
TTAAAGGTGCAATTACTGTCAAAGAGTTTCAGAAAATATAAGCATTAAAGTGTGAAGAGGATTTATT
TTGAAGACAAACATTACAAATATAAACGAAGTTGTGGTAACTTTACTTGAACACAGATCTTGTGTAT
TCATTCTGTATTAGAGCAAACCTGAGGCAATTGCTCAAACCTTTGTTCAACATCAGAGAATTTATATCA
AAGAAAAACCTGTAATGTAATAAATTTAGAAAAAACATTTA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

**Restriction Sites:** Sgfl-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).

**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:** [NM\\_001098626.2](#)



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**Summary:** May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function]  
**Locus ID:** 148198  
**MW:** 20.5