

## Product datasheet for **SC205622**

### ZNF274 (NM\_016324) Human 3' UTR Clone

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** ZNF274 (NM\_016324) Human 3' UTR Clone  
**Symbol:** ZNF274  
**Synonyms:** HFB101; ZF2; ZKSCAN19; ZSCAN51  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pMirTarget (PS100062)  
**ACCN:** NM\_016324  
**Insert Size:** 437 bp  
**Insert Sequence:** >SC205622 3' UTR clone of NM\_016324

The sequence shown below is from the reference sequence of NM\_016324. The complete sequence of this clone may contain minor differences, such as SNPs. **Red**=Cloning site  
**Blue**=Stop Codon

CAATTGGCAGAGCTCAGAATTCAA**GCGATCGC**

GAAACAGCCTACCTC**ATAG**CTCTCAAGCCAGTTGAAGAAACCTTGCCTTTTCAGCTTGACCCTGCAATAT  
AACATGCACAGGCCTGCTTGTGAATCAGGACTGAATGTGAAAGGGAAGTATTGAGTGAGGACATTCCCAA  
AACCAAAGGACA**ACTGAGGAGACTGCC**CAGCACATAATGAATAAATAAGAAAATGAGTGAGGAGTTATTA  
ACATCATTGGAAAAAGATTTCCCA**TTCACTT**GATATTGTTTGTCTACTCATTAGTCATTAAGTGA  
GATTAATAAAATCTGAAAATGTTATATAA**ACTTTAAAA**AGCCAGGTAATTAATCTGCACTGATAT  
TACATCCACAGTACCACAGTATTATGTGTATGAATTAAGGATTA**AAA**AGATAATGTGGATAAAATAAACTA  
TTGATCTATGTCTGTGT

**ACGCGT**AAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCG

**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).



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<b>Components:</b>	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.
<b>RefSeq:</b>	<a href="#">NM_016324.2</a>
<b>Summary:</b>	This gene encodes a zinc finger protein containing five C2H2-type zinc finger domains, one or two Kruppel-associated box A (KRAB A) domains, and a leucine-rich domain. The encoded protein has been suggested to be a transcriptional repressor. It localizes predominantly to the nucleolus. Alternatively spliced transcript variants encoding different isoforms exist. These variants utilize alternative polyadenylation signals. [provided by RefSeq, Jul 2008]
<b>Locus ID:</b>	10782