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# Product datasheet for SC205624

## ZNF274 (NM_016325) Human 3' UTR Clone

### Product data:

| | |
|---|---|
| **Product Type:** | 3' UTR Clones |
| **Product Name:** | ZNF274 (NM_016325) Human 3' UTR Clone |
| **Vector:** | pMirTarget (PS100062) |
| **Symbol:** | ZNF274 |
| **Synonyms:** | HFB101; ZF2; ZKSCAN19; ZSCAN51 |
| **ACCN:** | NM_016325 |
| **Insert Size:** | 437 bp |
| **Insert Sequence:** | >SC205624 3' UTR clone of NM_016325<br>The sequence shown below is from the reference sequence of NM_016325. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site Blue=Stop Codon<br><br>CAATTGGCAGAGCTCAGAATTCAAGCGATCGC<br><br>GAAACAGCCTACCTCA**TAG**CTCTCAAGCCAGTTGAAGAAACCTTGCCTTTTCAGCTTGACCCTGCAATAT<br>AACATGCACAGGCCTGCTTGTGAATCAGGACTGAATGTGAAAGGGAAGTATTGAGTGAGGACATTCCCAA<br>AACCAAAGGACAACTGAGGAGACTGCCCAGCACATAATGAATAAATAAGAAAATGAGTGAGGAGTTATTA<br>ACATCATTTGGAAAAAAGATTTCCCATTCACTTGATATTGTTTGTTCACTCATTTAGTCATTAAAAGTGA<br>GATTAATAAAATCTGAAAATGTTATATAATAACTTTAAAAAGCCAGGTAATTAATAATCTGCACTGATAT<br>TACATCCACAGTACCACAGTATTTATGTGTATGAATTAAGGATTAAAAGATAATGTGGATAAATAAACTA<br>TTGATCTATGTCTGTGT<br><br>ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCG |
| **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). |
| **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_016325.2 |

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**Summary:** 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]

**Locus ID:** 10782