

## Product datasheet for SC201468

### SCAND1 (NM\_016558) Human 3' UTR Clone

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** SCAND1 (NM\_016558) Human 3' UTR Clone  
**Symbol:** SCAND1  
**Synonyms:** RAZ1; SDP1  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pMirTarget (PS100062)  
**ACCN:** NM\_016558  
**Insert Size:** 151 bp  
**Insert Sequence:** >SC201468 3'UTR clone of NM\_016558  
 The sequence shown below is from the reference sequence of NM\_016558. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

```

GGCAAGTTGGACGCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CGCCGCACGGATGTGCGCATCACTGGCTGACGGTGGAGCTGCGGGCGGCCAGGGCCGGGCGCTCTGTG
CGGACTGGGGCCATGATCGGGCCCGGGGCGCTGAGCCTGGGACCCACCCCGTGTAAATGAAAAATGAG
TTTTGGCAGCGCC
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
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).  
**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\_016558.4


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Summary:	This gene encodes a SCAN box domain-containing protein. The SCAN domain is a highly conserved, leucine-rich motif of approximately 60 aa originally found within a subfamily of zinc finger proteins. This gene belongs to a family of genes that encode an isolated SCAN domain, but no zinc finger motif. This protein binds to and may regulate the function of the transcription factor myeloid zinc finger 1B. Alternate splicing results in multiple transcript variants.[provided by RefSeq, Jan 2011]
Locus ID:	51282
MW:	5.5