

## Product datasheet for SC201527

### C11orf2 (VPS51) (NM\_013265) Human 3' UTR Clone

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** C11orf2 (VPS51) (NM\_013265) Human 3' UTR Clone  
**Symbol:** C11orf2  
**Synonyms:** ANG2; ANG3; C11orf2; C11orf3; FFR; PCH13  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pMirTarget (PS100062)  
**ACCN:** NM\_013265  
**Insert Size:** 305 bp  
**Insert Sequence:** >SC201527 3'UTR clone of NM\_013265  
 The sequence shown below is from the reference sequence of NM\_013265. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

```
GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GTGGTTGAGGTCTCTGCGAGCGCGGCTAGGCGCAGCCGCTGCCATGCACCGGTCTGTCCCTGCACCCC
ATGGCACCCAGGATCTGGTCTCGGTGGTCTTCCCCGAGGCAGGTGTCAGGACCGGCTAATAAACAT
GTGTGGCCTCCTCTCTCGCTTGTGGCGGGCCTTCCGGGGGGGGTTTTGAAGCTGAGGCTTCTG
AGGCGCCCGCTCGGGTCCGCCCGAGCGCCGATTGGCTGGTGTCTGGGCCAGCATGGGCAGGGGG
CGGTTCCACTTAAAAACCTGGGACGAGA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
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\\_013265.4](#)



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**Summary:** This gene encodes a member of the vacuolar protein sorting-associated protein 51 family. The encoded protein is a component of the Golgi-associated retrograde protein complex which acts as a tethering factor for carriers in retrograde transport from the early and late endosomes to the trans-Golgi network. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Dec 2012]

**Locus ID:** 738

**MW:** 11.3