

Product datasheet for **SC201128**

ADAM32 (NM_145004) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: ADAM32 (NM_145004) Human 3' UTR Clone
Symbol: ADAM32
Mammalian Cell Selection: Neomycin
Vector: pMirTarget (PS100062)
ACCN: NM_145004
Insert Size: 148 bp
Insert Sequence: >SC201128 3'UTR clone of NM_145004
 The sequence shown below is from the reference sequence of NM_145004. The complete sequence of this clone may contain minor differences, such as SNPs.
 Blue=Stop Codon Red=Cloning site

```

GGCAAGTTGGACGCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAACGATCGCC
AGTACCCAAACACAAAGCAGTAGTAACAGTGATTCCTTCAGAAGGCAACGGATAACATCGAGAGTCTC
GCTAAGAAATGAAAATTCTGTCTTTCTTCCGTGGTCACAGCTGAAAGAAACAATAAATTGAGTGTGGA
TCAATTTGCA
ACGCGTAAGCGGCCGCGGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
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_145004.7](#)


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Summary:

This gene encodes a member of the disintegrin family of membrane-anchored proteins that play a role in diverse biological processes such as brain development, fertilization, tumor development and inflammation. This gene is predominantly expressed in the testis. The encoded protein undergoes proteolytic processing to generate a mature polypeptide comprised of an metalloprotease, disintegrin and epidermal growth factor-like domains. This gene is located in a cluster of other disintegrin and metallopeptidase family genes on chromosome 8. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Sep 2015]

Locus ID:

203102

MW:

5.5