

## **Product datasheet for SC202127**

## NME2 (NM 001018137) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: NME2 (NM\_001018137) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: NME2

Synonyms: NDKB; NDPK-B; NDPKB; NM23-H2; NM23B; PUF

**ACCN:** NM\_001018137

**Insert Size:** 170 bp

Insert Sequence: >SC202127 3'UTR clone of NM\_001018137

The sequence shown below is from the reference sequence of NM\_001018137. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCTTGTGCTCATGACTGGGTCTATGAATAAGAGGTGGACACAACAGCAGTCTCCTTCAGCACGGCGTGGTGTGTCCCTTGGACACAGCTCTTCATTCCATTGACTTAGAGGCAACAGGATTGATCATTCTTTTATAGAG

CATATTTGCCAATAAAGCTTTTGGAAGCCGGA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

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.

**RefSeg:** NM 001018137.3



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## NME2 (NM\_001018137) Human 3' UTR Clone - SC202127

Summary: Nucleoside diphosphate kinase (NDK) exists as a hexamer composed of 'A' (encoded by

NME1) and 'B' (encoded by this gene) isoforms. Multiple alternatively spliced transcript variants have been found for this gene. Read-through transcription from the neighboring upstream gene (NME1) generates naturally-occurring transcripts (NME1-NME2) that encode a fusion protein comprised of sequence sharing identity with each individual gene product.

[provided by RefSeq, Nov 2010]

**Locus ID:** 4831 **MW:** 6.5