

## Product datasheet for SC202526

## DDAH2 (NM 013974) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** DDAH2 (NM\_013974) Human 3' UTR Clone

Symbol:

DDAH; DDAHII; G6a; HEL-S-277; NG30 Synonyms:

**Mammalian Cell** 

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM 013974

**Insert Size:** 227 bp

**Insert Sequence:** >SC202526 3'UTR clone of NM\_013974

The sequence shown below is from the reference sequence of NM\_013974. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TTGGTGCTCAGCACACGCCCCACAGCTGAGGGCCTGGCCTTGGGGTACTGCTGGCCAGGGGTAGGATA GTATAGGAAGTAGAAGGGGAAGGAGGGTTAGATAGAGAATGCTGAATAGGCAGTAGTTGGGAGAGAGCC 

AAAATAGAATTGACCTTTTA

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.

NM 013974.3 RefSeq:



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



## DDAH2 (NM\_013974) Human 3' UTR Clone - SC202526

**Summary:** This gene encodes a dimethylarginine dimethylaminohydrolase. The encoded enzyme

functions in nitric oxide generation by regulating the cellular concentrations of

methylarginines, which in turn inhibit nitric oxide synthase activity. The protein may be localized to the mitochondria. Alternative splicing resulting in multiple transcript variants.

[provided by RefSeq, Dec 2014]

**Locus ID:** 23564

MW: 8