

Product datasheet for SC201020

NDUFB2 (NM 004546) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: NDUFB2 (NM_004546) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: NDUFB2

Synonyms: AGGG; CI-AGGG

ACCN: NM_004546

Insert Size: 140 bp

Insert Sequence: >SC201020 3'UTR clone of NM_004546

The sequence shown below is from the reference sequence of NM_004546. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TTAGGTATCCCTCCTGATGATGAAGACTGAAGGTGTAGACTCAGCCTCACTCTGTACAAGAGCCAGGTGAGAATTTCAAGGATTATCGACTTCATATTGCACATTAAAGTTACAAATTAAAGTGGCTTGGTCAAGAAT

GΑ

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 004546.3



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



NDUFB2 (NM_004546) Human 3' UTR Clone - SC201020

Summary: The protein encoded by this gene is a subunit of the multisubunit NADH:ubiquinone

oxidoreductase (complex I). Mammalian complex I is composed of 45 different subunits. This protein has NADH dehydrogenase activity and oxidoreductase activity. It plays a important role in transfering electrons from NADH to the respiratory chain. The immediate electron acceptor for the enzyme is believed to be ubiquinone. Hydropathy analysis revealed that this subunit and 4 other subunits have an overall hydrophilic pattern, even though they are found within the hydrophobic protein (HP) fraction of complex I. [provided by RefSeq, Jul 2008]

Locus ID: 4708

MW: 5.3