

Product datasheet for SC201243

ATP5MD (NM 032747) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: ATP5MD (NM 032747) Human 3' UTR Clone

Symbol: ATP5MD

Synonyms: bA792D24.4; DAPIT; HCVFTP2; MC5DN6; USMG5

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_032747

Insert Size: 150 bp

Insert Sequence: >SC201243 3'UTR clone of NM_032747

The sequence shown below is from the reference sequence of NM_032747. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAAAAACTCCAGCTGTGAAAGCAACATAAATGGATTTTAAACTGTCTACGGTTCTTAACCTCATCTGT TAAGTTCCCATGCCTGGAGAAGCTAATGCCAACTCATCATGTGATAATTCAATTTGTACAATAAATTAT

GAACCTGGAAAA

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.

RefSeq: <u>NM 032747.4</u>



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



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Summary:

Mitochondrial membrane ATP synthase (F(1)F(0) ATP synthase or Complex V) produces ATP from ADP in the presence of a proton gradient across the membrane which is generated by electron transport complexes of the respiratory chain. F-type ATPases consist of two structural domains, F(1) - containing the extramembraneous catalytic core and F(0) - containing the membrane proton channel, linked together by a central stalk and a peripheral stalk. During catalysis, ATP synthesis in the catalytic domain of F(1) is coupled via a rotary mechanism of the central stalk subunits to proton translocation (Probable). Minor subunit required to maintain the ATP synthase population in the mitochondria (PubMed:21345788). [UniProtKB/Swiss-Prot Function]

Locus ID: 84833

MW: 5.4