

Product datasheet for SC209090

NARS2 (NM_024678) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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3' UTR Clones
NARS2 (NM_024678) Human 3' UTR Clone
pMirTarget (PS100062)
NARS2
asnRS; DFNB94; SLM5
NM_024678
732 bp
<pre>>SC209090 3'UTR clone of NM_024678 The sequence shown below is from the reference sequence of NM_024678. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCACGCGAGATTCTCATTAAGGCCAAGAAGGCGCGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CCAAGGTTTCCTCATTCATGCCTTTTATAGCTGGAAGATTGGTTAAGGAAAAGCACCCCCCCATGGCAGA GACACTGCACATGATTGTGCATACAGCAGAATGCAGGTTGGTATAGGAAAAGCACCCCCCATGGCAGA GACACTGCACATGATTGTGCATACAGCAGAATGCAGATTGGATGTTGGATTTTCAGAAATGCAGGGTA TTTCACGTAAGGATAGCAGCAGAATGCAGGTACTTATAGCAAGGGGACTCTCAAATCGATGAGGGTA TTTCACGTAAGGATAGCAGAAGAGAGGTACTTATAGCAAGGGGACTCTCAAATCATAGCTAGAAATGAAATGAAGAGAAGAGAGAG</pre>
Sgfl-Mlul
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).



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	NARS2 (NM_024678) Human 3' UTR Clone – SC209090
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 024678.6</u>
Summary:	This gene encodes a putative member of the class II family of aminoacyl-tRNA synthetases. These enzymes play a critical role in protein biosynthesis by charging tRNAs with their cognate amino acids. This protein is encoded by the nuclear genome but is likely to be imported to the mitochondrion where it is thought to catalyze the ligation of asparagine to tRNA molecules. Mutations in this gene have been associated with combined oxidative phosphorylation deficiency 24 (COXPD24). [provided by RefSeq, Mar 2015]
Locus ID:	79731
MW:	28.9

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