

Product datasheet for SC329639

OriGene Technologies, Inc.

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Nucleotide binding protein like (NUBPL) (NM 001201574) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Nucleotide binding protein like (NUBPL) (NM 001201574) Human Untagged Clone

Tag: Tag Free
Symbol: NUBPL

Synonyms: C14orf127; hulnd1; IND1; MC1DN21

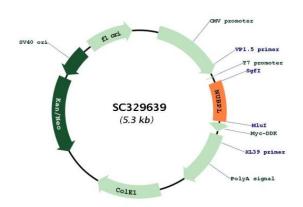
Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329639 representing NM_001201574.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

Restriction Sites: Sgfl-Mlul

Plasmid Map:



ACCN: NM 001201574



ORÏGENE

Insert Size: 411 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

RefSeq: <u>NM 001201574.1</u>

 RefSeq Size:
 2739 bp

 RefSeq ORF:
 411 bp

 Locus ID:
 80224

 Cytogenetics:
 14q12

 MW:
 14.7 kDa

Gene Summary: This gene encodes a member of the Mrp/NBP35 ATP-binding proteins family. The encoded

protein is required for the assembly of the respiratory chain NADH dehydrogenase (complex I), an oligomeric enzymatic complex located in the inner mitochondrial membrane. Mutations in this gene cause mitochondrial complex I deficiency. Alternative splicing results in multiple

transcript variants. [provided by RefSeq, May 2014]

Transcript Variant: This variant (3) lacks several exons from the 5' end and has an alternate 5' exon, as compared to variant 1. The resulting isoform (3) has a much shorter N-terminus, as compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome

assembly. The genomic coordinates used for the transcript record were based on transcript

alignments.