

Product datasheet for SC311349

NSL1 (NM 001042549) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: NSL1 (NM_001042549) Human Untagged Clone

Tag: Tag Free Symbol: NSL1

Synonyms: C1orf48; DC8; MIS14

Mammalian Cell Neomycin

Selection:

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC311349 representing NM_001042549.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGGCGGGGTCTCCTGAGTTGGTGGTCCTTGACCCTCCATGGGACAAGGAGCTCGCGGCTGGCACAGAG AGCCAGGCCTTGGTCTCCCGCACTCCCCGAGAAAGACTTTCGGGTGCGCTGCACCTCGAAGCGGGCTGTG ACCGAAATGCTACAACTGTGCGGCCGCTTCGTGCAAAAGCTCGGGGACGCTCTGCCGGAGGAGATTCGG GAGCCCGCTCTGCGAGATGCGCAGTGGACTTTTGAATCAGCTGTGCAAGAGAATATCAGCATTAATGGG CAAGCATGGCAGGAAGCTTCAGATAATTGTTTTATGGATTCTGACATCAAAGTACTTGAAGATCAGTTT GATGAAATCATAGTAGATATAGCCACAAAACGTAAGCAGTATCCCAGAAAGATCCTGGAATGTGTCATC AAAACCATAAAAGCAAAACAAGAAATTCTGAAGCAGTACCACCCTGTTGTACATCCACTGGACCTAAAA TATGACCCTGATCCAGTTCTCAACGGGAATGCTTTCAACTTTTCCCCATTCAACATGATGTTGGCTGTG GATTTTGTCATATATGGTTTTTATTACTTCGAGCCCCTCATATGGAAAATTTGAAATGCAGAGGGGAAAC

AGTAGCAAAGGAGATCAG<mark>TGA</mark>

ACGCGTACGCGCCCCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul

ACCN: NM 001042549

Insert Size: 642 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).



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



OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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: NM 001042549.1

 RefSeq Size:
 13233 bp

 RefSeq ORF:
 642 bp

 Locus ID:
 25936

 UniProt ID:
 Q96IY1

 Cytogenetics:
 1q32.3

MW: 24.1 kDa

Gene Summary: This gene encodes a protein with two coiled-coil domains that localizes to kinetochores,

which are chromosome-associated structures that attach to microtubules and mediate chromosome movements during cell division. The encoded protein is part of a conserved protein complex that includes two chromodomain-containing proteins and a component of the outer plate of the kinetochore. This protein complex is proposed to bridge centromeric heterochromatin with the outer kinetochore structure. Multiple transcript variants encoding different isoforms have been found for this gene. There is a pseudogene of the 3' UTR region

of this gene on chromosome X. [provided by RefSeq, Jul 2014]

Transcript Variant: This variant (2) contains an alternate exon, which results in a frameshift, compared to variant 1. The encoded isoform (2) has a distinct C-terminus and is shorter than 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.