

## Product datasheet for **RC235131**

### NLRP3 (NM\_001243133) Human Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	NLRP3 (NM_001243133) Human Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	NLRP3
Synonyms:	AGTAVPRL; All; AVP; C1orf7; CIAS1; CLR1.1; DFNA34; FCAS; FCAS1; FCU; KEFH; MWS; NALP3; PYPAF1
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
ORF Nucleotide Sequence:	>RC235131 representing NM_001243133 Red=Cloning site Blue=ORF Green=Tags(s)

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GCC**CGATCGCC**

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GGACTGCAGAAGCGGGATGTGTCTGCTTTCTGAGGATGAACCTGTTCCAAAAGGAAGTGGACTGCGAGA
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**Protein Sequence:**

>RC235131 representing NM\_001243133

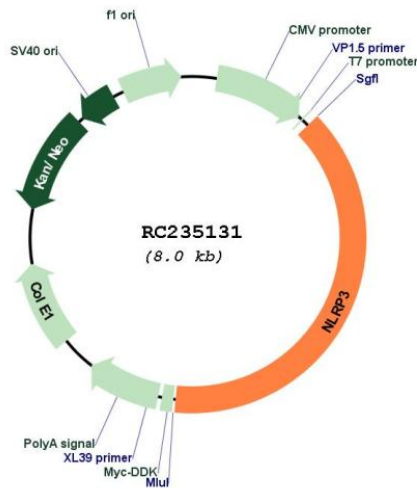
Red=Cloning site Green=Tags(s)

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YVRSRFQCIEDRNARLGESVSLNKRYTRLRIKEHRSQQEREQELLAIGKTKTCESPVSPKMELLFDPD
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PNPPIHKIVRKP SRILFLMDGFDELQGA FDEHIGPLCTDWQKAERGDILLSSLIRKKLLPEASLLITTRP
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CLYEMQEEDFVQRAMDYFPKIEINLSTRMDHMVSSF CIENCHRVESLSLGF LHNMPKEEEEEKEGRHLD
MVQCVL PSSSHAACSHGLVNSHL TSSFCRGLF SVLSTSQSL TELDLSDNSLGDGPMRVL CETLQHPGCNI
RRLWLGRGCL SHECCFDISLVLSSNQKLV ELDLSDNALGDFGIRLL CVGLKHL LCNLKKLWL VSCCLTSA
CCQDLASV LSTSHSL TRLYVGENALGDSGVA ILC EAKNPQC NLQKLG L VNSGL TSVCCSAL SSVLSTNQ
NLTHLYLRGNTLGDKG I KLLCEGLLHPDCKLQVLELDNCNL TSHCCWDLSTLL TSSQSLRKL SLGNNDLG
DLGVMMFCEVLKQQSCLLQNLGLSEMYFN YETKSALETLQEEKPELTVVFEP SW
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TRTRPLEQKLI SEEDLAANDILDYKDDDDKV
```

**Restriction Sites:**

SgfI-MluI

**Cloning Scheme:****Plasmid Map:****ACCN:**

NM\_001243133

**ORF Size:**

3102 bp

**OTI Disclaimer:**

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

<b>OTI Annotation:</b>	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<u><a href="#">NM_001243133.1</a></u> , <u><a href="#">NP_001230062.1</a></u>
<b>RefSeq Size:</b>	4470 bp
<b>RefSeq ORF:</b>	3105 bp
<b>Locus ID:</b>	114548
<b>UniProt ID:</b>	<u><a href="#">Q96P20</a></u>
<b>Cytogenetics:</b>	1q44
<b>Protein Families:</b>	Druggable Genome
<b>Protein Pathways:</b>	NOD-like receptor signaling pathway
<b>MW:</b>	118.4 kDa
<b>Gene Summary:</b>	<p>This gene encodes a pyrin-like protein containing a pyrin domain, a nucleotide-binding site (NBS) domain, and a leucine-rich repeat (LRR) motif. This protein interacts with the apoptosis-associated speck-like protein PYCARD/ASC, which contains a caspase recruitment domain, and is a member of the NLRP3 inflammasome complex. This complex functions as an upstream activator of NF-kappaB signaling, and it plays a role in the regulation of inflammation, the immune response, and apoptosis. The SARS-CoV 3a protein, a transmembrane pore-forming viroporin, has been shown to activate the NLRP3 inflammasome via the formation of ion channels in macrophages. Mutations in this gene are associated with familial cold autoinflammatory syndrome (FCAS), Muckle-Wells syndrome (MWS), chronic infantile neurological cutaneous and articular (CINCA) syndrome, neonatal-onset multisystem inflammatory disease (NOMID), keratoendotheliitis fugax hereditarian, and deafness, autosomal dominant 34, with or without inflammation. Multiple alternatively spliced transcript variants encoding distinct isoforms have been identified for this gene. Alternative 5' UTR structures are suggested by available data; however, insufficient evidence is available to determine if all of the represented 5' UTR splice patterns are biologically valid. [provided by RefSeq, Aug 2020]</p>