

Product datasheet for SC329936

NCR1 (NM 001242357) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: NCR1 (NM_001242357) Human Untagged Clone

Tag: Tag Free Symbol: NCR1

Synonyms: CD335; LY94; NK-p46; NKP46

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329936 representing NM_001242357.

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

AGAAGGCTGAACACACAGACTCTT<mark>TGA</mark>

Restriction Sites: Sgfl-Mlul

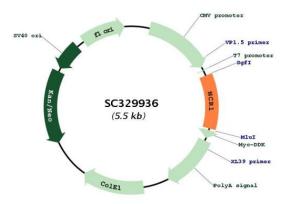
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Plasmid Map:



ACCN: NM_001242357

Insert Size: 579 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 001242357.2</u>

 RefSeq Size:
 817 bp

 RefSeq ORF:
 579 bp

 Locus ID:
 9437

 UniProt ID:
 076036

 Cytogenetics:
 19q13.42

Protein Families: Druggable Genome, Transmembrane
Protein Pathways: Natural killer cell mediated cytotoxicity

MW: 21.6 kDa

Gene Summary: Cytotoxicity-activating receptor that may contribute to the increased efficiency of activated

natural killer (NK) cells to mediate tumor cell lysis.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (5) lacks two alternate exons in the coding region compared to variant 1. The resulting protein (isoform e) is shorter but has the same N- and C-termini

compared to isoform a.