

Product datasheet for SC331322

NCR2 (NM 001199510) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: NCR2 (NM_001199510) Human Untagged Clone

Tag: Tag Free Symbol: NCR2

Synonyms: CD336; dJ149M18.1; LY95; NK-p44; NKP44

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC331322 representing NM_001199510.

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

TATGGTGGAAAACCATGA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001199510

Insert Size: 777 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).



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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 001199510.1</u>

RefSeq Size: 1048 bp
RefSeq ORF: 777 bp
Locus ID: 9436
UniProt ID: 095944
Cytogenetics: 6p21.1

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

MW: 28.4 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 (3) has an additional exon which results in frame-shift, as compared to variant 1. The resulting isoform (3) has a shorter and distinct C-terminus, as compared to isoform 1. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly.

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