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## Recombinant human NKp46/NCR1 protein

Catalog Number: NCR3001

#### PRODUCT INFORMATION

#### **Expression system**

E.coli

#### **Domain**

22-255aa

#### UniProt No.

076036

#### **NCBI Accession No.**

NP 004820.1

#### **Alternative Names**

NKp46 Extracellular Ig-like domain, NK-p46, NK cell-activating receptor, NCR1, NCR, Natural killer cell p46related protein, Natural cytotoxicity triggering receptor 1 isoform a, Natural cytotoxicity triggering receptor 1, Lymphocyte antigen 94 homolog, Ly96, LY94, hNKp46, CD335 antigen

## PRODUCT SPECIFICATION

#### **Molecular Weight**

26.6 kDa (235aa) confirmed by MALDI-TOF

#### Concentration

1mg/ml (determined by Bradford assay)

#### **Formulation**

Liquid in. Phosphate-Buffered Saline (pH 7.4) containing 10% glycerol, 1mM EDTA

#### **Purity**

> 95% by SDS-PAGE

#### **Endotoxin level**

< 1 EU per 1ug of protein (determined by LAL method)

#### Tag

Non-Tagged

## **Application**

SDS-PAGE

#### **Storage Condition**

Can be stored at +2C to +8C for 1 week. For long term storage, aliquot and store at -20C to -80C. Avoid repeated freezing and thawing cycles.

## **BACKGROUND**

## **Description**

A natural cytotoxicity receptor (NCR) NKp46 has been shown to represent a novel NK cell-specific molecule involved in human NK cell activation. The natural cytotoxicity receptors (NCRs) are a recently characterized



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family of Ig-like activation receptors that appear to be major triggering receptors in tumor cell recognition. The three known NCRs include NKp46 and NKp30, which are expressed on circulating NKcells, and NKp44, which is expressed only on activating NK cells. NKp46 has been implicated in NK cell-mediated lysis of several autologous tumor cells and pathogen-infected cell lines. NKp46 has two extracellular Ig-like domains followed by a ~40 residue stalk region, a type I transmembrane domain, and a short cytoplasmic tail. The extracellular Ig-like domain of NKp46 ( 22-255aa) was overexpressed in E. coli, and purified by FPLC gel-filtration chromatography, after refolding of the isolated inclusion bodies in a redox buffer.

#### **Amino acid Sequence**

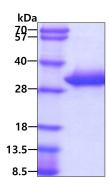
MQQQTLPKPF IWAEPHFMVP KEKQVTICCQ GNYGAVEYQL HFEGSLFAVD RPKPPERINK VKFYIPDMNS RMAGQYSCIY RVGELWSEPS NLLDLVVTEM YDTPTLSVHP GPEVISGEEV TFYCRLDTAT SMFLLLKEGR SSHVQRGYGK VQAEFPLGPV TTAHRGTYRC FGSYNNHAWS FPSEPVKLLV TGDIENTSLA PEDPTFPADT WGTYLLTTET GLQKDHALWD HTAQN

#### **General References**

Foster CE., et al. (2003) J.Biol.Chem. 278(46), 46081-6. Vankayalapati R., et al. (2002) J. Immunol. 168(7), 3451-7

## **DATA**

#### **SDS-PAGE**



3ug by SDS-PAGE under reducing condition and visualized by coomassie blue stain.

