

## Datasheet for ABIN3099649

# CD94 Protein (AA 1-179) (Strep Tag)



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Quantity:	250 μg
Target:	CD94 (KLRD1)
Protein Characteristics:	AA 1-179
Origin:	Human
Source:	Cell-free protein synthesis (CFPS)
Protein Type:	Recombinant
Purification tag / Conjugate:	This CD94 protein is labelled with Strep Tag.
Application:	Western Blotting (WB), SDS-PAGE (SDS), ELISA
Product Details	
Brand:	AliCE®
Brand: Sequence:	AliCE®  MAVFKTTLWR LISGTLGIIC LSLMSTLGIL LKNSFTKLSI EPAFTPGPNI ELQKDSDCCS  CQEKWVGYRC NCYFISSEQK TWNESRHLCA SQKSSLLQLQ NTDELDFMSS SQQFYWIGLS  YSEEHTAWLW ENGSALSQYL FPSFETFNTK NCIAYNPNGN ALDESCEDKN RYICKQQLI
	MAVFKTTLWR LISGTLGIIC LSLMSTLGIL LKNSFTKLSI EPAFTPGPNI ELQKDSDCCS CQEKWVGYRC NCYFISSEQK TWNESRHLCA SQKSSLLQLQ NTDELDFMSS SQQFYWIGLS
	MAVFKTTLWR LISGTLGIIC LSLMSTLGIL LKNSFTKLSI EPAFTPGPNI ELQKDSDCCS CQEKWVGYRC NCYFISSEQK TWNESRHLCA SQKSSLLQLQ NTDELDFMSS SQQFYWIGLS YSEEHTAWLW ENGSALSQYL FPSFETFNTK NCIAYNPNGN ALDESCEDKN RYICKQQLI
	MAVFKTTLWR LISGTLGIIC LSLMSTLGIL LKNSFTKLSI EPAFTPGPNI ELQKDSDCCS CQEKWVGYRC NCYFISSEQK TWNESRHLCA SQKSSLLQLQ NTDELDFMSS SQQFYWIGLS YSEEHTAWLW ENGSALSQYL FPSFETFNTK NCIAYNPNGN ALDESCEDKN RYICKQQLI Sequence without tag. The proposed Strep-Tag is based on experience s with the expression
	MAVFKTTLWR LISGTLGIIC LSLMSTLGIL LKNSFTKLSI EPAFTPGPNI ELQKDSDCCS CQEKWVGYRC NCYFISSEQK TWNESRHLCA SQKSSLLQLQ NTDELDFMSS SQQFYWIGLS YSEEHTAWLW ENGSALSQYL FPSFETFNTK NCIAYNPNGN ALDESCEDKN RYICKQQLI Sequence without tag. The proposed Strep-Tag is based on experience s with the expression system, a different complexity of the protein could make another tag necessary. In case you

reported (not tested by us and not guaranteed).

Durification:

• State-of-the-art algorithm used for plasmid design (Gene synthesis).

This protein is a **made-to-order protein** and will be made for the first time for your order. Our experts in the lab try to ensure that you receive soluble protein.

The big advantage of ordering our **made-to-order proteins** in comparison to ordering custom made proteins from other companies is that there is no financial obligation in case the protein cannot be expressed or purified.

#### Expression System:

- ALiCE®, our Almost Living Cell-Free Expression System is based on a lysate obtained from Nicotiana tabacum c.v.. This contains all the protein expression machinery needed to produce even the most difficult-to-express proteins, including those that require posttranslational modifications.
- During lysate production, the cell wall and other cellular components that are not required for
  protein production are removed, leaving only the protein production machinery and the
  mitochondria to drive the reaction. During our lysate completion steps, the additional
  components needed for protein production (amino acids, cofactors, etc.) are added to
  produce something that functions like a cell, but without the constraints of a living system all that's needed is the DNA that codes for the desired protein!

#### Concentration:

- The concentration of our recombinant proteins is measured using the absorbance at 280nm.
- The protein's absorbance will be measured against its specific reference buffer.
- · We use the Expasy's ProtParam tool to determine the absorption coefficient of each protein.

One step Strop tog purification of proteins expressed in Almost Living Call Free Expression

Purification:	One-step Strep-tag purification of proteins expressed in Almost Living Cell-Free Expression System (AliCE®).
Purity:	> 70-80 % as determined by SDS PAGE, Western Blot and analytical SEC (HPLC).
Grade:	custom-made
Target Details	
Target:	CD94 (KLRD1)
Alternative Name:	KLRD1 (KLRD1 Products)
Background:	Natural killer cells antigen CD94 (KP43) (Killer cell lectin-like receptor subfamily D member 1) (NK cell receptor) (CD antigen CD94),FUNCTION: Immune receptor involved in self-nonself

discrimination. In complex with KLRC1 or KLRC2 on cytotoxic and regulatory lymphocyte subsets, recognizes non-classical major histocompatibility (MHC) class lb molecule HLA-E loaded with self-peptides derived from the signal sequence of classical MHC class la and non-classical MHC class lb molecules (PubMed:9486650, PubMed:10023772, PubMed:18083576, PubMed:18064301, PubMed:9754572, PubMed:37264229). Enables cytotoxic cells to monitor the expression of MHC class I molecules in healthy cells and to tolerate self (PubMed:9430220, PubMed:12387742, PubMed:18064301). Primarily functions as a ligand binding subunit as it lacks the capacity to signal. {ECO:0000269|PubMed:10023772,

ECO:0000269|PubMed:12387742, ECO:0000269|PubMed:18064301,

ECO:0000269|PubMed:18083576, ECO:0000269|PubMed:37264229,

ECO:0000269|PubMed:9430220, ECO:0000269|PubMed:9486650,

ECO:0000269|PubMed:9754572}., FUNCTION: KLRD1-KLRC1 acts as an immune inhibitory receptor. Key inhibitory receptor on natural killer (NK) cells that regulates their activation and effector functions (PubMed:9486650, PubMed:9430220, PubMed:9485206,

PubMed:30860984). Dominantly counteracts T cell receptor signaling on a subset of memory/effector CD8-positive T cells as part of an antigen-driven response to avoid autoimmunity (PubMed:12387742). On intraepithelial CD8-positive gamma-delta regulatory T cells triggers TGFB1 secretion, which in turn limits the cytotoxic programming of intraepithelial CD8-positive alpha-beta T cells, distinguishing harmless from pathogenic antigens (PubMed:18064301). In HLA-E-rich tumor microenvironment, acts as an immune inhibitory checkpoint and may contribute to progressive loss of effector functions of NK cells and tumor-specific T cells, a state known as cell exhaustion (PubMed:30503213, PubMed:30860984). Upon HLA-E-peptide binding, transmits intracellular signals through KLRC1 immunoreceptor tyrosine-based inhibition motifs (ITIMs) by recruiting INPP5D/SHIP-1 and INPPL1/SHIP-2 tyrosine phosphatases to ITIMs, and ultimately opposing signals transmitted by activating receptors through dephosphorylation of proximal signaling molecules (PubMed:9485206, PubMed:12165520). {ECO:0000269|PubMed:12165520, ECO:0000269|PubMed:12387742, ECO:0000269|PubMed:18064301, ECO:0000269|PubMed:30503213,

ECO:0000269|PubMed:30860984, ECO:0000269|PubMed:9430220,

ECO:0000269|PubMed:9485206, ECO:0000269|PubMed:9486650}., FUNCTION: KLRD1-KLRC2 acts as an immune activating receptor (PubMed:9655483, PubMed:15940674). On cytotoxic lymphocyte subsets recognizes HLA-E loaded with signal sequence-derived peptides from non-classical MHC class Ib HLA-G molecules, likely playing a role in the generation and effector functions of adaptive NK cells and in maternal-fetal tolerance during pregnancy (PubMed:9754572, PubMed:30134159). Regulates the effector functions of terminally differentiated cytotoxic lymphocyte subsets, and in particular may play a role in adaptive NK

phosphorylation of proximal signaling molecules and cell activation (PubMed:9655483, PubMed:15940674). {ECO:0000269|PubMed:15940674, ECO:0000269|PubMed:20952657, ECO:0000269|PubMed:21825173, ECO:0000269|PubMed:30134159, ECO:0000269|PubMed:9655483, ECO:0000269|PubMed:9754572}., FUNCTION: (Microbial infection) Viruses like human cytomegalovirus have evolved an escape mechanism whereby virus-induced down-regulation of host MHC class I molecules is coupled to the binding of viral peptides to HLA-E, restoring HLA-E expression and inducing HLA-E-dependent NK cell immune

tolerance to infected cells. Recognizes HLA-E in complex with human cytomegalovirus UL40-

derived peptide (VMAPRTLIL) and inhibits NK cell cytotoxicity.

cell response to viral infection (PubMed:21825173, PubMed:20952657). Upon HLA-E-peptide binding, transmits intracellular signals via the adapter protein TYROBP/DAP12, triggering the

{ECO:0000269|PubMed:10669413, ECO:0000269|PubMed:23335510}., FUNCTION: (Microbial infection) May recognize HLA-E in complex with HIV-1 gag/Capsid protein p24-derived peptide (AISPRTLNA) on infected cells and may inhibit NK cell cytotoxicity, a mechanism that allows HIV-1 to escape immune recognition. {ECO:0000269|PubMed:15751767}., FUNCTION: (Microbial infection) Upon SARS-CoV-2 infection, may contribute to functional exhaustion of cytotoxic NK cells and CD8-positive T cells (PubMed:32859121). On NK cells, may recognize HLA-E in complex with SARS-CoV-2 S/Spike protein S1-derived peptide (LQPRTFLL) expressed on the surface of lung epithelial cells, inducing NK cell exhaustion and dampening antiviral immune surveillance (PubMed:32859121). {ECO:0000269|PubMed:32859121}.

Molecular Weight:

20.5 kDa

UniProt:

013241

### **Application Details**

Application Notes:

In addition to the applications listed above we expect the protein to work for functional studies as well. As the protein has not been tested for functional studies yet we cannot offer a guarantee though.

Comment:

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## **Application Details**

	components needed for protein production (amino acids, cofactors, etc.) are added to produce something that functions like a cell, but without the constraints of a living system - all that's needed is the DNA that codes for the desired protein!
Restrictions:	For Research Use only
Handling	
Format:	Liquid
Buffer:	The buffer composition is at the discretion of the manufacturer.  Standard Storage Buffer: PBS pH 7.4, 10 % Glycerol <b>Might differ depending on protein.</b>
Handling Advice:	Avoid repeated freeze-thaw cycles.
Storage:	-80 °C
Storage Comment:	Store at -80°C.
Expiry Date:	12 months