# abcam

## Product datasheet

## Recombinant Human CIRP protein ab106903

## 1 Image

**Description** 

Product name Recombinant Human CIRP protein

Purity > 95 % SDS-PAGE.

ab106903 is purified using conventional chromatography techniques.

**Expression system** Escherichia coli

Accession Q14011

Protein length Full length protein

Animal free No

Nature Recombinant

**Species** Human

**Sequence** MGSSHHHHHHSSGLVPRGSHMASDEGKLFVGGLSFDTN

**EQSLEQVFSKYG** 

QISEVVVVKDRETQRSRGFGFVTFENIDDAKDAMMAMNG

KSVDGRQIRVD

QAGKSSDNRSRGYRGGSAGGRGFFRGGRGRGRGFSRG

**GGDRGYGGNRFES** 

RSGGYGGSRDYYSSRSQSGGYSDRSSGGSYRDSYDSYA

**THNE** 

Predicted molecular weight 21 kDa including tags

Amino acids 1 to 172

Tags His tag N-Terminus

## **Specifications**

Our **Abpromise guarantee** covers the use of **ab106903** in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

**Applications** Mass Spectrometry

SDS-PAGE

Mass spectrometry MALDI-TOF

Form Liquid

**Preparation and Storage** 

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#### Stability and Storage

Shipped at  $4\,^\circ\text{C}$  . Upon delivery aliquot and store at -20 $^\circ\text{C}$  or -80 $^\circ\text{C}$  . Avoid repeated freeze / thaw

cycles.

pH: 8.00

Constituents: 0.02% DTT, 0.32% Tris HCI, 50% Glycerol (glycerin, glycerine), 0.58% Sodium

chloride

#### **General Info**

#### **Function** Cold-inducible mRNA binding protein that plays a protective role in the genotoxic stress response

by stabilizing transcripts of genes involved in cell survival. Acts as a translational activator. Seems to play an essential role in cold-induced suppression of cell proliferation. Binds specifically to the 3'-untranslated regions (3'-UTRs) of stress-responsive transcripts RPA2 and TXN. Acts as a translational repressor (By similarity). Promotes assembly of stress granules (SGs), when overexpressed.

Tissue specificity Ubiquitous.

Sequence similarities Contains 1 RRM (RNA recognition motif) domain.

**Domain**Both the RRM domain and the arginine, glycine (RGG) rich domain are necessary for binding to

the TXN 3'-untranslated region. Both the RRM domain and the arginine, glycine (RGG) rich domain (RGG repeats) are necessary for optimal recruitment into SGs upon cellular stress. The

C-terminal domain containing RGG repeats is necessary for translational repression.

Post-translational modifications

Methylated on arginine residues. Methylation of the RGG motifs is a prerequisite for recruitment

into SGs.

Phosphorylated by CK2, GSK3A and GSK3B. Phosphorylation by GSK3B increases RNA-

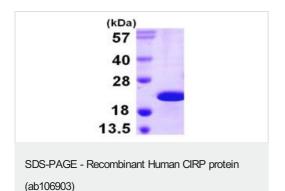
binding activity to the TXN 3'-UTR transcript upon exposure to UV radiation.

**Cellular localization** Nucleus > nucleoplasm. Cytoplasm. Translocates from the nucleus to the cytoplasm after

exposure to UV radiation. Translocates from the nucleus to the cytoplasm into stress granules upon various cytoplasmic stresses, such as osmotic and heat shocks. Its recruitment into stress

granules occurs in the absence of TIAR proteins.

#### **Images**



15% SDS-PAGE showing ab106903 at approximately 20.8kDa ( $3\mu g$ ).

 $\textbf{Please note:} \ \ \textbf{All products are "FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES"}$ 

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