# abcam

# Product datasheet

# Recombinant Human SIRT4 protein ab101713

# 1 Image

**Description** 

Product name Recombinant Human SIRT4 protein

Purity > 70 % SDS-PAGE.

Expression system Escherichia coli

Accession Q9Y6E7

Protein length Full length protein

Animal free No

Nature Recombinant

**Species** Human

**Sequence** MKMSFALTFRSAKGRWIANPSQPCSKASIGLFVPASPPL

**DPEKVKELQRF** 

 ${\tt ITLSKRLLVMTGAGISTESGIPDYRSEKVGLYARTDRRPIQH}$ 

**GDFVRSAP** 

IRQRYWARNFVGWPQFSSHQPNPAHWALSTWEKLGKLY

WLVTQNVDALHT

KAGSRRLTELHGCMDRVLCLDCGEQTPRGVLQERFQVL

**NPTWSAEAHGLA** 

PDGDVFLSEEQVRSFQVPTCVQCGGHLKPDVVFFGDTV

NPDKVDFVHKRV

KEADSLLVVGSSLQVYSGYRFILTAWEKKLPIAILNIGPTRS

DDLACLKL NSRCGELLPLIDPC

Predicted molecular weight 62 kDa including tags

Amino acids 1 to 314

Tags GST tag N-Terminus

#### **Specifications**

Our Abpromise guarantee covers the use of ab101713 in the following tested applications.

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

Applications SDS-PAGE

Form Liquid

1

#### **Preparation and Storage**

# **Stability and Storage**

Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles.

pH: 7.50

Constituents: 0.307% Glutathione, 0.00174% PMSF, 0.00385% DTT, 0.79% Tris HCI, 0.00292%

EDTA, 25% Glycerol (glycerin, glycerine), 0.87% Sodium chloride

#### **General Info**

#### **Function**

Acts as NAD-dependent protein lipoamidase, ADP-ribosyl transferase and deacetylase. Catalyzes more efficiently removal of lipoyl- and biotinyl- than acetyl-lysine modifications. Inhibits the pyruvate dehydrogenase complex (PDH) activity via the enzymatic hydrolysis of the lipoamide cofactor from the E2 component, DLAT, in a phosphorylation-independent manner (PubMed:25525879). Catalyzes the transfer of ADP-ribosyl groups onto target proteins, including mitochondrial GLUD1, inhibiting GLUD1 enzyme activity. Acts as a negative regulator of mitochondrial glutamine metabolism by mediating mono ADP-ribosylation of GLUD1: expressed in response to DNA damage and negatively regulates anaplerosis by inhibiting GLUD1, leading to block metabolism of glutamine into tricarboxylic acid cycle and promoting cell cycle arrest (PubMed:16959573, PubMed:17715127). In response to mTORC1 signal, SIRT4 expression is repressed, promoting anaplerosis and cell proliferation. Acts as a tumor suppressor (PubMed:23562301, PubMed:23663782). Also acts as a NAD-dependent protein deacetylase: mediates deacetylation of 'Lys-471' of MLYCD, inhibiting its activity, thereby acting as a regulator of lipid homeostasis (By similarity). Controls fatty acid oxidation by inhibiting PPARA transcriptional activation. Impairs SIRT1:PPARA interaction probably through the regulation of NAD(+) levels (PubMed:24043310). Down-regulates insulin secretion.

### Tissue specificity

Detected in vascular smooth muscle and striated muscle. Detected in insulin-producing beta-cells in pancreas islets of Langerhans (at protein level). Widely expressed. Weakly expressed in leukocytes and fetal thymus.

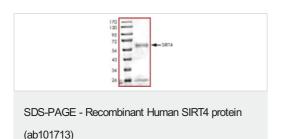
Sequence similarities

Belongs to the sirtuin family. Class II subfamily. Contains 1 deacetylase sirtuin-type domain.

**Cellular localization** 

Mitochondrion matrix.

#### **Images**



The putity of ab101713 was determined to be 90% by densitometry.

Approximate MWt: 62kDa

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