abcam

Product datasheet

Recombinant Human TRAF1 protein ab95858

1 Image

Description

Product name Recombinant Human TRAF1 protein

Purity > 95 % SDS-PAGE.

ab95858 is purified using conventional chromatography techniques.

Expression system Escherichia coli

Accession Q13077

Protein length Protein fragment

Animal free No

Nature Recombinant

Species Human

Sequence MGSSHHHHHH SSGLVPRGSH MDGTFLWKIT

NVTRRCHESA CGRTVSLFSP AFYTAKYGYK

LCLRLYLNGD GTGKRTHLSL FIVIMRGEYD ALLPWPFRNK

VTFMLLDQNN REHAIDAFRP DLSSASFQRP QSETNVASGC PLFFPLSKLQ SPKHAYVKDD

TMFLKCIVET ST

Predicted molecular weight 20 kDa including tags

Amino acids 266 to 416

Tags His tag N-Terminus

Specifications

Our **Abpromise guarantee** covers the use of **ab95858** 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

Mass Spectrometry

Form Liquid

Preparation and Storage

Stability and Storage Shipped at 4°C. Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze / thaw

cycles.

pH: 8.00

Constituents: 0.0154% DTT, 0.316% Tris HCl, 20% Glycerol (glycerin, glycerine), 0.58% Sodium chloride

General Info

Function	Adapter molecule that regulates the activation of NF-kappa-B and JNK. Plays a role in the

regulation of cell survival and apoptosis. The heterotrimer formed by TRAF1 and TRAF2 is part of a E3 ubiquitin-protein ligase complex that promotes ubiquitination of target proteins, such as MAP3K14. The TRAF1/TRAF2 complex recruits the antiapoptotic E3 protein-ubiquitin ligases

BIRC2 and BIRC3 to TNFRSF1B/TNFR2.

Sequence similarities Contains 1 MATH domain.

Domain The coiled coil domain mediates homo- and hetero-oligomerization.

The MATH/TRAF domain binds to receptor cytoplasmic domains.

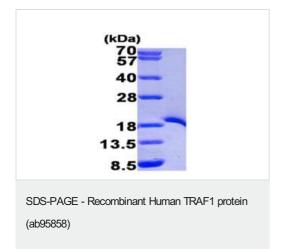
Cleavage by CASP8 liberates a C-terminal fragment that promotes apoptosis and inhibits the

activation of NF-kappa-B in response to TNF signaling.

Post-translational modifications

Polyubiquitinated by BIRC2 and/or BIRC3, leading to its subsequent proteasomal degradation.

Images



15% SDS-PAGE showing ab95858 at approximately 19.5kDa ($3\mu g$).

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