

#### ZFP36 Antibody (Center) Blocking Peptide Synthetic peptide

Catalog # BP8952c

## Specification

# ZFP36 Antibody (Center) Blocking Peptide - Product Information

Primary Accession

<u>P26651</u>

# ZFP36 Antibody (Center) Blocking Peptide - Additional Information

Gene ID 7538

#### **Other Names**

Tristetraprolin, TTP, G0/G1 switch regulatory protein 24, Growth factor-inducible nuclear protein NUP475, Protein TIS11A, TIS11, Zinc finger protein 36 homolog, Zfp-36, ZFP36, G0S24, RNF162A, TIS11A, TTP

## Target/Specificity

The synthetic peptide sequence used to generate the antibody <a href=/products/AP8952c>AP8952c</a> was selected from the Center region of human ZFP36. A 10 to 100 fold molar excess to antibody is recommended. Precise conditions should be optimized for a particular assay.

## Format

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

## Storage

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

## **Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

# ZFP36 Antibody (Center) Blocking Peptide - Protein Information

## Name ZFP36 (<u>HGNC:12862</u>)

Function

Zinc-finger RNA-binding protein that destabilizes several cytoplasmic AU-rich element (ARE)-containing mRNA transcripts by promoting their poly(A) tail removal or deadenylation, and hence provide a mechanism for attenuating protein synthesis (PubMed:<a href="http://www.uniprot.org/citations/10330172" target="\_blank">10330172</a>, PubMed:<a href="http://www.uniprot.org/citations/10751406" target="\_blank">10751406</a>, PubMed:<a href="http://www.uniprot.org/citations/10751406" target="\_blank">11279239</a>, PubMed:<a href="http://www.uniprot.org/citations/11279239" target="\_blank">11279239</a>, PubMed:<a href="http://www.uniprot.org/citations/12115244" target="\_blank">12115244</a>, PubMed:<a href="http://www.uniprot.org/citations/12115244" target="\_blank">12115244</a>, PubMed:<a href="http://www.uniprot.org/citations/12748283" target="\_blank">12748283</a>, PubMed:<a href="http://www.uniprot.org/citations/15187101" target="\_blank">15187101</a>, PubMed:<a href="http://www.uniprot.org/citations/15634918" target="\_blank">15634918</a>, PubMed:<a



href="http://www.uniprot.org/citations/16702957" target=" blank">16702957</a>, PubMed:<a href="http://www.uniprot.org/citations/17030620" target=" blank">17030620</a>, PubMed:<a href="http://www.uniprot.org/citations/20221403" target="\_blank">20221403</a>, PubMed:<a href="http://www.uniprot.org/citations/20702587" target="\_blank">20702587</a>, PubMed:<a href="http://www.uniprot.org/citations/21775632" target=" blank">21775632</a>, PubMed:<a href="http://www.uniprot.org/citations/23644599" target=" blank">23644599</a>, PubMed:<a href="http://www.uniprot.org/citations/25815583" target=" blank">25815583</a>, PubMed:<a href="http://www.uniprot.org/citations/27193233" target=" blank">27193233</a>, PubMed:<a href="http://www.uniprot.org/citations/31439631" target=" blank">31439631</a>, PubMed:<a href="http://www.uniprot.org/citations/9703499" target="\_blank">9703499</a>). Acts as an 3'-untranslated region (UTR) ARE mRNA-binding adapter protein to communicate signaling events to the mRNA decay machinery (PubMed:<a href="http://www.uniprot.org/citations/15687258" target=" blank">15687258</a>, PubMed:<a href="http://www.uniprot.org/citations/23644599" target=" blank">23644599</a>). Recruits deadenylase CNOT7 (and probably the CCR4-NOT complex) via association with CNOT1, and hence promotes ARE-mediated mRNA deadenylation (PubMed:<a href="http://www.uniprot.org/citations/23644599" target=" blank">23644599</a>). Functions also by recruiting components of the cytoplasmic RNA decay machinery to the bound ARE-containing mRNAs (PubMed:<a href="http://www.uniprot.org/citations/11719186" target=" blank">11719186</a>, PubMed:<a href="http://www.uniprot.org/citations/12748283" target=" blank">12748283</a>, PubMed:<a href="http://www.uniprot.org/citations/15687258" target=" blank">15687258</a>, PubMed:<a href="http://www.uniprot.org/citations/16364915" target=" blank">16364915</a>). Self regulates by destabilizing its own mRNA (PubMed:<a href="http://www.uniprot.org/citations/15187101" target=" blank">15187101</a>). Binds to 3'-UTR ARE of numerous mRNAs and of its own mRNA (PubMed: <a href="http://www.uniprot.org/citations/10330172" target="\_blank">10330172</a>, PubMed:<a href="http://www.uniprot.org/citations/10751406" target=" blank">10751406</a>, PubMed:<a href="http://www.uniprot.org/citations/12115244" target=" blank">12115244</a>, PubMed:<a href="http://www.uniprot.org/citations/15187101" target=" blank">15187101</a>, PubMed:<a href="http://www.uniprot.org/citations/15634918" target="\_blank">15634918</a>, PubMed:<a href="http://www.uniprot.org/citations/16702957" target=" blank">16702957</a>, PubMed:<a href="http://www.uniprot.org/citations/17030620" target=" blank">17030620</a>, PubMed:<a href="http://www.uniprot.org/citations/19188452" target="\_blank">19188452</a>, PubMed:<a href="http://www.uniprot.org/citations/20221403" target=" blank">20221403</a>, PubMed:<a href="http://www.uniprot.org/citations/20702587" target=" blank">20702587</a>, PubMed:<a href="http://www.uniprot.org/citations/21775632" target=" blank">21775632</a>, PubMed:<a href="http://www.uniprot.org/citations/25815583" target=" blank">25815583</a>). Plays a role in anti-inflammatory responses; suppresses tumor necrosis factor (TNF)-alpha production by stimulating ARE-mediated TNF-alpha mRNA decay and several other inflammatory ARE- containing mRNAs in interferon (IFN)- and/or lipopolysaccharide (LPS)- induced macrophages (By similarity). Also plays a role in the regulation of dendritic cell maturation at the post-transcriptional level, and hence operates as part of a negative feedback loop to limit the inflammatory response (PubMed:<a href="http://www.uniprot.org/citations/18367721" target=" blank">18367721</a>). Promotes ARE-mediated mRNA decay of hypoxia-inducible factor HIF1A mRNA during the response of endothelial cells to hypoxia (PubMed:<a href="http://www.uniprot.org/citations/21775632" target=" blank">21775632</a>). Positively regulates early adipogenesis of preadipocytes by promoting ARE-mediated mRNA decay of immediate early genes (IEGs) (By similarity). Negatively regulates hematopoietic/erythroid cell differentiation by promoting ARE-mediated mRNA decay of the transcription factor STAT5B mRNA (PubMed: <a href="http://www.uniprot.org/citations/20702587" target=" blank">20702587</a>). Plays a role in maintaining skeletal muscle satellite cell guiescence by promoting ARE-mediated mRNA decay of the myogenic determination factor MYOD1 mRNA (By similarity). Associates also with and regulates the expression of non-ARE-containing target mRNAs at the post-transcriptional level, such as MHC class I mRNAs (PubMed:<a href="http://www.uniprot.org/citations/18367721" target=" blank">18367721</a>). Participates in association with argonaute RISC catalytic

components in the ARE-mediated mRNA decay mechanism; assists microRNA (miRNA) targeting ARE-containing mRNAs (PubMed:<a href="http://www.uniprot.org/citations/15766526" target=" blank">15766526</a>). May also play a role in the regulation of cytoplasmic mRNA



decapping; enhances decapping of ARE-containing RNAs, in vitro (PubMed:<a href="http://www.uniprot.org/citations/16364915" target="\_blank">16364915</a>). Involved in the delivery of target ARE-mRNAs to processing bodies (PBs) (PubMed:<a href="http://www.uniprot.org/citations/17369404" target="\_blank">17369404</a>). In addition to its cytosolic mRNA-decay function, affects nuclear pre-mRNA processing (By similarity). Negatively regulates nuclear poly(A)-binding protein PABPN1-stimulated polyadenylation activity on ARE-containing pre-mRNA during LPS- stimulated macrophages (By similarity). Also involved in the regulation of stress granule (SG) and P-body (PB) formation and fusion (By similarity). Plays a role in the regulation of keratinocyte proliferation, differentiation and apoptosis (PubMed:<a href="http://www.uniprot.org/citations/27182009" target="\_blank">27182009</a>). Plays a role as a tumor suppressor by inhibiting cell proliferation in breast cancer cells (PubMed:<a href="http://www.uniprot.org/citations/26926077" target="\_blank">26926077</a>).

#### **Cellular Location**

Nucleus. Cytoplasm. Cytoplasmic granule. Cytoplasm, P-body. Note=Shuttles between nucleus and cytoplasm in a CRM1-dependent manner (By similarity). Localized predominantly in the cytoplasm in a p38 MAPK- and YWHAB-dependent manner (By similarity). Colocalizes with SH3KBP1 and MAP3K4 in the cytoplasm (PubMed:20221403). Component of cytoplasmic stress granules (SGs) (By similarity). Localizes to cytoplasmic stress granules upon energy starvation (PubMed:15014438). Localizes in processing bodies (PBs) (PubMed:17369404). Excluded from stress granules in a phosphorylation MAPKAPK2-dependent manner (By similarity). Shuttles in and out of both cytoplasmic P-body and SGs (By similarity) {ECO:0000250|UniProtKB:P22893, ECO:0000269|PubMed:15014438, ECO:0000269|PubMed:17369404, ECO:0000269|PubMed:20221403}

#### **Tissue Location**

Expressed in both basal and suprabasal epidermal layers (PubMed:27182009). Expressed in epidermal keratinocytes (PubMed:27182009). Expressed strongly in mature dendritic cells (PubMed:18367721). Expressed in immature dendritic cells (at protein level) (PubMed:18367721).

# ZFP36 Antibody (Center) Blocking Peptide - Protocols

Provided below are standard protocols that you may find useful for product applications.

#### <u>Blocking Peptides</u>

## ZFP36 Antibody (Center) Blocking Peptide - Images

## ZFP36 Antibody (Center) Blocking Peptide - Background

ZFP36 is probable regulatory protein with a novel zinc finger structure involved in regulating the response to growth factors. Has been experimentally shown to be able to bind zinc.

## ZFP36 Antibody (Center) Blocking Peptide - References

Lee,H.H., et.al., Int. J. Cancer 126 (8), 1817-1827 (2010)Datta,S., eet.al., J. Immunol. 184 (3), 1484-1491 (2010)