

## **ZBTB7A Antibody**

Catalog # ASC11264

## **Specification**

## **ZBTB7A Antibody - Product Information**

Application
Primary Accession
Other Accession
Reactivity
Host
Clonality
Isotype
Application Notes

WB, IHC-P, IF, E

095365

AAH84568, 7705375

Human, Mouse

Rabbit

Polyclonal

IgG

ZBTB7A antibody can be used for detection of ZBTB7A by Western blot at 1 - 2 μg/mL.

Antibody can also be used for immunohistochemistry starting at 2.5 μg/mL. For immunofluorescence start at 20 μg/mL.

#### **ZBTB7A Antibody - Additional Information**

Gene ID 51341

# **Target/Specificity**

ZBTB7A; At least two isoforms of ZBTB7A are known to exist; this antibody will recognize both isoforms. This antibody is predicted to not cross-react with other ZBTB protein family members.

#### **Reconstitution & Storage**

ZBTB7A antibody can be stored at 4°C for three months and -20°C, stable for up to one year. As with all antibodies care should be taken to avoid repeated freeze thaw cycles. Antibodies should not be exposed to prolonged high temperatures.

#### **Precautions**

ZBTB7A Antibody is for research use only and not for use in diagnostic or therapeutic procedures.

# **ZBTB7A Antibody - Protein Information**

Name ZBTB7A (HGNC:18078)

#### **Function**

Transcription factor that represses the transcription of a wide range of genes involved in cell proliferation and differentiation (PubMed:<a href="http://www.uniprot.org/citations/14701838" target="\_blank">14701838</a>, PubMed:<a href="http://www.uniprot.org/citations/17595526" target="\_blank">17595526</a>, PubMed:<a href="http://www.uniprot.org/citations/20812024" target="\_blank">20812024</a>, PubMed:<a href="http://www.uniprot.org/citations/25514493" target="\_blank">25514493</a>, PubMed:<a href="http://www.uniprot.org/citations/26455326" target="\_blank">26455326</a>, PubMed:<a href="http://www.uniprot.org/citations/26816381" target="\_blank">26816381</a>, PubMed:<a href="http://www.uniprot.org/citations/26816381" target="\_blank">26816381</a>). Directly and specifically binds to the consensus sequence 5'-[GA][CA]GACCCCCCCCC3' and represses transcription both by regulating the organization of



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chromatin and through the direct recruitment of transcription factors to gene regulatory regions (PubMed:<a href="http://www.uniprot.org/citations/12004059" target=" blank">12004059</a>, PubMed:<a href="http://www.uniprot.org/citations/17595526" target=" blank">17595526</a>, PubMed:<a href="http://www.uniprot.org/citations/20812024" target="\_blank">20812024</a>, PubMed:<a href="http://www.uniprot.org/citations/25514493" target="blank">25514493</a>, PubMed:<a href="http://www.uniprot.org/citations/26816381" target=" blank">26816381</a>). Negatively regulates SMAD4 transcriptional activity in the TGF-beta signaling pathway through these two mechanisms (PubMed: <a href="http://www.uniprot.org/citations/25514493" target=" blank">25514493</a>). That is, recruits the chromatin regulator HDAC1 to the SMAD4-DNA complex and in parallel prevents the recruitment of the transcriptional activators CREBBP and EP300 (PubMed:<a href="http://www.uniprot.org/citations/25514493" target=" blank">25514493</a>). Collaborates with transcription factors like RELA to modify the accessibility of gene transcription regulatory regions to secondary transcription factors (By similarity). Also directly interacts with transcription factors like SP1 to prevent their binding to DNA (PubMed:<a href="http://www.uniprot.org/citations/12004059" target=" blank">12004059</a>). Functions as an androgen receptor/AR transcriptional corepressor by recruiting NCOR1 and NCOR2 to the androgen response elements/ARE on target genes (PubMed: <a href="http://www.uniprot.org/citations/20812024" target="\_blank">20812024</a>). Thereby, negatively regulates androgen receptor signaling and androgen- induced cell proliferation (PubMed:<a href="http://www.uniprot.org/citations/20812024" target=" blank">20812024</a>). Involved in the switch between fetal and adult globin expression during erythroid cells maturation (PubMed:<a href="http://www.uniprot.org/citations/26816381" target=" blank">26816381</a>). Through its interaction with the NuRD complex regulates chromatin at the fetal globin genes to repress their transcription (PubMed: <a href="http://www.uniprot.org/citations/26816381" target=" blank">26816381</a>). Specifically represses the transcription of the tumor suppressor ARF isoform from the CDKN2A gene (By similarity). Efficiently abrogates E2F1-dependent CDKN2A transactivation (By similarity). Regulates chondrogenesis through the transcriptional repression of specific genes via a mechanism that also requires histone deacetylation (By similarity). Regulates cell proliferation through the transcriptional regulation of genes involved in glycolysis (PubMed: <a href="http://www.uniprot.org/citations/26455326" target=" blank">26455326</a>). Involved in adipogenesis through the regulation of genes involved in adipocyte differentiation (PubMed:<a href="http://www.uniprot.org/citations/14701838" target=" blank">14701838</a>). Plays a key role in the differentiation of lymphoid progenitors into B and T lineages (By similarity). Promotes differentiation towards the B lineage by inhibiting the T-cell instructive Notch signaling pathway through the specific transcriptional repression of Notch downstream target genes (By similarity). Also regulates osteoclast differentiation (By similarity). May also play a role, independently of its transcriptional activity, in double-strand break repair via classical non-homologous end joining/cNHEJ (By similarity). Recruited to double-strand break sites on damage DNA, interacts with the DNA-dependent protein kinase complex and directly regulates its stability and activity in DNA repair (By similarity). May also modulate the splicing activity of KHDRBS1 toward BCL2L1 in a mechanism which is histone deacetylase-dependent and thereby negatively regulates the pro-apoptotic effect of KHDRBS1 (PubMed: <a href="http://www.uniprot.org/citations/24514149" target=" blank">24514149</a>).

#### **Cellular Location**

Nucleus. Note=Recruited to double-strand break sites of damaged DNA. {ECO:0000250|UniProtKB:088939}

# **Tissue Location**

Widely expressed (PubMed:9927193). In normal thymus, expressed in medullary epithelial cells and Hassle's corpuscles (at protein level) (PubMed:15662416). In tonsil, expressed in squamous epithelium and germinal center lymphocytes (at protein level) (PubMed:15662416). Up-regulated in a subset of lymphomas, as well as in a subset of breast, lung, colon, prostate and bladder carcinomas (at protein level) (PubMed:15662416). Expressed in adipose tissues (PubMed:14701838).

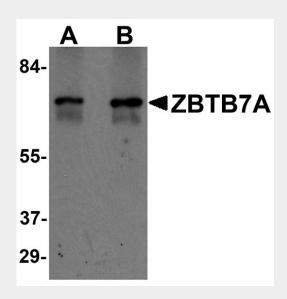


# **ZBTB7A Antibody - Protocols**

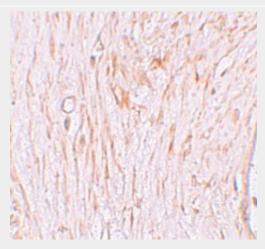
Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- <u>Immunohistochemistry</u>
- <u>Immunofluorescence</u>
- <u>Immunoprecipitation</u>
- Flow Cytomety
- Cell Culture

# **ZBTB7A Antibody - Images**

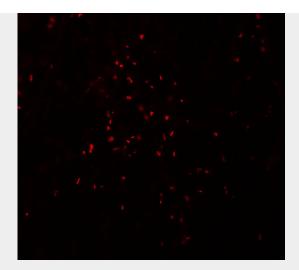


Western blot analysis of ZBTB7A in human ovary tissue lysate with ZBTB7A antibody at (A) 1 and (B) 2  $\mu$ g/mL.



Immunohistochemistry of ZBTB7A in human ovary tissue with ZBTB7A antibody at 2.5 μg/mL.





Immunofluorescence of ZBTB7A in human ovary tissue with ZBTB7A antibody at 20 µg/mL.

## **ZBTB7A Antibody - Background**

ZBTB7A Antibody: The ZBTB family of proteins is comprised of diverse zinc finger proteins that also contain a BTB (BR-C, ttk and bab) domain. ZBTB7A, also known as LRF, was initially identified as a potential target of the LAZ-3/BCL-6 oncogene. Later reports demonstrated that it could bind to the wild-type IST (inducer of short transcripts) elements of HIV-1 long terminal repeats (LTRs) and stimulate Tat activity on the HIV-1 LTR. More recently it ZBTB7A has been recognized as a proto-oncogene whose overexpression contributes to malignancy in breast cancer and functions as a co-repressor of the androgen receptor in prostate cancer cells.

### **ZBTB7A Antibody - References**

Filion GJP, Zhenilo S, Salozhin S, et al. A family of zinc finger proteins that bind methylated DNA and repress transcription. Mol. Cell. Biol. 2006; 26:169-81.

Davies JM, Hawe N, Kabarowski J, et al. Novel BTB/POZ domain zinc-finger protein, LRF, is a potential target of the LAZ-3/BCL-6 oncogene. Oncogene1999; 18:365-75.

Pendergrast PS, Wang C, Hernandez N, et al. FBI-1 can stimulate HIV-1 Tat activity and is targeted to a novel subnuclear domain that includes the Tat-P-TEFb-containing nuclear speckles. Mol. Cell. Biol.2002; 13:915-29.

Qu H, Qu D, Chen F, et al. ZBTB7 overexpression contributes to malignancy in breast cancer. Cancer Invest.2010; 28:672-8.