

RUVBL1 (TIP49) Antibody (Center)
Purified Rabbit Polyclonal Antibody (Pab)
Catalog # AP1922a**Specification**

RUVBL1 (TIP49) Antibody (Center) - Product Information

Application	FC, IHC-P, WB,E
Primary Accession	O9Y265
Other Accession	P60123 , P60122 , Q8AWW7
Reactivity	Human
Predicted	Zebrafish, Mouse, Rat
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	50228
Antigen Region	247-280

RUVBL1 (TIP49) Antibody (Center) - Additional Information**Gene ID** 8607**Other Names**

RuvB-like 1, 49 kDa TATA box-binding protein-interacting protein, 49 kDa TBP-interacting protein, 54 kDa erythrocyte cytosolic protein, ECP-54, INO80 complex subunit H, Nuclear matrix protein 238, NMP 238, Pontin 52, TIP49a, TIP60-associated protein 54-alpha, TAP54-alpha, RUVBL1, INO80H, NMP238, TIP49, TIP49A

Target/Specificity

This RUVBL1 (TIP49) antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 247-280 amino acids from the Central region of human RUVBL1 (TIP49).

Dilution

FC~~1:10~50

IHC-P~~1:10~50

WB~~1:1000

E~~Use at an assay dependent concentration.

Format

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is prepared by Saturated Ammonium Sulfate (SAS) precipitation followed by dialysis against PBS.

Storage

Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

Precautions

RUVBL1 (TIP49) Antibody (Center) is for research use only and not for use in diagnostic or therapeutic procedures.

RUVBL1 (TIP49) Antibody (Center) - Protein Information

Name RUVBL1 ([HGNC:10474](#))

Function Possesses single-stranded DNA-stimulated ATPase and ATP- dependent DNA helicase (3' to 5') activity; hexamerization is thought to be critical for ATP hydrolysis and adjacent subunits in the ring- like structure contribute to the ATPase activity (PubMed:[17157868](#), PubMed:[33205750](#)). Component of the NuA4 histone acetyltransferase complex which is involved in transcriptional activation of select genes principally by acetylation of nucleosomal histones H4 and H2A (PubMed:[14966270](#)). This modification may both alter nucleosome-DNA interactions and promote interaction of the modified histones with other proteins which positively regulate transcription (PubMed:[14966270](#)). This complex may be required for the activation of transcriptional programs associated with oncogene and proto-oncogene mediated growth induction, tumor suppressor mediated growth arrest and replicative senescence, apoptosis, and DNA repair (PubMed:[14966270](#)). The NuA4 complex ATPase and helicase activities seem to be, at least in part, contributed by the association of RUVBL1 and RUVBL2 with EP400. NuA4 may also play a direct role in DNA repair when recruited to sites of DNA damage (PubMed:[14966270](#)). Component of a SWR1-like complex that specifically mediates the removal of histone H2A.Z/H2AZ1 from the nucleosome (PubMed:[24463511](#)). Proposed core component of the chromatin remodeling INO80 complex which exhibits DNA- and nucleosome-activated ATPase activity and catalyzes ATP-dependent nucleosome sliding (PubMed:[16230350](#), PubMed:[21303910](#)). Plays an essential role in oncogenic transformation by MYC and also modulates transcriptional activation by the LEF1/TCF1-CTNNB1 complex (PubMed:[10882073](#), PubMed:[16014379](#)). Essential for cell proliferation (PubMed:[14506706](#)). May be able to bind plasminogen at cell surface and enhance plasminogen activation (PubMed:[11027681](#)).

Cellular Location

Nucleus matrix. Nucleus, nucleoplasm. Cytoplasm. Membrane Cytoplasm, cytoskeleton, microtubule organizing center, centrosome Dynein axonemal particle {ECO:0000250|UniProtKB:Q9DE26}. Note=Mainly localized in the nucleus, associated with nuclear matrix or in the nuclear cytosol, although it is also present in the cytoplasm and associated with the cell membranes. In prophase and prometaphase it is located at the centrosome and the branching microtubule spindles. After mitotic nuclear membrane disintegration it accumulates at the centrosome and sites of tubulin polymerization. As cells pass through metaphase and into telophase it is located close to the centrosome at the early phase of tubulin polymerization. In anaphase it accumulates at the zone of tubule interdigitation. In telophase it is found at polar tubule overlap, and it reappears at the site of chromosomal decondensation in the daughter cells

Tissue Location

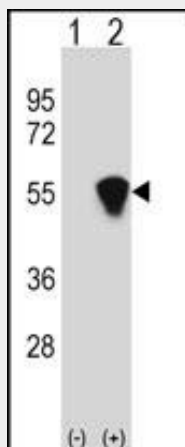
Ubiquitously expressed with high expression in heart, skeletal muscle and testis

RUVBL1 (TIP49) Antibody (Center) - Protocols

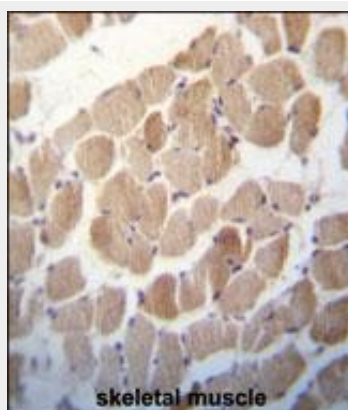
Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

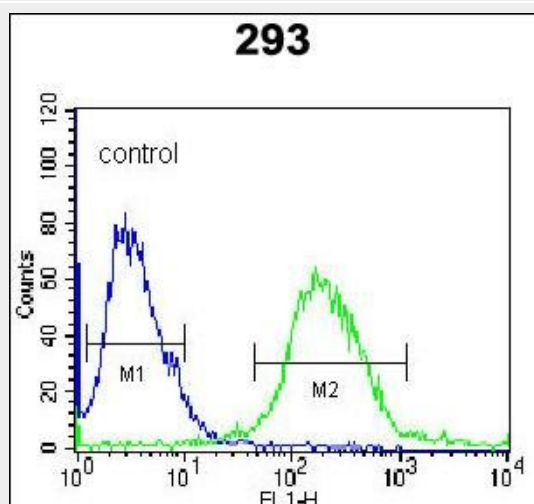
RUVBL1 (TIP49) Antibody (Center) - Images



Western blot analysis of RUVBL1 (arrow) using rabbit polyclonal RUVBL1 Antibody (Cat. #AP1922a). 293 cell lysates (2 ug/lane) either nontransfected (Lane 1) or transiently transfected (Lane 2) with the RUVBL1 gene.



RUVBL1 (TIP49) Antibody (Center) (Cat. #AP1922a) immunohistochemistry analysis in formalin fixed and paraffin embedded human skeletal muscle followed by peroxidase conjugation of the secondary antibody and DAB staining. This data demonstrates the use of RUVBL1 (TIP49) Antibody (Center) for immunohistochemistry. Clinical relevance has not been evaluated.



RUVBL1 (TIP49) Antibody (Center) (Cat. #AP1922a) flow cytometric analysis of 293 cells (right histogram) compared to a negative control cell (left histogram). FITC-conjugated goat-anti-rabbit secondary antibodies were used for the analysis.

RUVBL1 (TIP49) Antibody (Center) - Background

RUVBL1 possesses single-stranded DNA-stimulated ATPase and ATP-dependent DNA helicase (3' to 5') activity. It is a component of the NuA4 histone acetyltransferase complex which is involved in transcriptional activation of select genes principally by acetylation of nucleosomal histone H4 and H2A. This modification may both alter nucleosome - DNA interactions and promote interaction of the modified histones with other proteins which positively regulate transcription. This complex may be required for the activation of transcriptional programs associated with oncogene and proto-oncogene mediated growth induction, tumor suppressor mediated growth arrest and replicative senescence, apoptosis, and DNA repair. The NuA4 complex ATPase and helicase activities seem to be, at least in part, contributed by the association of RUVBL1 and RUVBL2 with EP400. NuA4 may also play a direct role in DNA repair when recruited to sites of DNA damage. RUVBL1 plays an essential role in oncogenic transformation by MYC and also modulates transcriptional activation by the LEF1/TCF1 -CTNNB1 complex. High levels of autoantibodies against RUVBL1 are detected in sera of patients with autoimmune diseases such as polymyositis/dermatomyositis and autoimmune hepatitis.

RUVBL1 (TIP49) Antibody (Center) - References

- Feng, Y., et al., Cancer Res. 63(24):8726-8734 (2003).
Hawley, S.B., et al., J. Biol. Chem. 276(1):179-186 (2001).
Ikura, T., et al., Cell 102(4):463-473 (2000).
Salzer, U., et al., Biochim. Biophys. Acta 1446(3):365-370 (1999).
Makino, Y., et al., Biochem. Biophys. Res. Commun. 245(3):819-823 (1998).