

**Anti-RNA Helicase A Rabbit Monoclonal Antibody**  
**Catalog # ABO14645****Specification****Anti-RNA Helicase A Rabbit Monoclonal Antibody - Product Information**

Application	WB, IHC, IF, ICC, FC
Primary Accession	<a href="#">Q08211</a>
Host	Rabbit
Isotype	Rabbit IgG
Reactivity	Rat, Human, Mouse
Clonality	Monoclonal
Format	Liquid

**Description**

Anti-RNA Helicase A Rabbit Monoclonal Antibody . Tested in WB, IHC, ICC/IF, Flow Cytometry applications. This antibody reacts with Human, Mouse, Rat.

**Anti-RNA Helicase A Rabbit Monoclonal Antibody - Additional Information**

**Gene ID** 1660

**Other Names**

ATP-dependent RNA helicase A, 3.6.4.13, DEAH box protein 9, DExH-box helicase 9 {ECO:0000312|HGNC:HGNC:2750}, Leukophysin, LKP, Nuclear DNA helicase II, NDH II, RNA helicase A, DHX9 ([http://www.genenames.org/cgi-bin/gene\\_symbol\\_report?hgnc\\_id=2750](http://www.genenames.org/cgi-bin/gene_symbol_report?hgnc_id=2750))

**Application Details**

WB 1:1000-1:5000<br>IHC 1:50-1:200<br>ICC/IF 1:50-1:200<br>FC 1:20

**Contents**

Rabbit IgG in phosphate buffered saline, pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol, 0.4-0.5mg/ml BSA.

**Immunogen**

A synthesized peptide derived from human RNA Helicase A

**Purification**

Affinity-chromatography

**Storage**

**Store at -20°C for one year. For short term storage and frequent use, store at 4°C for up to one month. Avoid repeated freeze-thaw cycles.**

**Anti-RNA Helicase A Rabbit Monoclonal Antibody - Protein Information**

**Name** DHX9 ([HGNC:2750](#))

## Function

Multifunctional ATP-dependent nucleic acid helicase that unwinds DNA and RNA in a 3' to 5' direction and that plays important roles in many processes, such as DNA replication, transcriptional activation, post-transcriptional RNA regulation, mRNA translation and RNA-mediated gene silencing (PubMed:<a href="http://www.uniprot.org/citations/11416126" target="\_blank">11416126</a>, PubMed:<a href="http://www.uniprot.org/citations/12711669" target="\_blank">12711669</a>, PubMed:<a href="http://www.uniprot.org/citations/15355351" target="\_blank">15355351</a>, PubMed:<a href="http://www.uniprot.org/citations/16680162" target="\_blank">16680162</a>, PubMed:<a href="http://www.uniprot.org/citations/17531811" target="\_blank">17531811</a>, PubMed:<a href="http://www.uniprot.org/citations/20669935" target="\_blank">20669935</a>, PubMed:<a href="http://www.uniprot.org/citations/21561811" target="\_blank">21561811</a>, PubMed:<a href="http://www.uniprot.org/citations/24049074" target="\_blank">24049074</a>, PubMed:<a href="http://www.uniprot.org/citations/24990949" target="\_blank">24990949</a>, PubMed:<a href="http://www.uniprot.org/citations/25062910" target="\_blank">25062910</a>, PubMed:<a href="http://www.uniprot.org/citations/28221134" target="\_blank">28221134</a>, PubMed:<a href="http://www.uniprot.org/citations/9111062" target="\_blank">9111062</a>, PubMed:<a href="http://www.uniprot.org/citations/37467750" target="\_blank">37467750</a>). Requires a 3'- single-stranded tail as entry site for acid nuclei unwinding activities as well as the binding and hydrolyzing of any of the four ribo- or deoxyribo-nucleotide triphosphates (NTPs) (PubMed:<a href="http://www.uniprot.org/citations/1537828" target="\_blank">1537828</a>). Unwinds numerous nucleic acid substrates such as double-stranded (ds) DNA and RNA, DNA:RNA hybrids, DNA and RNA forks composed of either partially complementary DNA duplexes or DNA:RNA hybrids, respectively, and also DNA and RNA displacement loops (D- and R-loops), triplex-helical DNA (H-DNA) structure and DNA and RNA-based G-quadruplexes (PubMed:<a href="http://www.uniprot.org/citations/20669935" target="\_blank">20669935</a>, PubMed:<a href="http://www.uniprot.org/citations/21561811" target="\_blank">21561811</a>, PubMed:<a href="http://www.uniprot.org/citations/24049074" target="\_blank">24049074</a>). Binds dsDNA, single-stranded DNA (ssDNA), dsRNA, ssRNA and poly(A)-containing RNA (PubMed:<a href="http://www.uniprot.org/citations/10198287" target="\_blank">10198287</a>, PubMed:<a href="http://www.uniprot.org/citations/9111062" target="\_blank">9111062</a>). Also binds to circular dsDNA or dsRNA of either linear and/or circular forms and stimulates the relaxation of supercoiled DNAs catalyzed by topoisomerase TOP2A (PubMed:<a href="http://www.uniprot.org/citations/12711669" target="\_blank">12711669</a>). Plays a role in DNA replication at origins of replication and cell cycle progression (PubMed:<a href="http://www.uniprot.org/citations/24990949" target="\_blank">24990949</a>). Plays a role as a transcriptional coactivator acting as a bridging factor between polymerase II holoenzyme and transcription factors or cofactors, such as BRCA1, CREBBP, RELA and SMN1 (PubMed:<a href="http://www.uniprot.org/citations/11038348" target="\_blank">11038348</a>, PubMed:<a href="http://www.uniprot.org/citations/11149922" target="\_blank">11149922</a>, PubMed:<a href="http://www.uniprot.org/citations/11416126" target="\_blank">11416126</a>, PubMed:<a href="http://www.uniprot.org/citations/15355351" target="\_blank">15355351</a>, PubMed:<a href="http://www.uniprot.org/citations/28221134" target="\_blank">28221134</a>, PubMed:<a href="http://www.uniprot.org/citations/9323138" target="\_blank">9323138</a>, PubMed:<a href="http://www.uniprot.org/citations/9662397" target="\_blank">9662397</a>). Binds to the CDKN2A promoter (PubMed:<a href="http://www.uniprot.org/citations/11038348" target="\_blank">11038348</a>). Plays several roles in post-transcriptional regulation of gene expression (PubMed:<a href="http://www.uniprot.org/citations/28221134" target="\_blank">28221134</a>, PubMed:<a href="http://www.uniprot.org/citations/28355180" target="\_blank">28355180</a>). In cooperation with NUP98, promotes pre-mRNA alternative splicing activities of a subset of genes (PubMed:<a href="http://www.uniprot.org/citations/11402034" target="\_blank">11402034</a>, PubMed:<a href="http://www.uniprot.org/citations/16680162" target="\_blank">16680162</a>, PubMed:<a href="http://www.uniprot.org/citations/28221134" target="\_blank">28221134</a>, PubMed:<a href="http://www.uniprot.org/citations/28355180" target="\_blank">28355180</a>). As component of a large PER complex, is involved in the negative regulation of 3' transcriptional termination of circadian target genes such as PER1 and NR1D1 and the control of the circadian

rhythms (By similarity). Also acts as a nuclear resolvase that is able to bind and neutralize harmful massive secondary double-stranded RNA structures formed by inverted-repeat Alu retrotransposon elements that are inserted and transcribed as parts of genes during the process of gene transposition (PubMed:<a href="http://www.uniprot.org/citations/28355180" target="\_blank">28355180</a>). Involved in the positive regulation of nuclear export of constitutive transport element (CTE)-containing unspliced mRNA (PubMed:<a href="http://www.uniprot.org/citations/10924507" target="\_blank">10924507</a>, PubMed:<a href="http://www.uniprot.org/citations/11402034" target="\_blank">11402034</a>, PubMed:<a href="http://www.uniprot.org/citations/9162007" target="\_blank">9162007</a>). Component of the coding region determinant (CRD)-mediated complex that promotes cytoplasmic MYC mRNA stability (PubMed:<a href="http://www.uniprot.org/citations/19029303" target="\_blank">19029303</a>). Plays a role in mRNA translation (PubMed:<a href="http://www.uniprot.org/citations/28355180" target="\_blank">28355180</a>). Positively regulates translation of selected mRNAs through its binding to post-transcriptional control element (PCE) in the 5'-untranslated region (UTR) (PubMed:<a href="http://www.uniprot.org/citations/16680162" target="\_blank">16680162</a>). Involved with LARP6 in the translation stimulation of type I collagen mRNAs for COL1A1 and COL1A2 through binding of a specific stem-loop structure in their 5'-UTRs (PubMed:<a href="http://www.uniprot.org/citations/22190748" target="\_blank">22190748</a>). Stimulates LIN28A- dependent mRNA translation probably by facilitating ribonucleoprotein remodeling during the process of translation (PubMed:<a href="http://www.uniprot.org/citations/21247876" target="\_blank">21247876</a>). Plays also a role as a small interfering (siRNA)-loading factor involved in the RNA-induced silencing complex (RISC) loading complex (RLC) assembly, and hence functions in the RISC-mediated gene silencing process (PubMed:<a href="http://www.uniprot.org/citations/17531811" target="\_blank">17531811</a>). Binds preferentially to short double- stranded RNA, such as those produced during rotavirus intestinal infection (PubMed:<a href="http://www.uniprot.org/citations/28636595" target="\_blank">28636595</a>). This interaction may mediate NLRP9 inflammasome activation and trigger inflammatory response, including IL18 release and pyroptosis (PubMed:<a href="http://www.uniprot.org/citations/28636595" target="\_blank">28636595</a>). Finally, mediates the attachment of heterogeneous nuclear ribonucleoproteins (hnRNPs) to actin filaments in the nucleus (PubMed:<a href="http://www.uniprot.org/citations/11687588" target="\_blank">11687588</a>).

### Cellular Location

Nucleus. Nucleus, nucleoplasm. Nucleus, nucleolus. Cytoplasm. Cytoplasm, cytoskeleton, microtubule organizing center, centrosome. Note=Nucleoplasmic shuttling protein (PubMed:10198287, PubMed:10207077, PubMed:16375861, PubMed:9162007) Its nuclear import involves the nucleocytoplasmic transport receptor Importin alpha/Importin beta receptor pathway in a Ran-dependent manner (PubMed:16375861). In interphase, localizes in nuclear stress granules and at perichromatin fibrils and in cytoplasmic ribonucleoprotein granules (PubMed:10198287). Colocalizes with WRN and H2AX at centrosomes in a microtubule-dependent manner following DNA damaging agent treatment (PubMed:17498979). Excluded from the mitotic nucleus as early as prophase and re-entered the nucleus at telophase (PubMed:10198287). Recruited in diffuse and discrete intranuclear foci (GLFG-body) in a NUP98-dependent manner (PubMed:28221134). Colocalizes with SP7 in the nucleus (PubMed:17303075). Colocalizes with ACTB at nuclear actin filaments inside the nucleus or at the nuclear pore (PubMed:11687588). Colocalizes with HNRNPC at nuclear ribonucleoprotein complex proteins in the nucleus (PubMed:11687588). Localized in cytoplasmic mRNP granules containing untranslated mRNAs (PubMed:17289661).

### Anti-RNA Helicase A Rabbit Monoclonal Antibody - 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](#)

## Anti-RNA Helicase A Rabbit Monoclonal Antibody - Images

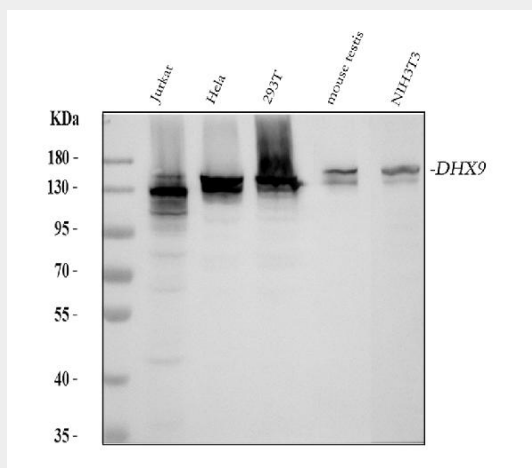


Figure 1. Western blot analysis of RNA Helicase A using anti-RNA Helicase A antibody (M02550). Electrophoresis was performed on a 5-20% SDS-PAGE gel at 70V (Stacking gel) / 90V (Resolving gel) for 2-3 hours. The sample well of each lane was loaded with 30 ug of sample under reducing conditions.

Lane 1: human Jurkat whole cell lysates,

Lane 2: human Hela whole cell lysates,

Lane 3: human 293T whole cell lysates,

Lane 4: mouse testis tissue lysates,

Lane 5: mouse NIH/3T3 whole cell lysates.

After electrophoresis, proteins were transferred to a nitrocellulose membrane at 150 mA for 50-90 minutes. Blocked the membrane with 5% non-fat milk/TBS for 1.5 hour at RT. The membrane was incubated with rabbit anti-RNA Helicase A antigen affinity purified monoclonal antibody (Catalog # M02550) at 1:1000 overnight at 4°C, then washed with TBS-0.1%Tween 3 times with 5 minutes each and probed with a goat anti-rabbit IgG-HRP secondary antibody at a dilution of 1:5000 for 1.5 hour at RT. The signal is developed using an Enhanced Chemiluminescent detection (ECL) kit (Catalog # EK1002) with Tanon 5200 system. A specific band was detected for RNA Helicase A at approximately 141 kDa. The expected band size for RNA Helicase A is at 141 kDa.