

**POLR1B Antibody (N-term)**  
**Affinity Purified Rabbit Polyclonal Antibody (Pab)**  
**Catalog # AP12495a**

**Specification**

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**POLR1B Antibody (N-term) - Product Information**

Application	WB, IHC-P, FC,E
Primary Accession	<a href="#">O9H9Y6</a>
Other Accession	<a href="#">O54888</a> , <a href="#">NP_061887.2</a> , <a href="#">NP_001131076.1</a>
Reactivity	Human
Predicted	Rat
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	128229
Antigen Region	89-117

**POLR1B Antibody (N-term) - Additional Information**

**Gene ID** 84172

**Other Names**

DNA-directed RNA polymerase I subunit RPA2, RNA polymerase I subunit 2, DNA-directed RNA polymerase I 135 kDa polypeptide, RPA135, POLR1B

**Target/Specificity**

This POLR1B antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 89-117 amino acids from the N-terminal region of human POLR1B.

**Dilution**

WB~~1:1000  
IHC-P~~1:10~50  
FC~~1:10~50

**Format**

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is purified through a protein A column, followed by peptide affinity purification.

**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**

POLR1B Antibody (N-term) is for research use only and not for use in diagnostic or therapeutic procedures.

**POLR1B Antibody (N-term) - Protein Information**

**Name** POLR1B {ECO:0000303|PubMed:31649276, ECO:0000312|HGNC:HGNC:20454}

**Function** Catalytic core component of RNA polymerase I (Pol I), a DNA- dependent RNA polymerase which synthesizes ribosomal RNA precursors using the four ribonucleoside triphosphates as substrates. Transcribes 47S pre-rRNAs from multicopy rRNA gene clusters, giving rise to 5.8S, 18S and 28S ribosomal RNAs (PubMed:[11250903](#), PubMed:[11283244](#), PubMed:[16858408](#), PubMed:[34671025](#), PubMed:[34887565](#), PubMed:[36271492](#)). Pol I-mediated transcription cycle proceeds through transcription initiation, transcription elongation and transcription termination stages. During transcription initiation, Pol I pre-initiation complex (PIC) is recruited by the selectivity factor 1 (SL1/TIF-IB) complex bound to the core promoter that precedes an rDNA repeat unit. The PIC assembly bends the promoter favoring the formation of the transcription bubble and promoter escape. Once the polymerase has escaped from the promoter it enters the elongation phase during which RNA is actively polymerized, based on complementarity with the template DNA strand. Highly processive, assembles in structures referred to as 'Miller trees' where many elongating Pol I complexes queue and transcribe the same rDNA coding regions. At terminator sequences downstream of the rDNA gene, PTRF interacts with Pol I and halts Pol I transcription leading to the release of the RNA transcript and polymerase from the DNA (PubMed:[11250903](#), PubMed:[11283244](#), PubMed:[16858408](#), PubMed:[34671025](#), PubMed:[34887565](#), PubMed:[36271492](#)). Forms Pol I active center together with the largest subunit POLR1A/RPA1. Appends one nucleotide at a time to the 3' end of the nascent RNA, with POLR1A/RPA1 contributing a Mg(2+)-coordinating DxDGD motif, and POLR1B/RPA2 participating in the coordination of a second Mg(2+) ion and providing lysine residues believed to facilitate Watson-Crick base pairing between the incoming nucleotide and the template base. Typically, Mg(2+) ions direct a 5' nucleoside triphosphate to form a phosphodiester bond with the 3' hydroxyl of the preceding nucleotide of the nascent RNA, with the elimination of pyrophosphate. Has proofreading activity: Pauses and backtracks to allow the cleavage of a missincorporated nucleotide via POLR1H/RPA12. High Pol I processivity is associated with decreased transcription fidelity (By similarity) (PubMed:[11250903](#), PubMed:[11283244](#), PubMed:[16809778](#), PubMed:[16858408](#), PubMed:[34671025](#), PubMed:[34887565](#), PubMed:[36271492](#)).

#### **Cellular Location**

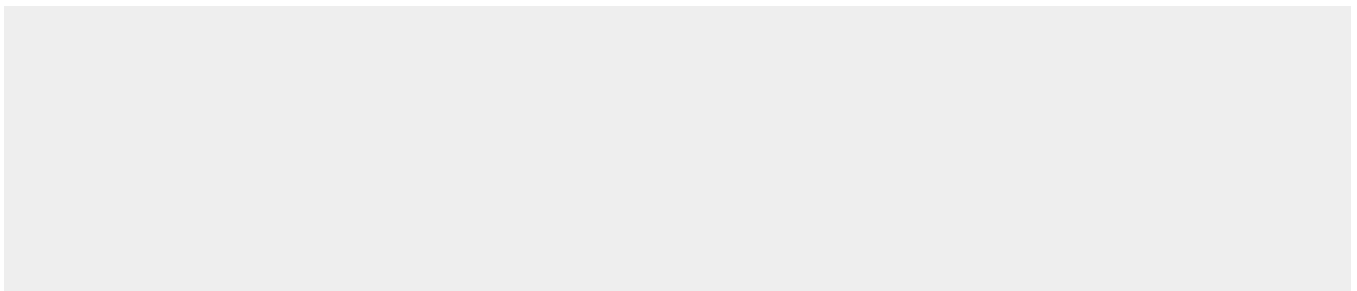
Nucleus, nucleolus. Chromosome {ECO:0000250|UniProtKB:P70700}

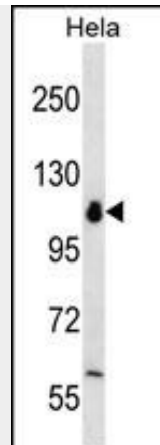
#### **POLR1B Antibody (N-term) - Protocols**

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

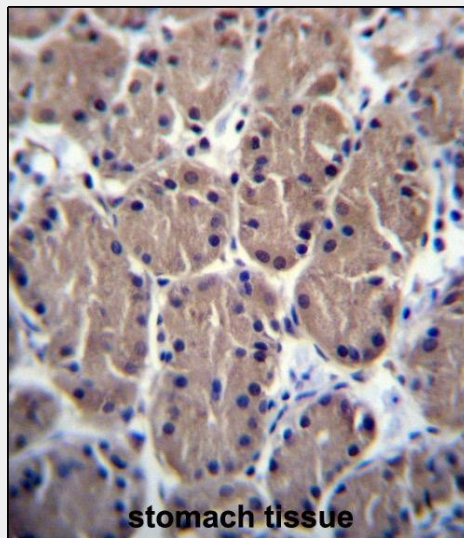
- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

#### **POLR1B Antibody (N-term) - Images**

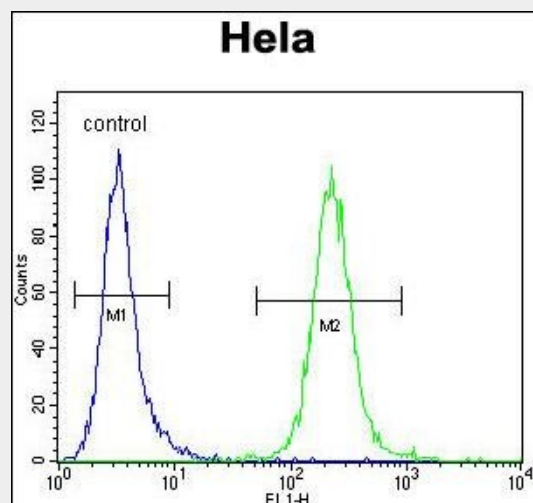




POLR1B Antibody (N-term) (Cat. #AP12495a) western blot analysis in HeLa cell line lysates (35ug/lane). This demonstrates the POLR1B antibody detected the POLR1B protein (arrow).



POLR1B Antibody (N-term) (Cat. #AP12495a) immunohistochemistry analysis in formalin fixed and paraffin embedded human stomach tissue followed by peroxidase conjugation of the secondary antibody and DAB staining. This data demonstrates the use of POLR1B Antibody (N-term) for immunohistochemistry. Clinical relevance has not been evaluated.



POLR1B Antibody (N-term) (Cat. #AP12495a) flow cytometric analysis of HeLa cells (right

histogram) compared to a negative control cell (left histogram).FITC-conjugated goat-anti-rabbit secondary antibodies were used for the analysis.

#### **POLR1B Antibody (N-term) - Background**

Eukaryotic RNA polymerase I (pol I) is responsible for the transcription of ribosomal RNA (rRNA) genes and production of rRNA, the primary component of ribosomes. Pol I is a multisubunit enzyme composed of 6 to 14 polypeptides, depending on the species. Most of the mass of the pol I complex derives from the 2 largest subunits, Rpa1 and Rpa2 in yeast. POLR1B is homologous to Rpa2 (Seither and Grummt, 1996 [PubMed 8921381]).

#### **POLR1B Antibody (N-term) - References**

Wen, L., et al. Biochem. Biophys. Res. Commun. 367(4):846-851(2008)  
Wu, C., et al. Proteomics 7(11):1775-1785(2007)  
Johnson, S.S., et al. Mol. Cell 26(3):367-379(2007)  
Panova, T.B., et al. Mol. Cell. Biol. 26(16):5957-5968(2006)  
Percipalle, P., et al. EMBO Rep. 7(5):525-530(2006)