Datasheet for ABIN2783467
anti-POLR2H antibody (N-Term)

## 1 Image



Overview

| Quantity: | $100 \mu \mathrm{~L}$ |
| :--- | :--- |
| Target: | POLR2H |
| Binding Specificity: | N-Term |
| Reactivity: | Human, Mouse, Rat, Cow, Guinea Pig, Horse, Rabbit, Zebrafish (Danio rerio) |
| Host: | Polyclonal |
| Clonality: | This POLR2H antibody is un-conjugated |
| Conjugate: | Western Blotting (WB) |

Product Details

| Immunogen: | The immunogen is a synthetic peptide directed towards the $N$ terminal region of human POLR2H |
| :---: | :---: |
| Sequence: | DLGDKFRLVI ASTLYEDGTL DDGEYNPTDD RPSRADQFEY VMYGKVYRIE |
| Predicted Reactivity: | Cow: $100 \%$, Guinea Pig: $100 \%$, Horse: $100 \%$, Human: $100 \%$, Mouse: $100 \%$, Rabbit: $100 \%$, Rat: 100\%, Zebrafish: 93\% |
| Characteristics: | This is a rabbit polyclonal antibody against POLR2H. It was validated on Western Blot using a cell lysate as a positive control. |
| Purification: | Protein A purified |

Target Details
Target:
POLR2H

| Alternative Name: | POLR2H (POLR2H Products) |
| :---: | :---: |
| Background: | POLR2H is one of the essential subunits of RNA polymerase II that is shared by the other two eukaryotic DNA-directed RNA polymerases, I and III.This gene encodes one of the essential subunits of RNA polymerase II that is shared by the other two eukaryotic DNA-directed RNA polymerases, I and III. Publication Note: This RefSeq record includes a subset of the publications that are available for this gene. Please see the Entrez Gene record to access additional publications. This gene encodes a member of the E2F transcription factor protein family. E2F family members play a crucial role in control of the cell cycle and of the action of tumor suppressor proteins. They are also a target of the transforming proteins of small DNA tumor viruses. Many E2F proteins contain several evolutionarily conserved domains: a DNA binding domain, a dimerization domain which determines interaction with the differentiation regulated transcription factor proteins (DP), a transactivation domain enriched in acidic amino acids, and a tumor suppressor protein association domain which is embedded within the transactivation domain. The encoded protein of this gene is atypical because it lacks the transactivation and tumor suppressor protein association domains. It contains a modular suppression domain and is an inhibitor of E2F-dependent transcription. The protein is part of a multimeric protein complex that contains a histone methyltransferase and the transcription factors Mga and Max. Multiple transcript variants have been reported for this gene, but it has not been clearly demonstrated that they encode valid isoforms. Publication Note: This RefSeq record includes a subset of the publications that are available for this gene. Please see the Entrez Gene record to access additional publications. PRIMARYREFSEQ_SPAN PRIMARY_IDENTIFIER PRIMARY_SPAN COMP 1-400 AU142999.1 1-400 401-907 BI772069.1 287-793 908-1792 BC008348.1 928-1812 1793-3185 AC099344.4 111461-112853 c Alias Symbols: RPABC3, RPB17, RPB8, hsRPB8 Protein Interaction Partner: TCEB3, SRC, PSMB9, ESR1, CEP57, RNF2, SNRNP200, SART3, POLR2E, DNMT3B, DNMT3A, BAG3, POLR2C, MED19, METTL18, MED26, UBC, POLR3D, POLR3H, POLR3B, POLR1A, POLR3A, POLR1C, NEDD8, Atf7ip, SUMO2, INTS10, POLR2D, POLR2A, BRCA1, BARD1, INTS5, INTS3, INTS6, INTS1 <br> Protein Size: 150 |
| Molecular Weight: | 17 kDa |
| Gene ID: | 5437 |

NCBI Accession: NM_006232, NP_006223

UniProt:
Pathways:

P52434

Regulatory RNA Pathways

## Application Details

| Application Notes: | Optimal working dilutions should be determined experimentally by the investigator. |
| :--- | :--- |
| Comment: | Antigen size: 150 AA |
| Restrictions: | For Research Use only |
| Handling | Liquid |
| Format: | Lot specific |
| Concentration: | Sucrose. |
| Buffer: | This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which |
| Preservative: | should be handled by trained staff only. |
| Precaution of Use: | Avoid repeated freeze-thaw cycles. |
| Storag in $1 \times$ PBS buffer with 0.09 \% (w/v) sodium azide and $2 \%$ |  |
| Storage Comment: | -20 ${ }^{\circ} \mathrm{C}$ |

Publications

Product cited in:
Hamada, Tashiro, Tada, Inazawa, Shirozu, Shibahara, Nakamura, Martina, Nakano, Honjo: " Isolation and characterization of a novel secretory protein, stromal cell-derived factor-2 (SDF-2) using the signal sequence trap method." in: Gene, Vol. 176, Issue 1-2, pp. 211-4, (1997) ( PubMed).


