Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 26, 2025

Rpb1 NTD (D8L4Y)

RRID:AB_2687876 Type: Antibody

Proper Citation

(Cell Signaling Technology Cat# 14958, RRID:AB_2687876)

Antibody Information

URL: http://antibodyregistry.org/AB_2687876

Proper Citation: (Cell Signaling Technology Cat# 14958, RRID:AB_2687876)

Target Antigen: RPB1 (N-terminal domain)

Host Organism: rabbit

Clonality: monoclonal

Comments: Applications: W, ChIP, ChIP-seq

Antibody Name: Rpb1 NTD (D8L4Y)

Description: This monoclonal targets RPB1 (N-terminal domain)

Target Organism: Human, Rat, Monkey, Mouse

Clone ID: D8L4Y

Antibody ID: AB_2687876

Vendor: Cell Signaling Technology

Catalog Number: 14958

Alternative Catalog Numbers: 14958S

Record Creation Time: 20231110T034040+0000

Record Last Update: 20240725T053921+0000

Ratings and Alerts

No rating or validation information has been found for Rpb1 NTD (D8L4Y).

No alerts have been found for Rpb1 NTD (D8L4Y).

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 46 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Aboreden NG, et al. (2025) LDB1 establishes multi-enhancer networks to regulate gene expression. Molecular cell, 85(2), 376.

Boddu PC, et al. (2024) Transcription elongation defects link oncogenic SF3B1 mutations to targetable alterations in chromatin landscape. Molecular cell, 84(8), 1475.

Bruno F, et al. (2024) Collisions of RNA polymerases behind the replication fork promote alternative RNA splicing in newly replicated chromatin. Molecular cell, 84(2), 221.

Žumer K, et al. (2024) FACT maintains chromatin architecture and thereby stimulates RNA polymerase II pausing during transcription in vivo. Molecular cell, 84(11), 2053.

Liang F, et al. (2024) DOT1L/H3K79me2 represses HIV-1 reactivation via recruiting DCAF1. Cell reports, 43(7), 114368.

Zhao H, et al. (2024) Pluripotency state transition of embryonic stem cells requires the turnover of histone chaperone FACT on chromatin. iScience, 27(1), 108537.

Blears D, et al. (2024) Redundant pathways for removal of defective RNA polymerase II complexes at a promoter-proximal pause checkpoint. Molecular cell, 84(24), 4790.

Hong Y, et al. (2024) SAFB restricts contact domain boundaries associated with L1 chimeric transcription. Molecular cell, 84(9), 1637.

Han X, et al. (2024) Nuclear RNA homeostasis promotes systems-level coordination of cell fate and senescence. Cell stem cell, 31(5), 694.

Phongbunchoo Y, et al. (2024) YY1-mediated enhancer-promoter communication in the immunoglobulin? locus is regulated by MSL/MOF recruitment. Cell reports, 43(7), 114456.

Lukoszek R, et al. (2024) CK2 phosphorylation of CMTR1 promotes RNA cap formation and

influenza virus infection. Cell reports, 43(7), 114405.

van den Heuvel D, et al. (2024) STK19 facilitates the clearance of lesion-stalled RNAPII during transcription-coupled DNA repair. Cell, 187(25), 7107.

Wang Z, et al. (2024) The phosphatase PP1 sustains global transcription by promoting RNA polymerase II pause release. Molecular cell, 84(24), 4824.

Li Y, et al. (2023) RNA Pol II preferentially regulates ribosomal protein expression by trapping disassociated subunits. Molecular cell, 83(8), 1280.

Ji YX, et al. (2023) Protocol for rapidly inducing genome-wide RNA Pol II hyperphosphorylation by selectively disrupting INTAC phosphatase activity. STAR protocols, 4(4), 102640.

Sun Z, et al. (2023) Chromatin regulation of transcriptional enhancers and cell fate by the Sotos syndrome gene NSD1. Molecular cell, 83(14), 2398.

Hu S, et al. (2023) INTAC endonuclease and phosphatase modules differentially regulate transcription by RNA polymerase II. Molecular cell, 83(10), 1588.

Yang N, et al. (2023) A hyper-quiescent chromatin state formed during aging is reversed by regeneration. Molecular cell, 83(10), 1659.

Marasco LE, et al. (2022) Counteracting chromatin effects of a splicing-correcting antisense oligonucleotide improves its therapeutic efficacy in spinal muscular atrophy. Cell, 185(12), 2057.

Wang X, et al. (2022) RPAP2 regulates a transcription initiation checkpoint by inhibiting assembly of pre-initiation complex. Cell reports, 39(4), 110732.