

Resource Summary Report

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Anti-RNA polymerase II subunit B1 (phospho-CTD Ser-5), clone 3E8

RRID:AB_10615822

Type: Antibody

Proper Citation

(Millipore Cat# 04-1572, RRID:AB_10615822)

Antibody Information

URL: http://antibodyregistry.org/AB_10615822

Proper Citation: (Millipore Cat# 04-1572, RRID:AB_10615822)

Target Antigen: RNA polymerase II subunit B1 (phospho-CTD Ser-5) clone 3E8

Clonality: monoclonal

Comments: seller recommendations: IgG2a; IgG2a Western Blot; Immunocytochemistry; ChIP; ELISA; WB, ELISA, ChIP

Antibody Name: Anti-RNA polymerase II subunit B1 (phospho-CTD Ser-5), clone 3E8

Description: This monoclonal targets RNA polymerase II subunit B1 (phospho-CTD Ser-5) clone 3E8

Target Organism: h, m

Antibody ID: AB_10615822

Vendor: Millipore

Catalog Number: 04-1572

Record Creation Time: 20231110T071140+0000

Record Last Update: 20241115T052021+0000

Ratings and Alerts

No rating or validation information has been found for Anti-RNA polymerase II subunit B1 (phospho-CTD Ser-5), clone 3E8.

No alerts have been found for Anti-RNA polymerase II subunit B1 (phospho-CTD Ser-5), clone 3E8.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 18 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Velychko T, et al. (2024) CDK7 kinase activity promotes RNA polymerase II promoter escape by facilitating initiation factor release. *Molecular cell*, 84(12), 2287.

Moreno RY, et al. (2023) Distinctive interactomes of RNA polymerase II phosphorylation during different stages of transcription. *iScience*, 26(9), 107581.

Pappas G, et al. (2023) MDC1 maintains active elongation complexes of RNA polymerase II. *Cell reports*, 42(1), 111979.

Perurena N, et al. (2023) USP9X mediates an acute adaptive response to MAPK suppression in pancreatic cancer but creates multiple actionable therapeutic vulnerabilities. *Cell reports. Medicine*, 4(4), 101007.

Lyons H, et al. (2023) Functional partitioning of transcriptional regulators by patterned charge blocks. *Cell*, 186(2), 327.

Berry S, et al. (2022) Feedback from nuclear RNA on transcription promotes robust RNA concentration homeostasis in human cells. *Cell systems*, 13(6), 454.

Allen BL, et al. (2022) Suppression of p53 response by targeting p53-Mediator binding with a stapled peptide. *Cell reports*, 39(1), 110630.

Aoi Y, et al. (2021) SPT5 stabilization of promoter-proximal RNA polymerase II. *Molecular cell*, 81(21), 4413.

Badjatia N, et al. (2021) Acute stress drives global repression through two independent RNA polymerase II stalling events in *Saccharomyces*. *Cell reports*, 34(3), 108640.

Richters A, et al. (2021) Modulating Androgen Receptor-Driven Transcription in Prostate Cancer with Selective CDK9 Inhibitors. *Cell chemical biology*, 28(2), 134.

Heo DH, et al. (2021) Transcription and chromatin-based surveillance mechanism controls suppression of cryptic antisense transcription. *Cell reports*, 36(10), 109671.

Nakazawa Y, et al. (2020) Ubiquitination of DNA Damage-Stalled RNAPII Promotes Transcription-Coupled Repair. *Cell*, 180(6), 1228.

Aoi Y, et al. (2020) NELF Regulates a Promoter-Proximal Step Distinct from RNA Pol II Pause-Release. *Molecular cell*, 78(2), 261.

Zhang H, et al. (2020) CDK7 Inhibition Potentiates Genome Instability Triggering Anti-tumor Immunity in Small Cell Lung Cancer. *Cancer cell*, 37(1), 37.

Olson CM, et al. (2019) Development of a Selective CDK7 Covalent Inhibitor Reveals Predominant Cell-Cycle Phenotype. *Cell chemical biology*, 26(6), 792.

Mayfield JE, et al. (2019) Tyr1 phosphorylation promotes phosphorylation of Ser2 on the C-terminal domain of eukaryotic RNA polymerase II by P-TEFb. *eLife*, 8.

Kecman T, et al. (2018) Elongation/Termination Factor Exchange Mediated by PP1 Phosphatase Orchestrates Transcription Termination. *Cell reports*, 25(1), 259.

Fischl H, et al. (2017) Paf1 Has Distinct Roles in Transcription Elongation and Differential Transcript Fate. *Molecular cell*, 65(4), 685.