

Resource Summary Report

Generated by ASWG on Aug 9, 2026

Cdk9 (D-7)

RRID:AB_627245

Type: Antibody

Proper Citation

Santa Cruz Biotechnology Cat# sc-13130, RRID:AB_627245

Antibody Information

URL: http://antibodyregistry.org/AB_627245

Proper Citation: Santa Cruz Biotechnology Cat# sc-13130, RRID:AB_627245

Target Antigen: CDK9

Host Organism: mouse

Clonality: monoclonal

Comments: validation status unknown check with seller; recommendations: ELISA; Immunocytochemistry; Immunofluorescence; Immunohistochemistry; Immunoprecipitation; Western Blot; Western Blotting, Immunoprecipitation, Immunofluorescence, Immunohistochemistry(P), ELISA

Antibody Name: Cdk9 (D-7)

Description: This monoclonal targets CDK9

Target Organism: rat, mouse, human

Clone ID: D-7

Antibody ID: AB_627245

Vendor: Santa Cruz Biotechnology

Catalog Number: sc-13130

Record Creation Time: 20250820T060910+0000

Record Last Update: 20250820T232440+0000

Ratings and Alerts

No rating or validation information has been found for Cdk9 (D-7).

No alerts have been found for Cdk9 (D-7).

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 15 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Tang J, et al. (2025) Targeting N-Myc in neuroblastoma with selective Aurora kinase A degraders. *Cell chemical biology*, 32(2), 352.

Puleo N, et al. (2025) Identification of a TNIK-CDK9 Axis as a Targetable Strategy for Platinum-Resistant Ovarian Cancer. *Molecular cancer therapeutics*, 24(4), 639.

Silva TF, et al. (2025) Extracellular vesicle heterogeneity through the lens of multiomics. *Cell reports. Medicine*, 6(7), 102161.

Cenik BK, et al. (2024) TurboCas: A method for locus-specific labeling of genomic regions and isolating their associated protein interactome. *Molecular cell*, 84(24), 4929.

Shan X, et al. (2023) Identification of a diketopiperazine-based O-GlcNAc transferase inhibitor sensitizing hepatocellular carcinoma to CDK9 inhibition. *The FEBS journal*.

Pei J, et al. (2023) Piperlongumine conjugates induce targeted protein degradation. *Cell chemical biology*, 30(2), 203.

Qiu Z, et al. (2022) Transcription Elongation Machinery Is a Druggable Dependency and Potentiates Immunotherapy in Glioblastoma Stem Cells. *Cancer discovery*, 12(2), 502.

Olsen SN, et al. (2022) MLL::AF9 degradation induces rapid changes in transcriptional elongation and subsequent loss of an active chromatin landscape. *Molecular cell*, 82(6), 1140.

Gonzalez-Teran B, et al. (2022) Transcription factor protein interactomes reveal genetic determinants in heart disease. *Cell*, 185(5), 794.

Ohno SI, et al. (2022) Nuclear microRNAs release paused Pol II via the DDX21-CDK9 complex. *Cell reports*, 39(2), 110673.

Wang X, et al. (2021) DDB1 binds histone reader BRWD3 to activate the transcriptional cascade in adipogenesis and promote onset of obesity. *Cell reports*, 35(12), 109281.

Studniarek C, et al. (2021) The 7SK/P-TEFb snRNP controls ultraviolet radiation-induced transcriptional reprogramming. *Cell reports*, 35(2), 108965.

Bacon CW, et al. (2020) KAP1 Is a Chromatin Reader that Couples Steps of RNA Polymerase II Transcription to Sustain Oncogenic Programs. *Molecular cell*, 78(6), 1133.

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

Faust TB, et al. (2018) The HIV-1 Tat protein recruits a ubiquitin ligase to reorganize the 7SK snRNP for transcriptional activation. *eLife*, 7.