

# Resource Summary Report

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## Rabbit Anti-Histone H3, trimethyl XP??? Monoclonal Antibody, Unconjugated, Clone D5A7

RRID:AB\_1950412

Type: Antibody

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### Proper Citation

(Cell Signaling Technology Cat# 4909, RRID:AB\_1950412)

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_1950412](http://antibodyregistry.org/AB_1950412)

**Proper Citation:** (Cell Signaling Technology Cat# 4909, RRID:AB\_1950412)

**Target Antigen:** Histone H3, trimethyl

**Host Organism:** rabbit

**Clonality:** monoclonal

**Comments:** Applications: W, IHC-P, IF-IC, F, ChIP, ChIP-seq. Consolidation on 11/2018: AB\_10499634, AB\_10500802, AB\_1950412, AB\_1950414, AB\_2616016.

**Antibody Name:** Rabbit Anti-Histone H3, trimethyl XP??? Monoclonal Antibody, Unconjugated, Clone D5A7

**Description:** This monoclonal targets Histone H3, trimethyl

**Target Organism:** chicken, monkey, chickenavian, rat, hamster, simian, xenopus, mouse, drosophila, fish, bovine, zebrafish, human

**Clone ID:** Clone D5A7

**Antibody ID:** AB\_1950412

**Vendor:** Cell Signaling Technology

**Catalog Number:** 4909

**Record Creation Time:** 20231110T051308+0000

**Record Last Update:** 20241115T132932+0000

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## Ratings and Alerts

- ENCODE PROJECT External validation for lot: 3 is available under ENCODE ID: ENCAB405MHV - ENCODE <https://www.encodeproject.org/antibodies/ENCAB405MHV>

No alerts have been found for Rabbit Anti-Histone H3, trimethyl XP??? Monoclonal Antibody, Unconjugated, Clone D5A7.

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## Data and Source Information

**Source:** [Antibody Registry](#)

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## Usage and Citation Metrics

We found 25 mentions in open access literature.

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

Fan S, et al. (2024) Promoter DNA methylation and transcription factor condensation are linked to transcriptional memory in mammalian cells. *Cell systems*, 15(9), 808.

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

Liu CC, et al. (2024) Targeting EMSY-mediated methionine metabolism is a potential therapeutic strategy for triple-negative breast cancer. *Cell reports. Medicine*, 5(2), 101396.

Zhu R, et al. (2024) ACSS2 acts as a lactyl-CoA synthetase and couples KAT2A to function as a lactyltransferase for histone lactylation and tumor immune evasion. *Cell metabolism*.

Wang G, et al. (2024) Ethanol changes Nestin-promoter induced neural stem cells to disturb newborn dendritic spine remodeling in the hippocampus of mice. *Neural regeneration research*, 19(2), 416.

Niu N, et al. (2024) Tumor cell-intrinsic epigenetic dysregulation shapes cancer-associated fibroblasts heterogeneity to metabolically support pancreatic cancer. *Cancer cell*, 42(5), 869.

Fang L, et al. (2023) Methionine restriction promotes cGAS activation and chromatin untethering through demethylation to enhance antitumor immunity. *Cancer cell*, 41(6), 1118.

Chandhasin C, et al. (2023) TACH101, a first-in-class pan-inhibitor of KDM4 histone demethylase. *Anti-cancer drugs*.

Li HT, et al. (2023) RNA mis-splicing drives viral mimicry response after DNMTi therapy in SETD2-mutant kidney cancer. *Cell reports*, 42(1), 112016.

Zhang Y, et al. (2022) An antibody-based proximity labeling map reveals mechanisms of SARS-CoV-2 inhibition of antiviral immunity. *Cell chemical biology*, 29(1), 5.

Harpaz N, et al. (2022) Single-cell epigenetic analysis reveals principles of chromatin states in H3.3-K27M gliomas. *Molecular cell*, 82(14), 2696.

Yamamoto J, et al. (2022) Linkage of methionine addiction, histone lysine hypermethylation, and malignancy. *iScience*, 25(4), 104162.

Schniewind I, et al. (2022) Cellular plasticity upon proton irradiation determines tumor cell radiosensitivity. *Cell reports*, 38(8), 110422.

Villa E, et al. (2021) mTORC1 stimulates cell growth through SAM synthesis and m6A mRNA-dependent control of protein synthesis. *Molecular cell*, 81(10), 2076.

Yan R, et al. (2021) Decoding dynamic epigenetic landscapes in human oocytes using single-cell multi-omics sequencing. *Cell stem cell*, 28(9), 1641.

Zhu C, et al. (2021) Cancer-associated exportin-6 upregulation inhibits the transcriptionally repressive and anticancer effects of nuclear profilin-1. *Cell reports*, 34(7), 108749.

Bado IL, et al. (2021) The bone microenvironment increases phenotypic plasticity of ER+ breast cancer cells. *Developmental cell*, 56(8), 1100.

Araki R, et al. (2021) Low folate induces abnormal neuronal maturation and DNA hypomethylation of neuronal differentiation-related genes in cultured mouse neural stem and progenitor cells. *Heliyon*, 7(9), e08071.

Michowski W, et al. (2020) Cdk1 Controls Global Epigenetic Landscape in Embryonic Stem Cells. *Molecular cell*, 78(3), 459.

González-Rodríguez P, et al. (2020) SETD2 mutation in renal clear cell carcinoma suppress autophagy via regulation of ATG12. *Cell death & disease*, 11(1), 69.