

Resource Summary Report

Generated by RRID on Aug 2, 2026

H3K9me2-dmelanogaster

RRID:AB_449854

Type: Antibody

Proper Citation

Abcam Cat# ab1220, RRID:AB_449854

Antibody Information

URL: http://antibodyregistry.org/AB_449854

Proper Citation: Abcam Cat# ab1220, RRID:AB_449854

Target Antigen: H3K9me2

Host Organism: mouse

Clonality: monoclonal

Comments: ENCODE PROJECT External validation for lot# GR38973-3 is available under ENCODE ID: ENCAB638TXJ

Antibody Name: H3K9me2-dmelanogaster

Description: This monoclonal targets H3K9me2

Target Organism: drosophila melanogaster

Clone ID: Clone mAbcam 1220

Antibody ID: AB_449854

Vendor: Abcam

Catalog Number: ab1220

Record Creation Time: 20250820T012819+0000

Record Last Update: 20250820T110056+0000

Ratings and Alerts

- ENCODE PROJECT External validation for lot: 320150 is available under ENCODE ID: ENCAB783AQT - ENCODE

<https://www.encodeproject.org/antibodies/ENCAB783AQT>

No alerts have been found for H3K9me2-dmelanogaster.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 118 mentions in open access literature.

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

Liu Y, et al. (2025) DNA hypomethylation promotes UHRF1-and SUV39H1/H2-dependent crosstalk between H3K18ub and H3K9me3 to reinforce heterochromatin states. *Molecular cell*, 85(2), 394.

Jiang S, et al. (2025) JMJD2 regulates enhancer-promoter interactions via biomolecular condensate formation. *Nature genetics*.

Basso M, et al. (2025) Actin-based deformations of the nucleus control mouse multiciliated ependymal cell differentiation. *Developmental cell*, 60(5), 749.

Gupta S, et al. (2025) The nucleolar granular component mediates genome-nucleolus interactions and establishes their repressive chromatin states. *Molecular cell*, 85(11), 2165.

Pal M, et al. (2025) The establishment of nuclear organization in mouse embryos is orchestrated by multiple epigenetic pathways. *Cell*, 188(13), 3583.

Miles A, et al. (2025) The homeobox gene, *dmbx1a*, is required for the development and maintenance of bipolar and photoreceptor cells in the zebrafish retina. *Experimental eye research*, 110809.

Liu L, et al. (2025) CXCR4-LASP1-G9a-SNAIL axis drives NEPC transdifferentiation via induction of EMT and downregulation of REST. *Cell genomics*, 5(8), 100916.

Sankhe CS, et al. (2025) Matrix Stiffness Regulates TGF β 1-Induced α SMA Expression via a G9a-LATS-YAP Signaling Cascade. *FASEB bioAdvances*, 7(7), e70035.

Dillingham CM, et al. (2025) KDM3A and KDM3B regulate alternative splicing in mouse pluripotent stem cells. *iScience*, 28(6), 112612.

Han G, et al. (2024) Selective translation of nuclear mitochondrial respiratory proteins reprograms succinate metabolism in AML development and chemoresistance. *Cell stem cell*, 31(12), 1777.

Shu J, et al. (2024) EMF1 functions as a 3D chromatin modulator in Arabidopsis. *Molecular cell*, 84(24), 4729.

Wang Z, et al. (2024) Histone demethylase PHF8 promotes prostate cancer metastasis via the E2F1-SNAI1 axis. *The Journal of pathology*, 264(1), 68.

Larkin A, et al. (2024) Mapping the dynamics of epigenetic adaptation in *S. pombe* during heterochromatin misregulation. *Developmental cell*, 59(16), 2222.

Del Vecchio A, et al. (2024) PCGF6 controls murine Tuft cell differentiation via H3K9me2 modification independently of Polycomb repression. *Developmental cell*, 59(3), 368.

Charlton SJ, et al. (2024) The fork protection complex promotes parental histone recycling and epigenetic memory. *Cell*, 187(18), 5029.

Yu J, et al. (2024) A replisome-associated histone H3-H4 chaperone required for epigenetic inheritance. *Cell*, 187(18), 5010.

Stutzman AV, et al. (2024) Heterochromatic 3D genome organization is directed by HP1a- and H3K9-dependent and independent mechanisms. *Molecular cell*, 84(11), 2017.

Khanduja JS, et al. (2024) RNA quality control factors nucleate Ctr4/SUV39H and trigger constitutive heterochromatin assembly. *Cell*, 187(13), 3262.

Alves-Lopes JP, et al. (2023) Specification of human germ cell fate with enhanced progression capability supported by hindgut organoids. *Cell reports*, 42(1), 111907.

Yang J, et al. (2023) Exposure to high-sugar diet induces transgenerational changes in sweet sensitivity and feeding behavior via H3K27me3 reprogramming. *eLife*, 12.