Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 10, 2025

Histone H3 (acetyl K9 + K14 + K18 + K23 + K27) antibody - ChIP Grade

RRID:AB_873860 Type: Antibody

Proper Citation

(Abcam Cat# ab47915, RRID:AB_873860)

Antibody Information

URL: http://antibodyregistry.org/AB_873860

Proper Citation: (Abcam Cat# ab47915, RRID:AB_873860)

Target Antigen: Histone H3 (acetyl K9 + K14 + K18 + K23 + K27) antibody - ChIP Grade

Host Organism: rabbit

Clonality: polyclonal

Comments: validation status unknown, seller recommendations provided in 2012: ChIP,

WB; Western Blot; ChIP

Antibody Name: Histone H3 (acetyl K9 + K14 + K18 + K23 + K27) antibody - ChIP Grade

Description: This polyclonal targets Histone H3 (acetyl K9 + K14 + K18 + K23 + K27)

antibody - ChIP Grade

Target Organism: human

Antibody ID: AB_873860

Vendor: Abcam

Catalog Number: ab47915

Record Creation Time: 20231110T075402+0000

Record Last Update: 20241115T010731+0000

Ratings and Alerts

No rating or validation information has been found for Histone H3 (acetyl K9 + K14 + K18 + K23 + K27) antibody - ChIP Grade.

No alerts have been found for Histone H3 (acetyl K9 + K14 + K18 + K23 + K27) antibody - ChIP Grade.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 17 mentions in open access literature.

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

Wang H, et al. (2024) Nucleo-cytosolic acetyl-CoA drives tumor immune evasion by regulating PD-L1 in melanoma. Cell reports, 43(12), 115015.

Bhaskar A, et al. (2023) SIRT2 inhibition by AGK2 enhances mycobacteria-specific stem cell memory responses by modulating beta-catenin and glycolysis. iScience, 26(5), 106644.

Zhao C, et al. (2023) A fiber-enriched diet alleviates Staphylococcus aureus-induced mastitis by activating the HDAC3-mediated antimicrobial program in macrophages via butyrate production in mice. PLoS pathogens, 19(1), e1011108.

Bandopadhyay S, et al. (2023) Oncogene-mediated nuclear accumulation of lactate promotes epigenetic alterations to induce cancer cell proliferation. Journal of cellular biochemistry, 124(4), 495.

Jamsheer K M, et al. (2022) A negative feedback loop of TOR signaling balances growth and stress-response trade-offs in plants. Cell reports, 39(1), 110631.

Wu D, et al. (2021) An acetyl-histone vulnerability in PI3K/AKT inhibition-resistant cancers is targetable by both BET and HDAC inhibitors. Cell reports, 34(7), 108744.

Bressan RB, et al. (2021) Regional identity of human neural stem cells determines oncogenic responses to histone H3.3 mutants. Cell stem cell, 28(5), 877.

Abeywickrama-Samarakoon N, et al. (2020) Hepatitis Delta Virus histone mimicry drives the recruitment of chromatin remodelers for viral RNA replication. Nature communications, 11(1), 419.

Sessa A, et al. (2019) SETD5 Regulates Chromatin Methylation State and Preserves Global Transcriptional Fidelity during Brain Development and Neuronal Wiring. Neuron, 104(2), 271.

Schulthess J, et al. (2019) The Short Chain Fatty Acid Butyrate Imprints an Antimicrobial Program in Macrophages. Immunity, 50(2), 432.

Wu Q, et al. (2019) A chemical toolbox for the study of bromodomains and epigenetic signaling. Nature communications, 10(1), 1915.

Merk DJ, et al. (2018) Opposing Effects of CREBBP Mutations Govern the Phenotype of Rubinstein-Taybi Syndrome and Adult SHH Medulloblastoma. Developmental cell, 44(6), 709.

Sechet E, et al. (2018) Natural molecules induce and synergize to boost expression of the human antimicrobial peptide ?-defensin-3. Proceedings of the National Academy of Sciences of the United States of America, 115(42), E9869.

Arango D, et al. (2018) Acetylation of Cytidine in mRNA Promotes Translation Efficiency. Cell, 175(7), 1872.

Kalkat M, et al. (2018) MYC Protein Interactome Profiling Reveals Functionally Distinct Regions that Cooperate to Drive Tumorigenesis. Molecular cell, 72(5), 836.

Langlet F, et al. (2017) Selective Inhibition of FOXO1 Activator/Repressor Balance Modulates Hepatic Glucose Handling. Cell, 171(4), 824.

D'Urso A, et al. (2016) Set1/COMPASS and Mediator are repurposed to promote epigenetic transcriptional memory. eLife, 5.