Resource Summary Report

Generated by nidm-terms on Aug 11, 2026

Rabbit Anti-Mouse IgG H&L

RRID:AB_2614925
Type: Antibody

Proper Citation

Abcam Cat# ab46540, RRID:AB_2614925

Antibody Information

URL: http://antibodyregistry.org/AB_2614925

Proper Citation: Abcam Cat# ab46540, RRID:AB_2614925

Target Antigen: IgG H&L

Host Organism: rabbit

Clonality: polyclonal secondary

Comments: Applications: ELISA, WB

Antibody Name: Rabbit Anti-Mouse IgG H&L

Description: This polyclonal secondary targets IgG H&L

Target Organism: mouse

Antibody ID: AB_2614925

Vendor: Abcam

Catalog Number: ab46540

Record Creation Time: 20250820T020521+0000

Record Last Update: 20250820T124008+0000

Ratings and Alerts

- ENCODE PROJECT External validation for lot: 587909 is available under ENCODE ID: ENCAB294YUD - ENCODE https://www.encodeproject.org/antibodies/ENCAB294YUD

No alerts have been found for Rabbit Anti-Mouse IgG H&L.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 60 mentions in open access literature.

Listed below are recent publications. The full list is available at nidm-terms .

Gong S, et al. (2026) CFP1 promotes B cell class switch recombination through the regulation of S region germline transcription and proliferation. iScience, 29(6), 116255.

Renaud-Pageot C, et al. (2026) Non-centromeric CENP-A regulates epithelial-mesenchymal plasticity and heterogeneity in human cells. Cell reports, 45(6), 117405.

Agrawal R, et al. (2025) p53 regulates DREAM complex-mediated repression in a p21-independent manner. The EMBO journal, 44(8), 2279.

Gall LG, et al. (2025) Zebrafish glial-vascular interactions progressively expand over the course of brain development. iScience, 28(1), 111549.

Lee H, et al. (2025) TCR signaling via NFATc1 constrains IL-15-induced bystander activation of human memory CD8+ T cells. Immunity, 58(12), 2957.

Day CR, et al. (2025) Altered cofactor recruitment and nucleosome dynamics underlie bisphenol A's impact on ER?-mediated transcriptional bursting. iScience, 28(7), 112864.

Semenova G, et al. (2025) Genotoxic antibody-drug conjugates combined with Bcl-xL inhibitors enhance therapeutic efficacy in metastatic castration-resistant prostate cancer. bioRxiv : the preprint server for biology.

Khyzha N, et al. (2025) Profiling transcriptome composition and dynamics within nuclear compartments using SLAM-RT&Tag. Molecular cell, 85(7), 1366.

Chakraborty S, et al. (2025) Deletion of a single CTCF motif at the boundary of a chromatin domain with three FGF genes disrupts gene expression and embryonic development. Developmental cell, 60(13), 1838.

Khyzha N, et al. (2025) Protocol for the spatiotemporal profiling of RNA within nuclear compartments in human cell lines using SLAM-RT&Tag. STAR protocols, 6(3), 104017.

Duffié R, et al. (2025) Mex3a-dependent post-transcriptional silencing ensures olfactory receptor diversity and axon guidance specificity. Cell reports, 44(8), 115979.

Li J, et al. (2025) CRISPR-based screening identifies the role of KRAB-containing transcription factors ZIM3 and ZNF394 in human major zygotic genome activation. Cell reports, 44(8), 116015.

Nguyen TF, et al. (2025) Dynamic regulation of murine RNA polymerase III transcription during heat shock stress. Genetics, 230(1).

Datar GK, et al. (2025) Disparate leukemia mutations converge on nuclear phase-separated condensates. Cell, 188(25), 7118.

Ziegler KC, et al. (2025) The brain neurovascular epigenome and its association with dementia. Neuron.

Kim S, et al. (2024) DNA-guided transcription factor cooperativity shapes face and limb mesenchyme. Cell, 187(3), 692.

Ma S, et al. (2024) Spatial transcriptomic landscape unveils immunoglobin-associated senescence as a hallmark of aging. Cell, 187(24), 7025.

Wang L, et al. (2024) Chromatin landscape instructs precise transcription factor regulome during embryonic lineage specification. Cell reports, 43(5), 114136.

Hu R, et al. (2024) Expanding GABAergic Neuronal Diversity in iPSC-Derived Disease Models. bioRxiv : the preprint server for biology.

Gao R, et al. (2024) Defining a TFAP2C-centered transcription factor network during murine peri-implantation. Developmental cell, 59(9), 1146.