

Resource Summary Report

Generated by [NIF](#) on Aug 5, 2026

HA tag antibody - ChIP Grade

RRID:AB_307019

Type: Antibody

Proper Citation

Abcam Cat# ab9110, RRID:AB_307019

Antibody Information

URL: http://antibodyregistry.org/AB_307019

Proper Citation: Abcam Cat# ab9110, RRID:AB_307019

Target Antigen: HA tag antibody - ChIP Grade

Host Organism: rabbit

Clonality: polyclonal

Comments: validation status unknown, seller recommendations provided in 2012: ChIP, ChIP/Chip, ELISA, ICC, ICC/IF, IP, WB; Western Blot; Immunoprecipitation; ChIP; ELISA; Immunocytochemistry; Immunofluorescence

Antibody Name: HA tag antibody - ChIP Grade

Description: This polyclonal targets HA tag antibody - ChIP Grade

Antibody ID: AB_307019

Vendor: Abcam

Catalog Number: ab9110

Record Creation Time: 20250820T041711+0000

Record Last Update: 20250820T183316+0000

Ratings and Alerts

No rating or validation information has been found for HA tag antibody - ChIP Grade.

No alerts have been found for HA tag antibody - ChIP Grade.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 269 mentions in open access literature.

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

Lupton CJ, et al. (2026) Structure of the lysosomal KICSTOR-GATOR1-SAMTOR nutrient-sensing supercomplex. *Cell*.

Lyu Y, et al. (2026) FKF1 nuclear condensates control anti-florigen turnover and flowering onset in response to nitrogen availability in monocots. *Developmental cell*.

Milovanovi? D, et al. (2026) Tissue-specific restriction of transposon-derived regulatory elements safeguards cell-type identity. *Cell reports*, 45(1), 116817.

Wu L, et al. (2025) The cell-surface shared proteome of astrocytes and neurons and the molecular foundations of their multicellular interactions. *Neuron*, 113(16), 2599.

Fang W, et al. (2025) Histone deacetylase complex 1 facilitates hypocotyl elongation by attenuating acetylation and transcription via a PIF4-dependent process. *Cell reports*, 44(8), 116027.

Wang H, et al. (2025) FTO SUMOylation regulates the differentiation of bone marrow mesenchymal stromal cells in inflammatory bowel disease-induced bone loss. *Cell reports*, 44(7), 115953.

Du Q, et al. (2025) Bilirubin metabolism in the liver orchestrates antiviral innate immunity in the body. *Cell reports*, 44(4), 115481.

Ee LS, et al. (2025) Enhancer remodeling by OTX2 directs specification and patterning of mammalian definitive endoderm. *Developmental cell*.

Bond C, et al. (2025) Heterogeneity of late endosome/lysosomes shown by multiplexed DNA-PAINT imaging. *The Journal of cell biology*, 224(1).

Bryan E, et al. (2025) Nucleosomal asymmetry shapes histone mark binding and promotes poising at bivalent domains. *Molecular cell*, 85(3), 471.

Perr J, et al. (2025) RNA-binding proteins and glycoRNAs form domains on the cell surface for cell-penetrating peptide entry. *Cell*, 188(7), 1878.

Manoim-Wolkovitz JE, et al. (2025) Nonlinear high-activity neuronal excitation enhances odor discrimination. *Current biology : CB*, 35(7), 1521.

Magnitov MD, et al. (2025) ZNF143 is a transcriptional regulator of nuclear-encoded mitochondrial genes that acts independently of looping and CTCF. *Molecular cell*, 85(1), 24.

Li S, et al. (2025) The m5C methyltransferase NSUN2 promotes progression of acute myeloid leukemia by regulating serine metabolism. *Cell reports*, 44(5), 115661.

Härting N, et al. (2025) Generation of hiPSC lines with fusion tagged thyroid hormone receptor α isoforms to study isoform-specific actions of TR α 1 and TR α 2. *Stem cell research*, 86, 103720.

Broix L, et al. (2025) m6A RNA methylation-mediated control of global APC expression is required for local translation of β -actin and axon development. *Cell reports*, 44(6), 115727.

Fu X, et al. (2025) A cancer-specific antigen drives histone acetylation by stabilizing the acetyltransferases. *Cell reports*, 44(10), 116408.

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

Li H, et al. (2025) Mapping chromatin structure at base-pair resolution unveils a unified model of cis-regulatory element interactions. *Cell*, 188(25), 7175.

Elango R, et al. (2025) Two-ended recombination at a Flp-nickase-broken replication fork. *Molecular cell*, 85(1), 78.