

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

ChIP-Atlas

RRID:SCR_015511

Type: Tool

Proper Citation

ChIP-Atlas (RRID:SCR_015511)

Resource Information

URL: <http://chip-atlas.org/>

Proper Citation: ChIP-Atlas (RRID:SCR_015511)

Description: Database for visualizing and making use of public ChIP-seq data. ChIP-Atlas covers almost all public ChIP-seq experiments and data submitted to the SRA (Sequence Read Archives) in NCBI, DDBJ, or ENA.

Resource Type: software resource, data or information resource, web application, database

Keywords: chip-seq, sequence read archive, sequencing, FASEB list

Availability: Freely Available

Resource Name: ChIP-Atlas

Resource ID: SCR_015511

License: Creative Commons Attribution-Share Alike 4.0 International

License URLs: <https://dbarchive.biosciencedbc.jp/en/chip-atlas/lic.html>

Record Creation Time: 20220129T080326+0000

Record Last Update: 20260801T042748+0000

Ratings and Alerts

No rating or validation information has been found for ChIP-Atlas.

No alerts have been found for ChIP-Atlas.

Data and Source Information

Usage and Citation Metrics

We found 376 mentions in open access literature.

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

Esposito DV, et al. (2026) A non-coding ABO regulatory variant associated with VWF levels, thrombosis risk, and COVID-19 severity is topologically linked to ADAMTS13 in endothelial cells. HGG advances, 7(1), 100550.

Silva-Llanes I, et al. (2026) GASDERMIN D-mediated pyroptosis as a therapeutic target in TAU-dependent frontotemporal dementia mouse model. Journal of biomedical science, 33(1), 6.

Wang C, et al. (2025) The Oncogenic Role of TNFRSF12A in Colorectal Cancer and Pan-Cancer Bioinformatics Analysis. Cancer research and treatment, 57(1), 212.

Paul S, et al. (2025) Cooperation between the Hippo and MAPK pathway activation drives acquired resistance to TEAD inhibition. Nature communications, 16(1), 1743.

Yin B, et al. (2025) Genetic compensation response contributes to *Pleurodeles waltl* limb regeneration. NPJ Regenerative medicine, 10(1), 25.

Shalaby O, et al. (2025) Folliculin Deletion in the Mouse Kidney Results in Cystogenesis of the Loops of Henle via Aberrant TFEB Activation. The American journal of pathology, 195(9), 1643.

Jatzlau J, et al. (2025) Rare but specific: 5-bp composite motifs define SMAD binding in BMP signaling. BMC biology, 23(1), 79.

Honda N, et al. (2025) Implications of Mutant SOD1 on RNA Processing and Interferon Responses in Amyotrophic Lateral Sclerosis: Omics Data Analysis. Cureus, 17(3), e81045.

Arase M, et al. (2025) Multi-omics uncovers transcriptional programs of gut-resident memory CD4+ T cells in Crohn's disease. The Journal of experimental medicine, 222(11).

Dutta S, et al. (2025) Tubulin hyperacetylation drives HMGB1 nuclear exit via the ROS-PARP1 axis, leading to rotenone-induced G2/M arrest. The Journal of biological chemistry, 301(10), 110695.

Inoki T, et al. (2025) N-acetyltransferase 10 promotes glioblastoma malignancy via mRNA stabilization of jumonji and AT-rich interaction domain containing 2. The Journal of biological chemistry, 301(6), 108544.

Nakajima T, et al. (2025) In silico screening system based on a transcription factors regulatory network only using transcriptomic data. PloS one, 20(4), e0319971.

Bashkatov A, et al. (2025) Deep learning deciphers the related role of master regulators and G-quadruplexes in tissue specification. *Scientific reports*, 15(1), 23119.

Sompub K, et al. (2025) NF- κ B RelA regulates temporal oligodendrocyte differentiation in the postnatal brains. *Frontiers in cellular neuroscience*, 19, 1622874.

Sakai S, et al. (2025) In vivo transition in chromatin accessibility during differentiation of deep-layer excitatory neurons in the neocortex. *Development (Cambridge, England)*, 152(13).

Brown AC, et al. (2025) The cAMP responsive element modulator (CREM) transcription factor influences susceptibility to undernutrition and infection. *mBio*, 16(8), e0139025.

Xu ??? X, et al. (2025) ScReNI: Single-cell Regulatory Network Inference Through Integrating scRNA-seq and scATAC-seq Data. *Genomics, proteomics & bioinformatics*, 23(4).

Wang W, et al. (2025) BET inhibitor in combination with BCG vaccine enhances antitumor efficacy and orchestrates T cell reprogramming for melanoma. *Cell reports. Medicine*, 6(3), 101995.

Johansen VBI, et al. (2025) Regulation of LEAP2 by insulin and glucagon in mice and humans. *Cell reports. Medicine*, 6(3), 101996.

He B, et al. (2025) *Helicobacter pylori* CagA elevates FTO to induce gastric cancer progression via a "hit-and-run" paradigm. *Cancer communications (London, England)*, 45(5), 608.