## **Resource Summary Report**

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# **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: database, data or information resource, software resource, web application

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

Funding:

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: 20250430T060015+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

Source: SciCrunch Registry

### **Usage and Citation Metrics**

We found 303 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

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.

Xu J, et al. (2025) miRStart 2.0: enhancing miRNA regulatory insights through deep learningbased TSS identification. Nucleic acids research, 53(D1), D138.

De Paolis V, et al. (2025) An antisense-long-noncoding-RNA modulates p75NTR expression levels during neuronal polarization. iScience, 28(1), 111566.

Wang H, et al. (2025) Targeting EGFR-binding protein SLC7A11 enhancing antitumor immunity of T cells via inducing MHC-I antigen presentation in nasopharyngeal carcinoma. Cell death & disease, 16(1), 21.

Liu N, et al. (2025) RBM39 Enhances Cholangiocarcinoma Growth Through EZH2-mediated WNT7B/?-catenin Pathway. Cellular and molecular gastroenterology and hepatology, 19(1), 101404.

Waisman A, et al. (2024) The transcription factor OCT6 promotes the dissolution of the naïve pluripotent state by repressing Nanog and activating a formative state gene regulatory network. Scientific reports, 14(1), 10420.

Yoshioka H, et al. (2024) Alteration of Gene Expression in Pathological Keratinization of the Ocular Surface. Investigative ophthalmology & visual science, 65(6), 37.

Wei T, et al. (2024) FOXM1/DEPDC1 feedback loop promotes hepatocarcinogenesis and represents promising targets for cancer therapy. Cancer science, 115(9), 3041.

Zhang B, et al. (2024) Deubiquitinase USP7 stabilizes KDM5B and promotes tumor progression and cisplatin resistance in nasopharyngeal carcinoma through the ZBTB16/TOP2A axis. Cell death and differentiation, 31(3), 309.

Statello L, et al. (2024) The chromatin-associated IncREST ensures effective replication stress response by promoting the assembly of fork signaling factors. Nature

communications, 15(1), 978.

Gu J, et al. (2024) Noncanonical functions of microRNAs in the nucleus. Acta biochimica et biophysica Sinica, 56(2), 151.

Gualdrini F, et al. (2024) An integrative epigenome-based strategy for unbiased functional profiling of clinical kinase inhibitors. Molecular systems biology, 20(6), 626.

Mishima K, et al. (2024) Splice-switching antisense oligonucleotide controlling tumor suppressor REST is a novel therapeutic medicine for neuroendocrine cancer. Molecular therapy. Nucleic acids, 35(3), 102250.

Ross MO, et al. (2024) PTPN2 copper-sensing relays copper level fluctuations into EGFR/CREB activation and associated CTR1 transcriptional repression. Nature communications, 15(1), 6947.

Sugimoto H, et al. (2024) iTraNet: a web-based platform for integrated trans-omics network visualization and analysis. Bioinformatics advances, 4(1), vbae141.

Weinstock JS, et al. (2024) Gene regulatory network inference from CRISPR perturbations in primary CD4+ T cells elucidates the genomic basis of immune disease. Cell genomics, 4(11), 100671.

Ueda H, et al. (2024) Glycolysis-mTORC1 crosstalk drives proliferation of patient-derived endometrial cancer spheroid cells with ALDH activity. Cell death discovery, 10(1), 435.

Leonard MR, et al. (2024) Aiolos promotes CXCR3 expression on Th1 cells via positive regulation of IFN-?/STAT1 signaling. JCI insight, 10(1).

Malik S, et al. (2024) Differential carbonic anhydrase activities control EBV-induced B-cell transformation and lytic cycle reactivation. PLoS pathogens, 20(3), e1011998.

Hosoi H, et al. (2024) IGLL5 controlled by super-enhancer affects cell survival and MYC expression in mature B-cell lymphoma. Leukemia research reports, 21, 100451.