

Resource Summary Report

Generated by PRECISE-TBI on Sep 7, 2026

NCBI Epigenomics

RRID:SCR_006151

Type: Tool

Proper Citation

NCBI Epigenomics (RRID:SCR_006151)

Resource Information

URL: <https://www.ncbi.nlm.nih.gov/geo/>

Proper Citation: NCBI Epigenomics (RRID:SCR_006151)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on January 19, 2022.

Abbreviations: Epigenomics

Synonyms: NCBI Epigenomic Gateway, National Center for Biotechnology Information Epigenomics, NCBI Epigenomic Hub

Resource Type: data or information resource, database

Keywords: genome-wide map, dna, histone, modification, epigenomic, genome, gold standard, gene mapping, gene amplification, genetic code, gene library, dna fingerprinting, chromatin, histone modification, dna methylation, dnaase footprinting, genome wide association study, gene expression

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: NCBI Epigenomics

Resource ID: SCR_006151

Alternate IDs: nlx_151643, OMICS_01848, r3d100010782

Alternate URLs: <https://doi.org/10.17616/R34K7J>

Old URLs: <http://www.ncbi.nlm.nih.gov/epigenomics>

Record Creation Time: 20220129T080234+0000

Record Last Update: 20260905T133133+0000

Ratings and Alerts

No rating or validation information has been found for NCBI Epigenomics.

No alerts have been found for NCBI Epigenomics.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 14631 mentions in open access literature.

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

Kim CY, et al. (2026) Integrative multi-omics of matched primary and liver metastatic colorectal cancer organoids identifies putative molecular targets. *Translational oncology*, 63, 102592.

Pan R, et al. (2026) Basement membrane-associated gene P3H2 regulates osteosarcoma progression and epithelial-mesenchymal transition through AKT/mTOR signaling axis. *Translational oncology*, 63, 102569.

Feng H, et al. (2026) SV2B Promotes the Progression of TFE3-Rearranged Renal Cell Carcinoma by Interacting with HERC2 to Impede the Degradation of NF- κ B Subunits. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 13(1), e09443.

Martinez-Romero A, et al. (2026) Aggressive prostate cancer is associated with pericyte dysfunction. *Molecular oncology*, 20(4), 947.

Li F, et al. (2026) SDMap: a comprehensive database of spatial drug perturbation maps. *Nucleic acids research*, 54(D1), D1634.

Lin J, et al. (2026) HBEGF/EGFR pathway activation by hUC-MSCs improves cognitive outcomes in anti-NMDAR encephalitis. *Molecular therapy : the journal of the American Society of Gene Therapy*, 34(2), 916.

Paliwal A, et al. (2026) Dysmorphic neurons express markers of inhibitory glycinergic signaling in focal cortical dysplasia IIb. *Brain pathology (Zurich, Switzerland)*, 36(2), e70043.

Wang S, et al. (2026) Multi-omics characterization of cachexia-related genes reveals prognostic signatures and immune landscape in breast cancer. *Clinical & translational oncology : official publication of the Federation of Spanish Oncology Societies and of the National Cancer Institute of Mexico*, 28(2), 557.

Wang X, et al. (2026) Transgelin defines pro-tumorigenic cancer-associated fibroblasts in pancreatic cancer. *British journal of cancer*, 134(4), 577.

Angelotti A, et al. (2026) Phenotypes and mechanisms of dysfunctional cardiac T-lymphocytes in dilated cardiomyopathy patients. *Journal of molecular and cellular cardiology*, 212, 16.

Li L, et al. (2026) Hyperoside prevents osteoporosis by activating the PI3K/AKT signaling pathway to inhibit oxidative stress and promote osteogenesis. *Molecular medicine reports*, 33(2).

Wang F, et al. (2026) Regulation and reversal of paclitaxel resistance via the STAT1-mediated apoptotic pathway in ovarian cancer. *International journal of oncology*, 68(2).

Li W, et al. (2026) Ubiquitination of Rhomboid 5 Homolog 2 by Constitutive Photomorphogenic 1 Alleviates Hepatic Ischemia-reperfusion Injury by Regulating the Transforming Growth Factor- β Activating Kinase 1-C-Jun N-terminal Kinase/p38 Signaling Pathway. *Cellular and molecular gastroenterology and hepatology*, 20(4), 101695.

Yu X, et al. (2026) Identification of common genetic features and pathways for osteoporosis with frozen shoulder by integrated bioinformatics analysis and machine learning. *The Korean journal of physiology & pharmacology : official journal of the Korean Physiological Society and the Korean Society of Pharmacology*, 30(2), 125.

Li M, et al. (2026) The novel endogenous micropeptide XLH-36 binds Gemin4 to promote triple-negative breast cancer metastasis. *Oncogene*, 45(2), 264.

Moyung K, et al. (2026) Genome-wide nucleosome and transcription factor responses to genetic perturbations reveal chromatin-mediated mechanisms of transcriptional regulation. *Genome research*, 36(1), 115.

Liu Z, et al. (2026) PBX1 Improves Cognition and Reduces Amyloid- β Pathology in APP/PS1 Mice by Transcriptionally Activating the CRTC2-CREB Pathway. *Aging cell*, 25(1), e70311.

Xiaoqiang W, et al. (2026) The Biomarkers and Underlying Mechanisms of Mitochondrial and Programmed Cell Death Related Genes in Epilepsy by Combining Transcriptome and Single-Cell Sequencing. *Journal of Korean Neurosurgical Society*, 69(2), 196.

Liu F, et al. (2026) The Positive Feedback of lncRNA ANRIL/miRNA-339-5p/ZBTB7A Suppresses Metastasis of Nasopharyngeal Carcinoma Cells via SREBP1-FASN. *Cancer science*, 117(1), 105.

Su K, et al. (2026) Mendelian Randomization and ScTranscriptomics Reveal a Lipid Low Malignant Cells Driving Immunosuppression and Matrix Remodeling in Oral squamous cell carcinoma. *International dental journal*, 76(1), 104009.