Resource Summary Report

Generated by RRID on Aug 5, 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: database, data or information resource

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: 20260805T044132+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 11568 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Huang X, et al. (2026) Contribution of ferroptosis and SLC7A11 to light-induced photoreceptor degeneration. Neural regeneration research, 21(1), 406.

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-?B Subunits. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 13(1), e09443.

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.

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.

Zhang M, et al. (2026) Spatial Transcriptomic Characteristics of the Aging Human Ovary. Aging cell, 25(1), e70288.

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

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

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 J, et al. (2026) Correlation of G protein-coupled receptor and tumor microenvironment with gastric cancer outcomes and therapies. Chinese medical journal, 139(1), 65.

Yu N, et al. (2026) Changes in border-associated macrophages after stroke: Single-cell sequencing analysis. Neural regeneration research, 21(1), 346.

Cai S, et al. (2026) Identification and validation of novel inflammatory response-related diagnostic biomarkers in diabetic foot ulcers. Journal of diabetes investigation, 17(1), 103.

Zhang Y, et al. (2026) Similarities and differences between Alzheimer's disease and schizophrenia: drug target Mendelian randomization and transcriptome analysis. International journal of medical sciences, 23(1), 133.

Liu Q, et al. (2026) Cell fusion-related proteins AoHam4, AoHam8 and AoPP2A regulate hyphal fusion, conidiation, trap morphogenesis, and secondary metabolism in Arthrobotrys oligospora. Virulence, 17(1), 2605779.

Wang Y, et al. (2026) METTL3 inhibits LINC00312 to suppress osteoporosis progression in a YTHDF2?dependent manner. International journal of molecular medicine, 57(1).

Zhou P, et al. (2026) Exploring the role of cytochrome P450 family 1 subfamily B member 1 and quercetin in modulating neuropathic pain after spinal cord injury. Molecular medicine reports, 33(1).

Nakamura M, et al. (2026) Chromatin-associated DEK proteins maintain H3K27me3 balance and coordinate developmental transitions in plants. The New phytologist, 249(2), 930.

Xu L, et al. (2026) Identification of cuproptosis-related subtypes, construction of a prognosis model, and tumor microenvironment landscape in multiple myeloma. Translational oncology, 63, 102601.

Wang Y, et al. (2026) Identification of Potential Ferroptosis Biomarkers in Multiple Myeloma via WGCNA and Experiments. Cancer reports (Hoboken, N.J.), 9(1), e70446.