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

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 4732 mentions in open access literature.

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


Li Y, et al. (2022) MiR-34a-3p suppresses pulmonary vascular proliferation in acute pulmonary embolism rat by targeting DUSP1. Bioscience reports, 42(1).


Liu F, et al. (2022) DNAJC10 correlates with tumor immune characteristics and predicts the prognosis of glioma patients. Bioscience reports, 42(1).


Della Verde G, et al. (2022) Fms-like tyrosine kinase 3 is a regulator of the cardiac side population in mice. Life science alliance, 5(3).