

Resource Summary Report

Generated by SPARC SAWG on Aug 14, 2026

Albert Einstein College of Medicine Epigenomics Shared Core Facility

RRID:SCR_023284

Type: Tool

Proper Citation

Albert Einstein College of Medicine Epigenomics Shared Core Facility
(RRID:SCR_023284)

Resource Information

URL: <https://einsteinmed.edu/research/shared-facilities/cores/53/epigenomics/>

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Description: Part of Einstein Center for Epigenomics and Illumina CPro (certified service provider) laboratory, offers massively-parallel sequencing (MPS) including fully-automated library preparation, quality control and assurance, and number of assays to study the genome/epigenome. Data analytical services are provided by Computational Genomics Facility.

Abbreviations: ESF

Synonyms: Albert Einstein College of Medicine Epigenomics Shared Facility (ESF), Epigenomics Shared Facility (ESF)

Resource Type: core facility, service resource, access service resource

Keywords: USEdit, ABRF, massively-parallel sequencing, fully automated library preparation, quality control and assurance, genome, epigenome, data analytical services,

Resource Name: Albert Einstein College of Medicine Epigenomics Shared Core Facility

Resource ID: SCR_023284

Alternate IDs: ABRF_1686

Alternate URLs: <https://coremarketplace.org/?FacilityID=1686&citation=1>

Old URLs: <https://www.einsteinmed.edu/research/shared-facilities/cores/53/epigenomics/>

Record Creation Time: 20230216T050210+0000

Record Last Update: 20260814T043043+0000

Ratings and Alerts

No rating or validation information has been found for Albert Einstein College of Medicine Epigenomics Shared Core Facility.

No alerts have been found for Albert Einstein College of Medicine Epigenomics Shared Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Kaur M, et al. (2026) RASP: rapid antibody functional screening by pentavalent phage display. mAbs, 18(1), 2627708.

Molero AE, et al. (2026) Aberrant medial ganglionic eminence (MGE) GABAergic neurogenesis contributes to Huntington's disease pathogenesis. Neurobiology of disease, 221, 107297.

Shrivastava A, et al. (2026) Mutant KRAS-driven selective mRNA translation reveals mechanisms and therapeutic vulnerabilities in cancer. Cell reports, 45(6), 117520.

Dubey A, et al. (2025) Brain tumors induce widespread disruption of calvarial bone and alteration of skull marrow immune landscape. Nature neuroscience, 28(11), 2231.

Cunningham JL, et al. (2025) Ultrarare Variants in DNA Damage Repair Genes in Pediatric Acute-Onset Neuropsychiatric Syndrome or Acute Behavioral Regression in Neurodevelopmental Disorders. Developmental neuroscience, 47(4), 231.