Resource Summary Report

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Harvard University Chan School of Public Health Bioinformatics Core Facility

RRID:SCR_025373
Type: Tool

Proper Citation

Harvard University Chan School of Public Health Bioinformatics Core Facility (RRID:SCR_025373)

Resource Information

URL: https://bioinformatics.sph.harvard.edu/

Proper Citation: Harvard University Chan School of Public Health Bioinformatics Core Facility (RRID:SCR_025373)

Description: Provides single point of contact for Harvard researchers interested in bioinformatics support, applying genomic approaches together with established and developing methodologies from epidemiology, environmental health, biostatistics and bioinformatics to improve human health.Services include analysis all types of next-generation sequencing data from RNA-seq and single cell RNA-seq to variant sequencing (exome or whole genome), to ChIP-seq or bisulfite sequencing. Offers bioinformatics training program for Harvard researchers interested in learning basic data skills and analysis of high-throughput sequencing data.

Abbreviations: HSPH

Synonyms: HSPH Harvard Chan Bioinformatics Core, Harvard University HSPH Harvard Chan Bioinformatics Core

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

Keywords: ABRF, bioinformatics support, next-generation sequencing data, ChIP-seq data, bisulfite sequencing, data analysis,

Availability: Restricted

Resource Name: Harvard University Chan School of Public Health Bioinformatics Core Facility

Resource ID: SCR_025373

Alternate IDs: ABRF_2789

Alternate URLs: https://coremarketplace.org/?FacilityID=2789&citation=1

Record Creation Time: 20240525T053230+0000

Record Last Update: 20260801T191321+0000

Ratings and Alerts

No rating or validation information has been found for Harvard University Chan School of Public Health Bioinformatics Core Facility.

No alerts have been found for Harvard University Chan School of Public Health Bioinformatics Core Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Jachim SK, et al. (2026) Leveraging single cell multiomic analyses to identify gene regulatory networks that drive human articular cartilage cell fate. Osteoarthritis and cartilage.

Sullivan ZA, et al. (2025) Brain-immune interactions generate pathogen-specific sickness states. bioRxiv : the preprint server for biology.

Iuchi S, et al. (2025) The role of MKI67 in the regulation of 60S pre-ribosome nucleolar export, transcripts, energy supply, and apoptosis. bioRxiv : the preprint server for biology.

Pignatti E, et al. (2025) CXXC Finger Protein 1 drives BMP signaling and progenitor cell differentiation during limb development. Developmental biology.