

# Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

## University of Virginia Bioinformatics Core

RRID:SCR\_012718

Type: Tool

### Proper Citation

University of Virginia Bioinformatics Core (RRID:SCR\_012718)

### Resource Information

**URL:** <https://med.virginia.edu/bioinformatics-core/>

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**Description:** Core serves as a centralized resource for providing expert and timely bioinformatics consulting and data analysis solutions. The core offers services to investigators both within and outside UVA on both grant-funded and chargeback-based projects for management and analysis of large-scale biological datasets produced by high-throughput genomics experiments. The Core's mission is to build and maintain an infrastructure that enables the application of strong bioinformatics analysis with a measurable impact on the ability of UVA investigators to both publish their work and obtain new funding.

**Abbreviations:** UVa Bioinformatics Core

**Synonyms:** , UVa, University of Virginia, core facility, School of Medicine, Bioinformatics

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

**Keywords:** core facility, data analysis, infrastructure,

**Availability:** Available to external user

**Resource Name:** University of Virginia Bioinformatics Core

**Resource ID:** SCR\_012718

**Alternate IDs:** SciEx\_8923

**Old URLs:** <http://www.scienceexchange.com/facilities/bioinformatics-core-virginia>

**Record Creation Time:** 20220129T080311+0000

**Record Last Update:** 20260801T191203+0000

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## Ratings and Alerts

No rating or validation information has been found for University of Virginia Bioinformatics Core.

No alerts have been found for University of Virginia Bioinformatics Core.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Cook SR, et al. (2025) A 3D-printed multi-compartment organ-on-chip platform with a tubing-free pump models communication with the lymph node. Lab on a chip, 25(2), 155.

Boulton A, et al. (2025) The MLLT3 YEATS domain is a dual reader of histone marks (H3K9/18/27ac/cr) and ncRNA (7SK), linking epigenetic and RNA signaling to regulate hematopoiesis. bioRxiv : the preprint server for biology.

Uddin MJ, et al. (2024) Investigating the impact of antibiotic-induced dysbiosis on protection from Clostridium difficile colitis by mouse colonic innate lymphoid cells. mBio, 15(3), e0333823.

Pierce R, et al. (2024) Persistent dysbiosis of duodenal microbiota in patients with controlled pediatric Crohn's disease after resolution of inflammation. Scientific reports, 14(1), 12668.

Cook SR, et al. (2024) A 3D-printed multi-compartment organ-on-chip platform with a tubing-free pump models communication with the lymph node. bioRxiv : the preprint server for biology.

Bingham GC, et al. (2023) High-dimensional comparison of monocytes and T cells in post-COVID and idiopathic pulmonary fibrosis. Frontiers in immunology, 14, 1308594.