

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 8, 2026

UVa Genomics Core Facility

RRID:SCR_012197

Type: Tool

Proper Citation

UVa Genomics Core Facility (RRID:SCR_012197)

Resource Information

URL: <http://www.scienceexchange.com/facilities/genomics-core-facility-virginia>

Proper Citation: UVa Genomics Core Facility (RRID:SCR_012197)

Description: The GCF provides second generation genome sequencing service to all members of the scientific community. Using the 454 Life Sciences/Roche Genome Sequencer FLX, we facilitate data collection and analysis.

Abbreviations: UVa GCF

Synonyms: University of Virginia Genomics Core Facility

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

Resource Name: UVa Genomics Core Facility

Resource ID: SCR_012197

Alternate IDs: SciEx_10443

Record Creation Time: 20220129T080309+0000

Record Last Update: 20260808T190616+0000

Ratings and Alerts

No rating or validation information has been found for UVa Genomics Core Facility.

No alerts have been found for UVa Genomics Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.

Jan N, et al. (2021) Fecal Microbiota Transplantation Increases Colonic IL-25 and Dampens Tissue Inflammation in Patients with Recurrent Clostridioides difficile. mSphere, 6(5), e0066921.