## **Resource Summary Report**

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# Indiana University Center for Genomics and Bioinformatics Core Facility

RRID:SCR\_017165 Type: Tool

**Proper Citation** 

Indiana University Center for Genomics and Bioinformatics Core Facility (RRID:SCR\_017165)

#### **Resource Information**

URL: https://cgb.indiana.edu

**Proper Citation:** Indiana University Center for Genomics and Bioinformatics Core Facility (RRID:SCR\_017165)

**Description:** CGB offers range of genomic services, including high-throughput DNA/RNA extraction, library preparation, next-generation sequencing, and bioinformatic analysis.

Abbreviations: CGB

Synonyms: Indiana University Bloomington Center for Genomics and Bioinformatics

**Resource Type:** data or information resource, core facility, service resource, access service resource

**Keywords:** ABRF, genomic services, DNA/RNA extraction, library preparation, nextgeneration sequencing, bioinformatic analysis,

Funding:

Availability: Restricted

Resource Name: Indiana University Center for Genomics and Bioinformatics Core Facility

Resource ID: SCR\_017165

Alternate IDs: SCR\_025534, ABRF\_2841

Alternate URLs: https://indianactsi.org/servicecores/core/20/, https://coremarketplace.org/?FacilityID=2841&citation=1

Record Creation Time: 20220129T080334+0000

Record Last Update: 20250502T060429+0000

### **Ratings and Alerts**

No rating or validation information has been found for Indiana University Center for Genomics and Bioinformatics Core Facility.

No alerts have been found for Indiana University Center for Genomics and Bioinformatics Core Facility.

Data and Source Information

Source: <u>SciCrunch Registry</u>

#### **Usage and Citation Metrics**

We found 3 mentions in open access literature.

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

Kelley BR, et al. (2020) Whole-Genome Sequencing and Bioinformatic Analysis of Environmental, Agricultural, and Human Campylobacter jejuni Isolates From East Tennessee. Frontiers in microbiology, 11, 571064.

Luhur A, et al. (2020) Adapting Drosophila melanogaster Cell Lines to Serum-Free Culture Conditions. G3 (Bethesda, Md.), 10(12), 4541.

Mushinski RM, et al. (2020) Nitrogen cycling microbiomes are structured by plant mycorrhizal associations with consequences for nitrogen oxide fluxes in forests. Global change biology, 27(5), 1068.