Resource Summary Report

Generated by dkNET on Apr 26, 2025

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, access service resource, service resource

Keywords: ABRF, genomic services, DNA/RNA extraction, library preparation, next-generation 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: 20250426T060619+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: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

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.