

Resource Summary Report

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Cleveland Clinic Lerner Research Institute Microbial Sequencing and Analytics Core Facility

RRID:SCR_026609

Type: Tool

Proper Citation

Cleveland Clinic Lerner Research Institute Microbial Sequencing and Analytics Core Facility (RRID:SCR_026609)

Resource Information

URL: <https://www.lerner.ccf.org/cores/microbial-sequencing/>

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Description: Core provides services, from DNA and RNA extraction from diverse sample types such as tissue, stool, and bacterial isolates to next-generation sequencing, including whole-genome sequencing, total RNA-seq, metatranscriptomics, and dual RNA-seq, along with expert bioinformatics support. Core expertise spans 16S rRNA amplicon sequencing, shotgun metagenomics, metatranscriptomics, meta-omics analysis of dual RNA-seq, and single-cell RNA-seq data. Core also assists with experimental design and power calculations to ensure statistically robust results, offering consultation at every stage, from experimental design to data interpretation and troubleshooting.

Abbreviations: MSAAR

Synonyms: Microbial Sequencing and Analytics Resource (MSAAR)

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

Keywords: ABRF, DNA and RNA extraction, next-generation sequencing, whole-genome sequencing, total RNA-seq, metatranscriptomics, dual RNA-seq, bioinformatics support,

Resource Name: Cleveland Clinic Lerner Research Institute Microbial Sequencing and Analytics Core Facility

Resource ID: SCR_026609

Alternate IDs: ABRF_3099

Alternate URLs: <https://coremarketplace.org/?FacilityID=3099&citation=1>

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Ratings and Alerts

No rating or validation information has been found for Cleveland Clinic Lerner Research Institute Microbial Sequencing and Analytics Core Facility.

No alerts have been found for Cleveland Clinic Lerner Research Institute Microbial Sequencing and Analytics Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.