

Resource Summary Report

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University of Texas at Austin Genomic Sequencing and Analysis Core Facility

RRID:SCR_021713

Type: Tool

Proper Citation

University of Texas at Austin Genomic Sequencing and Analysis Core Facility
(RRID:SCR_021713)

Resource Information

URL: <https://wikis.utexas.edu/display/GSAF/Home+Page>

Proper Citation: University of Texas at Austin Genomic Sequencing and Analysis Core Facility (RRID:SCR_021713)

Description: Full service sequencing core that provides both Sanger and Next Generation Sequencing services, in addition to library prep and quality check services, open access to equipment and training. Works directly with the bioinformatics core who can provide assist in any of your analysis needs to data storage solutions.

Abbreviations: GSAF

Synonyms: The University of Texas at Austin Genomic Sequencing and Analysis Facility, Genomic Sequencing and Analysis Facility

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

Keywords: USEDit, ABRF, sequencing core, Sanger Sequencing service, Next Generation Sequencing service, library preparation, quality check service

Resource Name: University of Texas at Austin Genomic Sequencing and Analysis Core Facility

Resource ID: SCR_021713

Alternate IDs: ABRF_1209

Alternate URLs: <https://coremarketplace.org/?FacilityID=1209>

Record Creation Time: 20220129T080357+0000

Record Last Update: 20261017T203700+0000

Ratings and Alerts

No rating or validation information has been found for University of Texas at Austin Genomic Sequencing and Analysis Core Facility.

No alerts have been found for University of Texas at Austin Genomic Sequencing and Analysis Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.