

# Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

## University of Texas at Austin Computational Biology and Bioinformatics Core Facility

RRID:SCR\_022688

Type: Tool

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### Proper Citation

University of Texas at Austin Computational Biology and Bioinformatics Core Facility  
(RRID:SCR\_022688)

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### Resource Information

**URL:** <https://research.utexas.edu/cbrs/cores/cbb/>

**Proper Citation:** University of Texas at Austin Computational Biology and Bioinformatics Core Facility (RRID:SCR\_022688)

**Description:** Provides support for students, postdoctoral fellows, and faculty interested in the use of computational approaches to solving biological problems. Our goal is to lower as much as possible the threshold to enter the -omics area of the life sciences.

**Synonyms:** CBRS-Bioinformatics Consulting Group

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

**Keywords:** USEDit, ABRF,

**Resource Name:** University of Texas at Austin Computational Biology and Bioinformatics Core Facility

**Resource ID:** SCR\_022688

**Alternate IDs:** ABRF\_1505

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=1505&citation=1>

**Record Creation Time:** 20220819T050145+0000

**Record Last Update:** 20260829T183331+0000

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### Ratings and Alerts

No rating or validation information has been found for University of Texas at Austin Computational Biology and Bioinformatics Core Facility.

No alerts have been found for University of Texas at Austin Computational Biology and Bioinformatics Core Facility.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Engels SM, et al. (2024) Particulate matter composition drives differential molecular and morphological responses in lung epithelial cells. PNAS nexus, 3(1), pgad415.

Zhao P, et al. (2023) Femtosecond laser microdissection for isolation of regenerating C. elegans neurons for single-cell RNA sequencing. Nature methods, 20(4), 590.