# **Resource Summary Report**

Generated by NIF on Apr 15, 2025

# University of Texas at Austin Cryo Electron Microscopy Core Facility

RRID:SCR\_022951

Type: Tool

## **Proper Citation**

University of Texas at Austin Cryo Electron Microscopy Core Facility (RRID:SCR\_022951)

#### **Resource Information**

URL: https://research.utexas.edu/cbrs/cores/cem/

**Proper Citation:** University of Texas at Austin Cryo Electron Microscopy Core Facility (RRID:SCR\_022951)

**Description:** Offers training, assisted and unassisted use of microscopes and equipment, as well as consultations and collaborations. Support includes single particle cryo-EM, Cryo-ET, Micro-ED, simple characterization of samples in vitrified aqueous mixtures (e.g. nanoparticle size, etc.).

**Synonyms:** UTA CBRS Sauer Structural Biology Laboratory, University of Texas at Austin UTA CBRS Sauer Structural Biology Laboratory

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

**Keywords:** USEDit, ABRF, single particle cryo-EM, Cryo-ET, Electron Microscopy, Micro-ED, characterization of samples in vitrified aqueous mixtures, nanoparticle size

**Funding:** 

Resource Name: University of Texas at Austin Cryo Electron Microscopy Core Facility

Resource ID: SCR\_022951

Alternate IDs: ABRF\_1625

Alternate URLs: https://coremarketplace.org/?FacilityID=1625&citation=1

**Record Creation Time:** 20221109T050203+0000

Record Last Update: 20250412T060521+0000

## **Ratings and Alerts**

No rating or validation information has been found for University of Texas at Austin Cryo Electron Microscopy Core Facility.

No alerts have been found for University of Texas at Austin Cryo Electron Microscopy Core Facility.

#### Data and Source Information

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We found 4 mentions in open access literature.

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

Afsar M, et al. (2023) Cryo-EM structures of Uba7 reveal the molecular basis for ISG15 activation and E1-E2 thioester transfer. Nature communications, 14(1), 4786.

Rawal Y, et al. (2023) Structural insights into BCDX2 complex function in homologous recombination. Nature, 619(7970), 640.

Musalgaonkar S, et al. (2023) The Ribosome Assembly Factor Reh1 is Released from the Polypeptide Exit Tunnel in the Pioneer Round of Translation. bioRxiv: the preprint server for biology.

Jara MO, et al. (2022) Oral Delivery of Niclosamide as an Amorphous Solid Dispersion That Generates Amorphous Nanoparticles during Dissolution. Pharmaceutics, 14(12).