

Resource Summary Report

Generated by [NIF](#) on Aug 8, 2026

UTSA RCMI Computational Systems Biology Core

RRID:SCR_010148

Type: Tool

Proper Citation

UTSA RCMI Computational Systems Biology Core (RRID:SCR_010148)

Resource Information

URL: <http://utsa.eagle-i.net/i/00000135-2f42-aa68-d7c8-cf3780000000>

Proper Citation: UTSA RCMI Computational Systems Biology Core (RRID:SCR_010148)

Description: Core facility that provides the following services: Computational systems biology data analysis assistance, Biological computational software installation service. The objective of the Computational Systems Biology Core facility (CSBC) is to provide computational support for basic and translational health research at UTSA with the following specific aims: - Build the computational infrastructure to support modeling and simulation of biological systems - Live cell imaging - Protein Biomarker research

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

Keywords: nucleic acid microarray assay, pathway data analysis, protein assay, gene expression analysis assay, computational modeling technique

Resource Name: UTSA RCMI Computational Systems Biology Core

Resource ID: SCR_010148

Alternate IDs: nlx_156628

Record Creation Time: 20220129T080257+0000

Record Last Update: 20260808T042713+0000

Ratings and Alerts

No rating or validation information has been found for UTSA RCMI Computational Systems Biology Core.

No alerts have been found for UTSA RCMI Computational Systems Biology Core.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.