

Resource Summary Report

Generated by SPARC SAWG on Sep 14, 2026

University of Florida UF Scripps Biomedical Research Bioinformatics and Statistics Core Facility

RRID:SCR_023048

Type: Tool

Proper Citation

University of Florida UF Scripps Biomedical Research Bioinformatics and Statistics Core Facility (RRID:SCR_023048)

Resource Information

URL: <https://scripps.ufl.edu/cores-and-technologies/bioinformatics/>

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Description: Core provides data management and analysis services for UF Scripps community as well as for external collaborators. Offers scientific support to assist in experimental design, analysis, integration and dissemination of biomedical data and knowledge. Supports data analysis for samples processed at Genomics, Proteomics, and Neurobehavior cores.

Synonyms: University of Florida UF Scripps Biomedical Research Bioinformatics and Statistics Core, UF Scripps Biomedical Research-Bioinformatics and Statistics Core

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

Keywords: USEDit, ABRF, Bioinformatics and Statistics Core, biomedical data analysis, data management and analysis services

Resource Name: University of Florida UF Scripps Biomedical Research Bioinformatics and Statistics Core Facility

Resource ID: SCR_023048

Alternate IDs: ABRF_1648

Alternate URLs: <https://coremarketplace.org/?FacilityID=1648&citation=1>

Record Creation Time: 20221215T050202+0000

Record Last Update: 20260912T200422+0000

Ratings and Alerts

No rating or validation information has been found for University of Florida UF Scripps Biomedical Research Bioinformatics and Statistics Core Facility.

No alerts have been found for University of Florida UF Scripps Biomedical Research Bioinformatics and Statistics Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Eldeeb MH, et al. (2026) Mitochondrial respirasome-like supercomplexes support metabolic flexibility in yeast. Nature communications, 17(1).

Wang S, et al. (2026) Structural basis of TACO1-mediated efficient mitochondrial translation. Nature communications, 17(1).

Zhao Z, et al. (2025) Direct interoceptive input to the insular cortex shapes learned feeding behavior. bioRxiv : the preprint server for biology.

Dagar S, et al. (2024) Ribosome Profiling and Mass Spectrometry Reveal Widespread Mitochondrial Translation Defects in a Striatal Cell Model of Huntington Disease. Molecular & cellular proteomics : MCP, 100746.

Brischigliaro M, et al. (2024) The human mitochondrial translation factor TACO1 alleviates mitoribosome stalling at polyproline stretches. Nucleic acids research, 52(16), 9710.