

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core Facility

RRID:SCR_026404

Type: Tool

Proper Citation

University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core Facility (RRID:SCR_026404)

Resource Information

URL: <https://touchstonelabs.org/metabolic-phenotyping-core>

Proper Citation: University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core Facility (RRID:SCR_026404)

Description: Academic core facility provides analytical and phenotypical measures to the scientific community. Provides services related to metabolic disorders (diabetes and obesity), cancer, aging neurological disorders, etc. Provides letters of support for grant proposals and research applications.

Synonyms: , UTSW Metabolic Phenotyping Core, University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core

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

Keywords: ABRF, analytical and phenotypical services, metabolic disorders,

Availability: Open

Resource Name: University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core Facility

Resource ID: SCR_026404

Alternate IDs: ABRF_3056

Alternate URLs: <https://coremarketplace.org/?FacilityID=3056&citation=1>

Record Creation Time: 20250211T060340+0000

Record Last Update: 20260905T133521+0000

Ratings and Alerts

No rating or validation information has been found for University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core Facility.

No alerts have been found for University of Texas Southwestern Medical Center Dallas Metabolic Phenotyping Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Crewe C, et al. (2026) Adipocytes signal to recruit specific mRNAs from surrounding cells to restore expression deficits. Nature communications, 17(1).