

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

Kendrick Labs Core Facility

RRID:SCR_018271

Type: Tool

Proper Citation

Kendrick Labs Core Facility (RRID:SCR_018271)

Resource Information

URL: <http://www.kendricklabs.com>

Proper Citation: Kendrick Labs Core Facility (RRID:SCR_018271)

Description: Contract Research Org. specializing in custom protein analysis. Services Provided include SDS PAGE analysis, Western Blotting, Mass Spectrometry, protein staining (silver,coomassie) for academia, federal institutions and industry.

Synonyms: Private Company - Kendrick Labs, Inc., Kendrick Labs, Inc

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

Keywords: custom protein analysis, SDS Page analysis, Western Blotting, Mass Spectrometry, silver protein staining, coomassie protein staining, USEDit, ABRF

Availability: Open

Resource Name: Kendrick Labs Core Facility

Resource ID: SCR_018271

Alternate IDs: SCR_002914, ABRF_514, nif-0000-30014

Alternate URLs: <https://coremarketplace.org/?FacilityID=514>

Record Creation Time: 20220129T080339+0000

Record Last Update: 20260731T043032+0000

Ratings and Alerts

No rating or validation information has been found for Kendrick Labs Core Facility.

No alerts have been found for Kendrick Labs 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 [PRECISE-TBI](#) .

Ou L, et al. (2017) Proteomic analysis of mucopolysaccharidosis I mouse brain with two-dimensional polyacrylamide gel electrophoresis. Molecular genetics and metabolism, 120(1-2), 101.

Ngounou Wetie AG, et al. (2015) Comparative two-dimensional polyacrylamide gel electrophoresis of the salivary proteome of children with autism spectrum disorder. Journal of cellular and molecular medicine, 19(11), 2664.

Nguyen AT, et al. (2011) The prototype HIV-1 maturation inhibitor, bevirimat, binds to the CA-SP1 cleavage site in immature Gag particles. Retrovirology, 8, 101.

Wong CC, et al. (2007) Global analysis of posttranslational protein arginylation. PLoS biology, 5(10), e258.

Pearce MJ, et al. (2006) Identification of substrates of the Mycobacterium tuberculosis proteasome. The EMBO journal, 25(22), 5423.