

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

Yale Center for Genome Analysis Core Facility

RRID:SCR_027324

Type: Tool

Proper Citation

Yale Center for Genome Analysis Core Facility (RRID:SCR_027324)

Resource Information

URL: <https://medicine.yale.edu/genetics/research/ycga/>

Proper Citation: Yale Center for Genome Analysis Core Facility (RRID:SCR_027324)

Description: Core performs next generation sequencing for RNA and DNA samples, as well as provides support for downstream analysis.

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

Keywords: ABRF, next generation sequencing, RNA, DNA

Resource Name: Yale Center for Genome Analysis Core Facility

Resource ID: SCR_027324

Alternate IDs: ABRF_4092

Alternate URLs: <https://coremarketplace.org/?FacilityID=4092>

Record Creation Time: 20250815T054452+0000

Record Last Update: 20260905T133545+0000

Ratings and Alerts

No rating or validation information has been found for Yale Center for Genome Analysis Core Facility.

No alerts have been found for Yale Center for Genome Analysis 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 [PRECISE-TBI](#) .

Kelley VM, et al. (2026) Loss of JAK1 Function Causes G2-M Cell-Cycle Defects Vulnerable to KIF18A Inhibition. Cancer research, 86(9), 2301.