

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

MGH CCIB DNA Sequencing Core

RRID:SCR_009915

Type: Tool

Proper Citation

MGH CCIB DNA Sequencing Core (RRID:SCR_009915)

Resource Information

URL: <http://harvard.eagle-i.net/i/0000012d-ee0d-d869-b2b9-4d8780000000>

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Description: Core facility that provides the following services: DNA sequencing service, High volume DNA sequencing, Plasmid verification and primer walking service, PCR purification service, Microsatellite analysis.

The DNA Sequencing Core at the Massachusetts General Hospital functions both as a small-scale sequencing facility and a high-throughput center for large-scale sequencing projects.

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

Keywords: dna sequencing

Resource Name: MGH CCIB DNA Sequencing Core

Resource ID: SCR_009915

Alternate IDs: nlx_156380

Alternate URLs:

http://www.partners.org/researchcores/DNA%20Synthesis/ccib_dna_mgh.asp,
<https://dnacore.mgh.harvard.edu/>, http://ccib.mgh.harvard.edu/dna_core

Record Creation Time: 20220129T080255+0000

Record Last Update: 20260905T133336+0000

Ratings and Alerts

No rating or validation information has been found for MGH CCIB DNA Sequencing Core.

No alerts have been found for MGH CCIB DNA Sequencing Core.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Bolli E, et al. (2026) CCL3 is produced by aged neutrophils across cancers and promotes tumor growth. Cancer cell, 44(3), 624.

Armani-Tourret M, et al. (2024) Selection of epigenetically privileged HIV-1 proviruses during treatment with panobinostat and interferon- γ . Cell, 187(5), 1238.