## **Resource Summary Report**

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# 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

**Proper Citation:** MGH CCIB DNA Sequencing Core (RRID:SCR\_009915)

**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: core facility, access service resource, service resource

Keywords: dna sequencing

**Funding:** 

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: 20250411T055406+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 1 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Armani-Tourret M, et al. (2024) Selection of epigenetically privileged HIV-1 proviruses during treatment with panobinostat and interferon-?2a. Cell, 187(5), 1238.