

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

Massachusetts General Hospital NGS Core Facility

RRID:SCR_026362

Type: Tool

Proper Citation

Massachusetts General Hospital NGS Core Facility (RRID:SCR_026362)

Resource Information

URL: <https://nextgen.mgh.harvard.edu/Pricing.html>

Proper Citation: Massachusetts General Hospital NGS Core Facility (RRID:SCR_026362)

Description: Core offers sequencing, upstream library construction (bulk RNA/DNA, single-cell), and post-sequencing analysis. Core uses Illumina NextSeq 2000 sequencers (and a MiSeq). Single-cell (fresh, frozen, FFPE) libraries (3' & 5' RNA-Seq, CITE-Seq, ATAC-Seq, Multiome, FLEX) are made with 10X Chromium X.

Synonyms: , Massachusetts General Hospital MGH NGS Core, Massachusetts General Hospital NGS Core, MGH NGS Core

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

Keywords: ABRF, sequencing, upstream library construction, post-sequencing analysis,

Resource Name: Massachusetts General Hospital NGS Core Facility

Resource ID: SCR_026362

Alternate IDs: ABRF_3023

Alternate URLs: <https://coremarketplace.org/?FacilityID=3023&citation=1>

Record Creation Time: 20250201T053323+0000

Record Last Update: 20260729T044637+0000

Ratings and Alerts

No rating or validation information has been found for Massachusetts General Hospital NGS Core Facility.

No alerts have been found for Massachusetts General Hospital NGS Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.