

# Resource Summary Report

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## University of California at San Diego Institute for Genomic Medicine Genomics Center Core Facility

RRID:SCR\_022740

Type: Tool

### Proper Citation

University of California at San Diego Institute for Genomic Medicine Genomics Center Core Facility (RRID:SCR\_022740)

### Resource Information

**URL:** <http://igm.ucsd.edu/genomics/>

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**Description:** Provides services including sequencing library preparation and sequencing on Illumina MiSeq and NovaSeq 6000 platforms. Supports single cell sequencing on 10X Genomics Chromium Controller. Provides Illumina Infinium Beadchips, which includes variety of whole genome genotyping arrays as well as Infinium MethylationEPIC BeadChip.

**Synonyms:** UCSD IGM Genomics Center

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

**Keywords:** USEEdit, ABRF, sequencing library preparation, sequencing, single cell sequencing, whole genome genotyping arrays

**Funding:** NIH Office of the Director S10 OD026929;  
NIDDK P30DK063491;  
NCI P30CA023100;  
NIDDK P30DK120515

**Resource Name:** University of California at San Diego Institute for Genomic Medicine Genomics Center Core Facility

**Resource ID:** SCR\_022740

**Alternate IDs:** ABRF\_1025

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=1025&citation=1>

**Record Creation Time:** 20220915T050143+0000

**Record Last Update:** 20260808T190706+0000

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## Ratings and Alerts

No rating or validation information has been found for University of California at San Diego Institute for Genomic Medicine Genomics Center Core Facility.

No alerts have been found for University of California at San Diego Institute for Genomic Medicine Genomics Center Core Facility.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [NIF](#) .

Hakim A, et al. (2026) A high-throughput, end-to-end pipeline for extracellular miRNA biomarker discovery from human biofluids. Cell reports methods, 101502.

Chang H, et al. (2025) Single-cell transcriptomics reveals immune remodeling of the murine lung microenvironment following chronic house dust mite exposure. Frontiers in immunology, 16, 1652539.