

# Resource Summary Report

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

## Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility (IGC)

RRID:SCR\_014842

Type: Tool

---

### Proper Citation

Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility (IGC) (RRID:SCR\_014842)

---

### Resource Information

**URL:** <http://www.salk.edu/science/core-facilities/integrative-genomics-and-bioinformatics-core/>

**Proper Citation:** Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility (IGC) (RRID:SCR\_014842)

**Description:** Core facility established to assist the Salk community with integrating genomics data into their research. The primary focus of the core is to provide analysis support for next-generation sequencing applications.

**Abbreviations:** SALK IGC, IGC

**Synonyms:** , Integrative Genomics, Salk, Core Facility, Institute, Razavi Newman, UCSD, Bioinformatics

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

**Keywords:** core facility, gene, genomic, genomic data, analysis, consultation, applications

**Funding:** NCI CA014195;  
Helmsley Trust ;  
Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility

**Availability:** Open

**Resource Name:** Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility (IGC)

**Resource ID:** SCR\_014842

**Record Creation Time:** 20220129T080322+0000

## Ratings and Alerts

No rating or validation information has been found for Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility (IGC).

No alerts have been found for Salk Institute Razavi Newman Integrative Genomics and Bioinformatics Core Facility (IGC).

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We found 940 mentions in open access literature.

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

Adams P, et al. (2025) Nucleosome stability safeguards cell identity, stress resilience and healthy aging. Research square.

Miller CN, et al. (2025) A single-nuclei transcriptome census of the Arabidopsis maturing root identifies that MYB67 controls phellem cell maturation. Developmental cell, 60(9), 1377.

Dufresne S, et al. (2025) Leveraging autophagy and pyrimidine metabolism to target pancreatic cancer. bioRxiv : the preprint server for biology.

Tanaka H, et al. (2025) Nucleosome stability safeguards cell identity, stress resilience and healthy aging. bioRxiv : the preprint server for biology.

Gallant RM, et al. (2025) Fluoxetine promotes IL-10-dependent metabolic defenses to protect from sepsis-induced lethality. Science advances, 11(7), eadu4034.

Wang HJ, et al. (2025) BCL6 coordinates muscle mass homeostasis with nutritional states. Proceedings of the National Academy of Sciences of the United States of America, 122(4), e2408896122.

Gatti P, et al. (2025) Mitochondria- and ER-associated actin are required for mitochondrial fusion. Nature communications, 16(1), 451.

Huang P, et al. (2025) Clinical functional proteomics of intercellular signalling in pancreatic cancer. Nature, 637(8046), 726.

Fan W, et al. (2025) Estrogen-related receptors regulate innate and adaptive muscle mitochondrial energetics through cooperative and distinct actions. Proceedings of the National Academy of Sciences of the United States of America, 122(20), e2426179122.

Wang X, et al. (2025) PIN1 Prolyl Isomerase Promotes Initiation and Progression of Bladder Cancer through the SREBP2-Mediated Cholesterol Biosynthesis Pathway. *Cancer discovery*, 15(3), 633.

Agrawal A, et al. (2025) Identification of microproteins with transactivation activity by polyalanine motif selection. *RSC chemical biology*, 6(5), 800.

Lorenzini MH, et al. (2025) Joint single-cell profiling of Cas9 edits and transcriptomes reveals widespread off-target events and effects on gene expression. *bioRxiv : the preprint server for biology*.

Xu Z, et al. (2025) Scavenger Receptor CD36 in Tumor-Associated Macrophages Promotes Cancer Progression by Dampening Type-I IFN Signaling. *Cancer research*, 85(3), 462.

Siguenza N, et al. (2025) Gut Competition Dynamics of Live Bacterial Therapeutics Are Shaped by Microbiome Complexity, Diet, and Therapeutic Transgenes. *bioRxiv : the preprint server for biology*.

Bosworth AP, et al. (2025) Astrocyte glypican 5 regulates synapse maturation and stabilization. *Cell reports*, 44(3), 115374.

Roos M, et al. (2025) Optimizing mouse metatranscriptome profiling by selective removal of redundant nucleic acid sequences. *bioRxiv : the preprint server for biology*.

Zhang J, et al. (2024) Live bacterial therapeutics for detection and treatment of colorectal cancer. *Nature reviews. Gastroenterology & hepatology*, 21(5), 295.

Newman LE, et al. (2024) Mitochondrial DNA replication stress triggers a pro-inflammatory endosomal pathway of nucleoid disposal. *Nature cell biology*, 26(2), 194.

Mendiratta G, et al. (2024) Cancer research is not correlated with driver gene mutation burdens. *Med (New York, N.Y.)*, 5(7), 832.

Buswinka CJ, et al. (2024) Large-scale annotated dataset for cochlear hair cell detection and classification. *Scientific data*, 11(1), 416.