

Resource Summary Report

Generated by SPARC SAWG on Sep 19, 2026

University of California at Irvine Genomics Research and Technology Hub Core Facility

RRID:SCR_026615

Type: Tool

Proper Citation

University of California at Irvine Genomics Research and Technology Hub Core Facility
(RRID:SCR_026615)

Resource Information

URL: <https://genomics.uci.edu/about-us/>

Proper Citation: University of California at Irvine Genomics Research and Technology Hub Core Facility (RRID:SCR_026615)

Description: Core provides services ranging from DNA/RNA QC analysis, library construction and sequencing on various platforms. Offers several sequencing platforms from Illumina's MiSeq and NovaSeq to PacBio's Sequel II System.

Synonyms: , GRT Hub, University of California at Irvine-Genomics Research and Technology Hub, UCI GRT Hub, UCI Genomics Research and Technology Hub

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

Keywords: ABRF, DNA/RNA QC analysis, library construction, sequencing,

Resource Name: University of California at Irvine Genomics Research and Technology Hub Core Facility

Resource ID: SCR_026615

Record Creation Time: 20250322T060326+0000

Record Last Update: 20260919T200053+0000

Ratings and Alerts

No rating or validation information has been found for University of California at Irvine Genomics Research and Technology Hub Core Facility.

No alerts have been found for University of California at Irvine Genomics Research and Technology Hub Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Subramanian A, et al. (2026) Analysis of cranial tenocyte heterogeneity reveals a role for Wnt signaling in tendon attachments. *Development (Cambridge, England)*, 153(2).

Hudson C, et al. (2026) Human neural stem cell-derived extracellular vesicles improve cognitive function following glioma chemoradiation therapy. *Cancer letters*, 653, 218564.

Pirbhoy PS, et al. (2026) A Workflow for Spatial Transcriptomic Analysis from Intra-operative Human Skeletal Muscle Biopsies. *bioRxiv : the preprint server for biology*.

Nayak PK, et al. (2025) Transcriptome profiling of tendon fibroblasts at the onset of embryonic muscle contraction reveals novel force-responsive genes. *eLife*, 14.