

# Resource Summary Report

Generated by [dkNET](#) on Aug 17, 2026

## University of California, Berkeley Biosciences Divisional Services Cell Culture Core Facility

RRID:SCR\_017924

Type: Tool

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### Proper Citation

University of California, Berkeley Biosciences Divisional Services Cell Culture Core Facility (RRID:SCR\_017924)

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### Resource Information

**URL:** <https://bds.berkeley.edu/facilities/cell-culture>

**Proper Citation:** University of California, Berkeley Biosciences Divisional Services Cell Culture Core Facility (RRID:SCR\_017924)

**Description:** Core provides cell cultures (mammalian and insect), media, supplies and expertise and consultation regarding cell culture needs, media, and experimental design. Provides tested media, serum, and cell lines at exceptionally low prices, large scale production of cells, expansion and harvesting of hybridoma lines, cryopreservation of cell lines, Cell Authentication by STR analysis provided in conjunction with DNA Sequencing Facility, Mycoplasma screening, Use of 2 laminar flow hoods for mammalian or insect cell manipulation.

**Synonyms:** Cell Culture Facility

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

**Keywords:** Cell, culture, mammalian, insect, media, supplies, expertise, hybridoma, line, cryopreservation, authentication, DNA, sequencing, mycoplasma, screening, , service, core, ABRF

**Availability:** Open

**Resource Name:** University of California, Berkeley Biosciences Divisional Services Cell Culture Core Facility

**Resource ID:** SCR\_017924

**Alternate IDs:** ABRF\_816

**Record Creation Time:** 20220129T080337+0000

## Ratings and Alerts

No rating or validation information has been found for University of California, Berkeley Biosciences Divisional Services Cell Culture Core Facility.

No alerts have been found for University of California, Berkeley Biosciences Divisional Services Cell Culture Core Facility.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 17 mentions in open access literature.

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

Heider M, et al. (2026) Localized heme sensing through a ternary molecular glue. bioRxiv : the preprint server for biology.

Gonzalez-Valero A, et al. (2026) Chiral methionine oxidation reagents reveal stereospecific proteome modifications. bioRxiv : the preprint server for biology.

Elleman AV, et al. (2026) Tension TRAAKer: a chemigenetic fluorescent membrane tension reporter. bioRxiv : the preprint server for biology.

Witus SR, et al. (2026) Metabolite glues as a means of purine sensing and chemotherapeutic response. Nature, 655(8125), 1300.

Arkinson C, et al. (2025) NUB1 traps unfolded FAT10 for ubiquitin-independent degradation by the 26S proteasome. Nature structural & molecular biology, 32(9), 1752.

Arkinson C, et al. (2025) Structural landscape of the degrading 26S proteasome reveals conformation-specific binding of TXNL1. Nature structural & molecular biology, 32(12), 2403.

Voke E, et al. (2025) Protein corona formed on lipid nanoparticles compromises delivery efficiency of mRNA cargo. Nature communications, 16(1), 8699.

Arkinson C, et al. (2024) Nub1 traps unfolded FAT10 for ubiquitin-independent degradation by the 26S proteasome. bioRxiv : the preprint server for biology.

Arkinson C, et al. (2024) Structural landscape of AAA+ ATPase motor states in the substrate-degrading human 26S proteasome reveals conformation-specific binding of TXNL1. bioRxiv : the preprint server for biology.

Lee RJ, et al. (2024) Conserved and divergent DNA recognition specificities and functions of R2 retrotransposon N-terminal domains. *Cell reports*, 43(5), 114239.

Morgens DW, et al. (2023) Coding and non-coding elements comprise a regulatory network controlling transcription in Kaposi's sarcoma-associated herpesvirus. *bioRxiv : the preprint server for biology*.

Mark KG, et al. (2023) Orphan quality control shapes network dynamics and gene expression. *Cell*, 186(16), 3460.

Chen Y, et al. (2022) Mechanisms governing target search and binding dynamics of hypoxia-inducible factors. *eLife*, 11.

Hamilton JR, et al. (2021) Targeted delivery of CRISPR-Cas9 and transgenes enables complex immune cell engineering. *Cell reports*, 35(9), 109207.

Tsu BV, et al. (2021) Diverse viral proteases activate the NLRP1 inflammasome. *eLife*, 10.

Suter EC, et al. (2021) Antibody:CD47 ratio regulates macrophage phagocytosis through competitive receptor phosphorylation. *Cell reports*, 36(8), 109587.

Belardi B, et al. (2020) A Weak Link with Actin Organizes Tight Junctions to Control Epithelial Permeability. *Developmental cell*, 54(6), 792.