

# Resource Summary Report

Generated by Kravitz on Sep 10, 2026

## Stanford University Community of Shared Advanced Research Platforms

RRID:SCR\_022986

Type: Tool

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### Proper Citation

Stanford University Community of Shared Advanced Research Platforms  
(RRID:SCR\_022986)

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### Resource Information

**URL:** <https://csharp.stanford.edu/>

**Proper Citation:** Stanford University Community of Shared Advanced Research Platforms  
(RRID:SCR\_022986)

**Description:** Center coordinates ideas and investments to advance and democratize access to Stanford cores, instrumentation, scientific and technical staff, allowing users to perform innovative, rigorous experiments.

**Abbreviations:** C-ShARP

**Synonyms:** Community of Shared Advanced Research Platforms

**Resource Type:** data or information resource, organization portal, portal

**Keywords:** USEDit, ABRF, access to Stanford cores, access to Stanford instrumentation, access to Stanford scientific and technical staff

**Availability:** Free, Freely available

**Resource Name:** Stanford University Community of Shared Advanced Research Platforms

**Resource ID:** SCR\_022986

**Alternate IDs:** ABRF\_2461

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=2461&citation=1>

**Record Creation Time:** 20221119T050156+0000

**Record Last Update:** 20260905T133101+0000

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## Ratings and Alerts

No rating or validation information has been found for Stanford University Community of Shared Advanced Research Platforms.

No alerts have been found for Stanford University Community of Shared Advanced Research Platforms.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 13 mentions in open access literature.

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

Zakrzewska S, et al. (2026) Saxiphilin is a broad-spectrum toxin sponge for C13-modified saxitoxins. bioRxiv : the preprint server for biology.

Liu Q, et al. (2026) Structural basis for selective thymidine binding by the *Borrelia burgdorferi* substrate-binding protein BmpA. The Journal of biological chemistry, 302(7), 113206.

Liu Q, et al. (2026) Structure and ligand specificity of *Borrelia burgdorferi* BmpA. bioRxiv : the preprint server for biology.

Zakrzewska S, et al. (2026) Saxiphilin is a broad-spectrum toxin sponge for C13-modified saxitoxins. Structure (London, England : 1993).

Feist WN, et al. (2025) Multilayered HIV-1 resistance in HSPCs through CCR5 Knockout and B cell secretion of HIV-inhibiting antibodies. Nature communications, 16(1), 3103.

Franklin JM, et al. (2025) Human Satellite 3 DNA encodes megabase-scale transcription factor binding platforms. bioRxiv : the preprint server for biology.

Gudinas AP, et al. (2025) Ion-selective conformational stabilization of a disordered repeats-in-toxin protein domain. Biophysical journal, 124(23), 4243.

Park ER, et al. (2025) A chemogenetic ligand-receptor pair for voltage-gated sodium channel subtype-selective inhibition. bioRxiv : the preprint server for biology.

Yuan L, et al. (2025) Defining the Antitumor Mechanism of Action of a Clinical-stage Compound as a Selective Degradator of the Nuclear Pore Complex. Cancer discovery, 15(12), 2505.

Gao Y, et al. (2025) Convenient syntheses of isotopically labeled pyrimidine 2'-deoxynucleosides and their 5-hydroxy oxidation products. Nucleosides, nucleotides & nucleic acids, 44(11), 915.

Hansen ME, et al. (2025) Unidirectional Chain Transfer of Enol Ethers Yields Heterotelechelic Degradable Poly(Enol Ether). ACS macro letters, 14(10), 1492.

Chosy MB, et al. (2024) Vancomycin-Polyguanidino Dendrimer Conjugates Inhibit Growth of Antibiotic-Resistant Gram-Positive and Gram-Negative Bacteria and Eradicate Biofilm-Associated *S. aureus*. ACS infectious diseases, 10(2), 384.

Horst M, et al. (2024) Fluorination Affects the Force Sensitivity and Nonequilibrium Dynamics of the Mechanochemical Unzipping of Ladderanes. Journal of the American Chemical Society, 146(47), 32651.