

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## Leginon

RRID:SCR\_016731

Type: Tool

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### Proper Citation

Leginon (RRID:SCR\_016731)

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### Resource Information

**URL:** [http://emg.nysbc.org/redmine/projects/leginon/wiki/Leginon\\_Homepage](http://emg.nysbc.org/redmine/projects/leginon/wiki/Leginon_Homepage)

**Proper Citation:** Leginon (RRID:SCR\_016731)

**Description:** System designed for automated collection of images from a transmission electron microscope.

**Resource Type:** data acquisition software, data or information resource, data processing software, data repository, image acquisition software, portal, service resource, software application, software resource, storage service resource

**Defining Citation:** [PMID:15890530](#)

**Keywords:** automated, collection, acquisition, data, image, electron, microscope

**Funding:** NCRR RR17573;  
NIGMS GM61939;  
NSF DBI0352386;  
NSF DBI9730056;  
NSF DBI9904547

**Availability:** Free, Available for download, Freely available, Registration suggested

**Resource Name:** Leginon

**Resource ID:** SCR\_016731

**License:** Apache License, Version 2.0

**Record Creation Time:** 20220129T080332+0000

**Record Last Update:** 20260905T132813+0000

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### Ratings and Alerts

No rating or validation information has been found for Leginon.

No alerts have been found for Leginon.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 45 mentions in open access literature.

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

Monteiro AC, et al. (2026) OPG-Producing B Cells and RANKL-Expressing T Cells Define Immune Signatures Predictive of Bone Metastases in Breast Cancer. Cancer research communications, 6(1), 85.

Bader DLV, et al. (2026) Structural and immunogenetic signatures guide CD4-mimetic HIV vaccine development. Cell reports, 45(4), 117180.

Li SH, et al. (2026) Childhood immunological imprinting of cross-subtype antibodies targeting the hemagglutinin head domain of influenza viruses. Cell host & microbe, 34(5), 873.

Abiona OM, et al. (2025) MERS-CoV spike vaccine-induced N-terminal domain-specific antibodies are more protective than receptor binding domain-specific antibodies. iScience, 28(2), 111632.

Schleich FA, et al. (2025) Vaccination of nonhuman primates elicits a broadly neutralizing antibody lineage targeting a quaternary epitope on the HIV-1 Env trimer. Immunity, 58(6), 1598.

Dederer V, et al. (2024) A designed ankyrin-repeat protein that targets Parkinson's disease-associated LRRK2. The Journal of biological chemistry, 300(7), 107469.

Reimer JM, et al. (2023) Structure of LRRK1 and mechanisms of autoinhibition and activation. Nature structural & molecular biology, 30(11), 1735.

Tan ZY, et al. (2023) Heterogeneous non-canonical nucleosomes predominate in yeast cells in situ. eLife, 12.

Huang P, et al. (2022) Structural basis for catalyzed assembly of the Sonic hedgehog-Patched1 signaling complex. Developmental cell, 57(5), 670.

Van Tilbeurgh M, et al. (2022) Innate cell markers that predict anti-HIV neutralizing antibody titers in vaccinated macaques. Cell reports. Medicine, 3(10), 100751.

Snead DM, et al. (2022) Structural basis for Parkinson's disease-linked LRRK2's binding to microtubules. Nature structural & molecular biology, 29(12), 1196.

Melville Z, et al. (2022) High-resolution structure of the membrane-embedded skeletal muscle ryanodine receptor. *Structure (London, England : 1993)*, 30(1), 172.

Puno MR, et al. (2022) Structural basis for RNA surveillance by the human nuclear exosome targeting (NEXT) complex. *Cell*, 185(12), 2132.

Sahoo A, et al. (2022) Structure-guided changes at the V2 apex of HIV-1 clade C trimer enhance elicitation of autologous neutralizing and broad V1V2-scaffold antibodies. *Cell reports*, 38(9), 110436.

Kirchdoerfer RN, et al. (2021) Structure and immune recognition of the porcine epidemic diarrhea virus spike protein. *Structure (London, England : 1993)*, 29(4), 385.

Jiang D, et al. (2021) Open-state structure and pore gating mechanism of the cardiac sodium channel. *Cell*, 184(20), 5151.

Wisedchaisri G, et al. (2021) Structural Basis for High-Affinity Trapping of the NaV1.7 Channel in Its Resting State by Tarantula Toxin. *Molecular cell*, 81(1), 38.

Nikolay R, et al. (2021) Snapshots of native pre-50S ribosomes reveal a biogenesis factor network and evolutionary specialization. *Molecular cell*, 81(6), 1200.

Wang C, et al. (2020) Structures reveal gatekeeping of the mitochondrial Ca<sup>2+</sup> uniporter by MICU1-MICU2. *eLife*, 9.

Walls AC, et al. (2020) Structure, Function, and Antigenicity of the SARS-CoV-2 Spike Glycoprotein. *Cell*, 181(2), 281.