Resource Summary Report

Generated by dkNET on Sep 4, 2026

Napari

RRID:SCR_022765
Type: Tool

Proper Citation

Napari (RRID:SCR_022765)

Resource Information

URL: https://napari.org/

Proper Citation: Napari (RRID:SCR_022765)

Description: Multi dimensional image viewer for Python. Used for browsing, annotating, and analyzing large multi dimensional images. Can be coupled to machine learning and image analysis tools enabling more user friendly automated analysis.

Resource Type: data access protocol, software resource, web service

Keywords: Multi dimensional image viewer, large multi dimensional images, automated analysis

Availability: Free, Available for download, Freely available

Resource Name: Napari

Resource ID: SCR_022765

Alternate URLs: https://github.com/napari/napari, https://zenodo.org/record/6598542#.YykVuHbMK3A

License: BSD 3-Clause "New" or "Revised" License

Record Creation Time: 20220920T050146+0000

Record Last Update: 20260903T235854+0000

Ratings and Alerts

No rating or validation information has been found for Napari.

No alerts have been found for Napari.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 46 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Jung F, et al. (2026) DMC-BrainMap is an open-source, end-to-end tool for multi-feature brain mapping in different species. Cell reports methods, 6(2), 101302.

Naert T, et al. (2026) Precise, predictable genome integrations by deep-learning-assisted design of microhomology-based templates. Nature biotechnology, 44(6), 1023.

Park CJ, et al. (2026) Validating DevCheck: a semi-automated computer algorithm for measuring foveal immaturity in premature infants. Quantitative imaging in medicine and surgery, 16(1), 5.

Wafer R, et al. (2026) A quantitative in vivo CRISPR-imaging platform identifies regulators of hyperplastic and hypertrophic adipose morphology in zebrafish. eLife, 14.

Pérez-Verdugo F, et al. (2026) Mechanosensitive feedback organizes cell shape and motion during hindbrain neuropore morphogenesis. Current biology : CB, 36(8), 1903.

Pileggi CA, et al. (2026) Functional and metabolomic analyses of brown adipose tissue during cold-deacclimation reveal rapid N-acetylated amino acid adaptations. iScience, 29(4), 115146.

Cutts EE, et al. (2025) Molecular mechanism of condensin I activation by KIF4A. The EMBO journal, 44(3), 682.

Marcotti S, et al. (2025) Enhancing Research Through Image Analysis Workshops: Experiences and Best Practices. Microscopy research and technique, 88(4), 922.

Pylvänäinen JW, et al. (2025) Practical considerations for data exploration in quantitative cell biology. Journal of cell science, 138(7).

Ahmad S, et al. (2025) Visualizing the transcription and replication of influenza A viral RNAs in cells by multiple direct RNA padlock probing and in situ sequencing (mudRapp-seq). Nucleic acids research, 53(11).

Sendra M, et al. (2025) Myocardium and endocardium of the early mammalian heart tube arise from independent multipotent lineages specified at the primitive streak. Developmental cell.

Coughlin GM, et al. (2025) Spatial genomics of AAV vectors reveals mechanism of transcriptional crosstalk that enables targeted delivery of large genetic cargo. Nature biotechnology.

Nicolas WJ, et al. (2025) Structure of the lens MP20 mediated adhesive junction. Nature communications, 16(1), 2977.

Glaser A, et al. (2025) Expansion-assisted selective plane illumination microscopy for nanoscale imaging of centimeter-scale tissues. eLife, 12.

Autio M, et al. (2025) Spatial Analysis of the Tumor Microenvironment in Diffuse Large B-cell Lymphoma Reveals Clinically Relevant Cell Interactions and Recurrent Cellular Neighborhoods. Cancer immunology research, 13(10), 1674.

Pollington HQ, et al. (2025) The Hunchback transcription factor determines interneuron molecular identity, morphology, and presynapse targeting in the Drosophila NB5-2 lineage. PLoS biology, 23(3), e3002881.

Ullrich S, et al. (2025) Co-transcriptional splicing is delayed in the highly expressed thyroglobulin gene. Journal of cell science, 138(6).

Vanspauwen SK, et al. (2025) A correlation-based tool for quantifying membrane periodic skeleton associated periodicity. Frontiers in neuroinformatics, 19, 1628538.

Randel N, et al. (2025) Effects of food availability on larval development during ontogenetic niche shift in a marine annelid. Open biology, 15(10), 250135.

Cramer EM, et al. (2025) Temporal reassignment and correspondence evaluation with quality control for time-course imaging of 3D cell culture. Cell reports methods, 5(12), 101237.