

Resource Summary Report

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

Papaya

RRID:SCR_014188

Type: Tool

Proper Citation

Papaya (RRID:SCR_014188)

Resource Information

URL: <http://www.nitrc.org/projects/papaya>

Proper Citation: Papaya (RRID:SCR_014188)

Description: A pure JavaScript medical research image viewer, compatible across a range of popular web browsers. The orthogonal viewer supports NIFTI and DICOM files, overlays and atlas labels. It requires Firefox (7+), Chrome (7+), Safari (6+), MobileSafari (iOS 6+), or IE (10+).

Resource Type: software resource, source code

Keywords: source code, javascript, medical research, image viewer

Availability: Public, Available to the research community

Resource Name: Papaya

Resource ID: SCR_014188

Alternate URLs: <https://github.com/rii-mango/Papaya/>

License: BSD License

Record Creation Time: 20220129T080319+0000

Record Last Update: 20260905T133359+0000

Ratings and Alerts

No rating or validation information has been found for Papaya.

No alerts have been found for Papaya.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

G??ttler C, et al. (2025) Toward a standard preoperative MRI protocol for functional neurosurgery. Imaging neuroscience (Cambridge, Mass.), 3.

Alaqra AS, et al. (2021) Machine Learning-Based Analysis of Encrypted Medical Data in the Cloud: Qualitative Study of Expert Stakeholders' Perspectives. JMIR human factors, 8(3), e21810.

Wilts BD, et al. (2017) Butterfly gyroid nanostructures as a time-frozen glimpse of intracellular membrane development. Science advances, 3(4), e1603119.

Berghoff BA, et al. (2017) RNA-sequence data normalization through in silico prediction of reference genes: the bacterial response to DNA damage as case study. BioData mining, 10, 30.

Janevski A, et al. (2009) PAPAYa: a platform for breast cancer biomarker signature discovery, evaluation and assessment. BMC bioinformatics, 10 Suppl 9(Suppl 9), S7.