

Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

Harvard Medical School, Department of Systems Biology: The Megason Lab - MegaCapture Software

RRID:SCR_001732

Type: Tool

Proper Citation

Harvard Medical School, Department of Systems Biology: The Megason Lab - MegaCapture Software (RRID:SCR_001732)

Resource Information

URL: <https://wiki.med.harvard.edu/SysBio/Megason/MegaCapture>

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Description: MegaCapture is a VisualBasic macro developed for automating the acquisition of in toto image sets. It is useable on Zeiss microscopes using LSM v4.x software (pre-Zen). MegaCapture can automatically acquire image sets across any combination of dimensions including x, y, z, time, color, x-tile, y-tile, row, and column. Images are exported on the fly and can be compressed allowing very large (100,000 images) image sets to be captured. Sponsors: MegaCapture is supported by Harvard University.

Synonyms: MegaCapture

Resource Type: data visualization software, data or information resource, portal, topical portal, data acquisition software, software application, data processing software, software resource

Keywords: fly, acquisition, automating, dimention, macro, microscope, software, toto image, image

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Harvard Medical School, Department of Systems Biology: The Megason Lab - MegaCapture Software

Resource ID: SCR_001732

Alternate IDs: nif-0000-10242

Record Creation Time: 20220129T080209+0000

Record Last Update: 20260801T190148+0000

Ratings and Alerts

No rating or validation information has been found for Harvard Medical School, Department of Systems Biology: The Megason Lab - MegaCapture Software.

No alerts have been found for Harvard Medical School, Department of Systems Biology: The Megason Lab - MegaCapture Software.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.