

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

PAM

RRID:SCR_020966

Type: Tool

Proper Citation

PAM (RRID:SCR_020966)

Resource Information

URL: <https://gitlab.com/PAM-PIE/PAM>

Proper Citation: PAM (RRID:SCR_020966)

Description: Software package for quantitative analysis of fluorescence microscopy and spectroscopy data, with focus on experiments using pulsed interleaved excitation. Open source software package written in MATLAB that offers workflow through its graphical user interface. Framework for integrated analysis of imaging, single-molecule, and ensemble fluorescence data. Supports most types of data collection modalities.

Synonyms: Pulsed interleaved excitation Analysis with Matlab, Pulsed Interleaved Excitation analysis with MATLAB, PIE analysis with MATLAB

Resource Type: data analysis software, data processing software, software application, software resource

Defining Citation: [PMID:29642023](#)

Keywords: Quantitative data analysis, fluorescence microscopy data, data analysis, spectroscopy data analysis, pulsed interleaved excitation, imaging integrated analysis

Funding: Deutsche Forschungsgemeinschaft ;
Ludwig-Maximilians-University; Munich; Germany ;
Research Foundation Flanders

Availability: Free, Available for download, Freely available

Resource Name: PAM

Resource ID: SCR_020966

Alternate URLs: <http://www.cup.uni-muenchen.de/pc/lamb/software/pam.html>

License: GNU General Public License v3.0

Record Creation Time: 20220129T080353+0000

Record Last Update: 20260912T195949+0000

Ratings and Alerts

No rating or validation information has been found for PAM.

No alerts have been found for PAM.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Fenn KL, et al. (2024) Outer membrane protein assembly mediated by BAM-SurA complexes. Nature communications, 15(1), 7612.

Coucke Q, et al. (2023) Particle-based phasor-FLIM-FRET resolves protein-protein interactions inside single viral particles. Biophysical reports, 3(3), 100122.

Sefer A, et al. (2022) Structural dynamics of DNA strand break sensing by PARP-1 at a single-molecule level. Nature communications, 13(1), 6569.

Lerner E, et al. (2021) FRET-based dynamic structural biology: Challenges, perspectives and an appeal for open-science practices. eLife, 10.