

Resource Summary Report

Generated by SPARC SAWG on Sep 16, 2026

FBIRN Image Processing Scripts

RRID:SCR_009471

Type: Tool

Proper Citation

FBIRN Image Processing Scripts (RRID:SCR_009471)

Resource Information

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

Proper Citation: FBIRN Image Processing Scripts (RRID:SCR_009471)

Description: A FSL package for the comprehensive management of large-scale multi-site fMRI projects, including data storage, retrieval, calibration, analysis, multi-modal integration, and quality control.

Abbreviations: FIPS

Synonyms: FBIRN Image Processing Scripts (FIPS)

Resource Type: data management software, software application, software resource

Keywords: analyze, csh/tcsh, database application, dicom, linux, macos, magnetic resonance, posix/unix-like, statistical operation, unix shell, web environment

Availability: BIRN License, BSD License

Resource Name: FBIRN Image Processing Scripts

Resource ID: SCR_009471

Alternate IDs: nlx_155618

Record Creation Time: 20220129T080253+0000

Record Last Update: 20260912T195712+0000

Ratings and Alerts

No rating or validation information has been found for FBIRN Image Processing Scripts.

No alerts have been found for FBIRN Image Processing Scripts.

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 [SPARC SAWG](#) .

Amaro TMMM, et al. (2018) Random mutagenesis screen shows that *Phytophthora capsici* CRN83_152-mediated cell death is not required for its virulence function(s). *Molecular plant pathology*, 19(5), 1114.

Wilson JJ, et al. (2018) Three-Dimensional Representation of Motor Space in the Mouse Superior Colliculus. *Current biology : CB*, 28(11), 1744.

Jamei M, et al. (2013) The simcyp population based simulator: architecture, implementation, and quality assurance. *In silico pharmacology*, 1, 9.

O'Farrell B, et al. (2012) Transforming microbial genotyping: a robotic pipeline for genotyping bacterial strains. *PloS one*, 7(10), e48022.

Turner JA, et al. (2010) Application of neuroanatomical ontologies for neuroimaging data annotation. *Frontiers in neuroinformatics*, 4.