

Resource Summary Report

Generated by RRID on Aug 15, 2026

SCIPI

RRID:SCR_027939

Type: Tool

Proper Citation

SCIPI (RRID:SCR_027939)

Resource Information

URL: https://github.com/JeffLuo9/Hep_Ploidy_protocol/

Proper Citation: SCIPI (RRID:SCR_027939)

Description: Software hepatocyte ploidy identification pipeline on the stereo-cell platform for accurate liver polyploidy classification via fluorescence staining and deep learning tools.

Resource Type: software toolkit, source code, software resource

Keywords: hepatocyte ploidy identification, liver polyploidy classification, fluorescence staining, deep learning tools,

Availability: Free, Available for download, Freely available

Resource Name: SCIPI

Resource ID: SCR_027939

License: MIT license

Record Creation Time: 20260206T055600+0000

Record Last Update: 20260815T183314+0000

Ratings and Alerts

No rating or validation information has been found for SCIPI.

No alerts have been found for SCIPI.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Yang Y, et al. (2026) Stereo-cell deciphers the spatial and functional heterogeneity of polyploid hepatocytes. GigaScience, 15.

Berta J, et al. (2026) Catching the Mardi Gras fever: Quantifying the impact of mass gathering tourism on local bacterial prevalence and community diversity in municipal wastewater. Research square.