

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

syndRomics

RRID:SCR_019312

Type: Tool

Proper Citation

syndRomics (RRID:SCR_019312)

Resource Information

URL: <https://github.com/ucsf-ferguson-lab/syndRomics>

Proper Citation: syndRomics (RRID:SCR_019312)

Description: Software R package implements functions for helping in process of disease patterns analysis by means of principal components. These include component visualization, interpretation and stability analysis.

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

Keywords: disease patterns analysis, principal components, component visualization, component interpretation, component stability analysis

Availability: Free, Available for download, Freely available

Resource Name: syndRomics

Resource ID: SCR_019312

License: MIT License

Record Creation Time: 20220129T080344+0000

Record Last Update: 20260905T132848+0000

Ratings and Alerts

No rating or validation information has been found for syndRomics.

No alerts have been found for syndRomics.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Torres-Esp??n A, et al. (2021) Reproducible analysis of disease space via principal components using the novel R package syndRomics. eLife, 10.