# **Resource Summary Report**

Generated by NIF on May 4, 2025

# **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 processing software, software resource, data analysis software, software application

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

**Funding:** 

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:** 20250503T060843+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 NIF.

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