

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

[hypeR](#)

RRID:SCR_027229

Type: Tool

Proper Citation

hypeR (RRID:SCR_027229)

Resource Information

URL: <https://montilab.github.io/hypeR-docs/>

Proper Citation: hypeR (RRID:SCR_027229)

Description: Software R package that supports different types of gene set enrichment workflows. Offers multiple enrichment, visualization, and sharing methods in addition to novel features such as hierarchical geneset analysis and built-in markdown reporting.

Resource Type: software resource, software toolkit, source code

Defining Citation: [PMID:31498385](#)

Keywords: gene set enrichment workflows, hierarchical geneset analysis, built-in markdown reporting,

Funding: NIA 2U19AG023122;
William M. Wood Foundation

Availability: Free, Available for download, Freely available

Resource Name: hypeR

Resource ID: SCR_027229

License: GNU GPLv3

Record Creation Time: 20250723T062011+0000

Record Last Update: 20260805T044610+0000

Ratings and Alerts

No rating or validation information has been found for hypeR.

No alerts have been found for hypeR.

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 [RRID](#) .

Dado T, et al. (2022) Hyperrealistic neural decoding for reconstructing faces from fMRI activations via the GAN latent space. Scientific reports, 12(1), 141.