

Resource Summary Report

Generated by [RRID](#) on Aug 11, 2026

ActivePathways

RRID:SCR_027736

Type: Tool

Proper Citation

ActivePathways (RRID:SCR_027736)

Resource Information

URL: <https://cran.r-project.org/web/packages/ActivePathways/index.html>

Proper Citation: ActivePathways (RRID:SCR_027736)

Description: Software R package for analysing multiple omics datasets in the context of molecular pathways, biological processes and other types of gene sets. Method that first prioritises genes through multi-omics data fusion and then identifies enriched pathways with gene-level evidence from input datasets.

Resource Type: software toolkit, software resource

Defining Citation: [DOI:10.1038/s41467-024-49986-4](https://doi.org/10.1038/s41467-024-49986-4)

Keywords: Pathway, genes, analysing multiple omics datasets, molecular pathways, biological processes, gene sets,

Availability: Free, Available for download, Freely available

Resource Name: ActivePathways

Resource ID: SCR_027736

License: GNU GPL v3

Record Creation Time: 20251209T055453+0000

Record Last Update: 20260810T040930+0000

Ratings and Alerts

No rating or validation information has been found for ActivePathways.

No alerts have been found for ActivePathways.

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](#) .

Culliford R, et al. (2026) Contrasting Features of Papillary and Chromophobe Renal Cell Carcinoma Revealed by Whole-Genome Sequencing. Molecular cancer research : MCR, 24(5), 327.