

Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

SigFuge

RRID:SCR_000444

Type: Tool

Proper Citation

SigFuge (RRID:SCR_000444)

Resource Information

URL: <http://www.bioconductor.org/packages/release/bioc/html/SigFuge.html>

Proper Citation: SigFuge (RRID:SCR_000444)

Description: Algorithm for testing significance of clustering in RNA-seq data.

Resource Type: software resource

Defining Citation: [PMID:25030904](#)

Keywords: software package, unix/linux, mac os x, windows, r, clustering, rna-seq, visualization

Availability: Free, Available for download, Freely available

Resource Name: SigFuge

Resource ID: SCR_000444

Alternate IDs: OMICS_05201

License: GNU General Public License, v3

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260725T190446+0000

Ratings and Alerts

No rating or validation information has been found for SigFuge.

No alerts have been found for SigFuge.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.