

# Resource Summary Report

Generated by T1D on Aug 9, 2026

## SigFuge

RRID:SCR\_000444

Type: Tool

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### Proper Citation

SigFuge (RRID:SCR\_000444)

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### 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:** 20260808T185716+0000

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### Ratings and Alerts

No rating or validation information has been found for SigFuge.

No alerts have been found for SigFuge.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.