

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## [rgsepd](#)

RRID:SCR\_024256

Type: Tool

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### Proper Citation

rgsepd (RRID:SCR\_024256)

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### Resource Information

**URL:** <https://bioconductor.org/packages/release/bioc/html/rgsepd.html>

**Proper Citation:** rgsepd (RRID:SCR\_024256)

**Description:** Software R package to help disambiguate transcriptome samples by automating differential expression, then gene set enrichment, and finally N-dimensional projection to quantify in which ways each sample is like either treatment group.

**Resource Type:** software toolkit, software resource

**Keywords:** disambiguate transcriptome samples, automating differential expression, gene set enrichment, N-dimensional projection, treatment group,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** rgsepd

**Resource ID:** SCR\_024256

**Alternate URLs:** <https://sources.debian.org/src/r-bioc-rgsepd/>

**Record Creation Time:** 20230830T050217+0000

**Record Last Update:** 20260729T044607+0000

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### Ratings and Alerts

No rating or validation information has been found for rgsepd.

No alerts have been found for rgsepd.

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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.