

# Resource Summary Report

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## regioneReloaded

RRID:SCR\_028252

Type: Tool

### Proper Citation

regioneReloaded (RRID:SCR\_028252)

### Resource Information

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

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**Description:** Software package that allows simultaneous analysis of associations between genomic region sets, enabling clustering of data and creation of graphs. Incorporates strategy to improve p-value calculations and normalize z-scores coming from multiple analysis to allow for their direct comparison. Builds upon regioneR by adding new plotting functions for obtaining publication-ready graphs.

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

**Defining Citation:** [PMID:37988135](#)

**Keywords:** analysis of associations between genomic region sets, data clustering, graphs creation, p-value calculations, normalize z-scores, multiple analysis direct comparison,

**Funding:** European Regional Development Fund ;  
Marie Skłodowska Curie Training network ;  
Fundació Internacional Josep Carreras

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

**Resource Name:** regioneReloaded

**Resource ID:** SCR\_028252

**Alternate URLs:** <https://github.com/RMalinverni/regioneReloaded>

**License:** Artistic-2.0

**Record Creation Time:** 20260404T065920+0000

**Record Last Update:** 20260810T040953+0000

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## Ratings and Alerts

No rating or validation information has been found for regioneReloaded.

No alerts have been found for regioneReloaded.

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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.