

Resource Summary Report

Generated by SPARC SAWG on Jul 28, 2026

robustrankaggreg

RRID:SCR_024299

Type: Tool

Proper Citation

robustrankaggreg (RRID:SCR_024299)

Resource Information

URL: <https://cran.r-project.org/package=RobustRankAggreg>

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Description: Software R package for aggregating ranked lists, especially lists of genes. RRA method uses a probabilistic model for aggregation that is robust to noise and also facilitates the calculation of significance probabilities for all the elements in the final ranking.

Abbreviations: RRA

Synonyms: Robust Rank Aggregation

Resource Type: software resource, software toolkit

Keywords: aggregating ranked lists, lists of genes aggregation,

Availability: Free, Available for download, Freely available,

Resource Name: robustrankaggreg

Resource ID: SCR_024299

Alternate URLs: <https://sources.debian.org/src/r-cran-robustrankaggreg/>

License: GPL-2

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260725T191111+0000

Ratings and Alerts

No rating or validation information has been found for robustrankaggreg.

No alerts have been found for robustrankaggreg.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Liu X, et al. (2025) Identification of the Hub Gene LDB3 in Stanford Type A Aortic Dissection Based on Comprehensive Bioinformatics Analysis. Journal of cellular and molecular medicine, 29(6), e70471.

Javanmardifard Z, et al. (2024) A comprehensive in silico analysis and experimental validation of miRNAs capable of discriminating between lung adenocarcinoma and squamous cell carcinoma. Frontiers in genetics, 15, 1419099.

Dunn SL, et al. (2016) Gene expression changes in damaged osteoarthritic cartilage identify a signature of non-chondrogenic and mechanical responses. Osteoarthritis and cartilage, 24(8), 1431.