## **Resource Summary Report**

Generated by NIF on Apr 28, 2025

# robustrankaggreg

RRID:SCR\_024299

Type: Tool

### **Proper Citation**

robustrankaggreg (RRID:SCR\_024299)

#### **Resource Information**

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

**Proper Citation:** robustrankaggreg (RRID:SCR\_024299)

**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 toolkit, software resource

**Keywords:** aggregating ranked lists, lists of genes aggregation,

**Funding:** 

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: 20250425T060603+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 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

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.