

# Resource Summary Report

Generated by [NIF](#) on Sep 12, 2026

## pairedBayes

RRID:SCR\_001738

Type: Tool

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### Proper Citation

pairedBayes (RRID:SCR\_001738)

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### Resource Information

**URL:** <http://bioinformatics.med.yale.edu/group/>

**Proper Citation:** pairedBayes (RRID:SCR\_001738)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on April 10th,2023. An R code for Bayesian modeling of paired RNA-seq experiments.

**Abbreviations:** pairedBayes

**Resource Type:** software resource

**Keywords:** r, rna-seq

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE.

**Resource Name:** pairedBayes

**Resource ID:** SCR\_001738

**Alternate IDs:** OMICS\_01958

**Record Creation Time:** 20220129T080209+0000

**Record Last Update:** 20260912T195530+0000

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### Ratings and Alerts

No rating or validation information has been found for pairedBayes.

No alerts have been found for pairedBayes.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Walsh CJ, et al. (2016) Discovering MicroRNA-Regulatory Modules in Multi-Dimensional Cancer Genomic Data: A Survey of Computational Methods. Cancer informatics, 15(Suppl 2), 25.