

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

pairedBayes

RRID:SCR_001738

Type: Tool

Proper Citation

pairedBayes (RRID:SCR_001738)

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: 20260725T190511+0000

Ratings and Alerts

No rating or validation information has been found for pairedBayes.

No alerts have been found for pairedBayes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

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.