

Resource Summary Report

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casper

RRID:SCR_006613

Type: Tool

Proper Citation

casper (RRID:SCR_006613)

Resource Information

URL: <http://bioconductor.org/packages/release/bioc/html/casper.html>

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Description: Software to infer alternative splicing from paired-end RNA-seq data. The model is based on counting paths across exons, rather than pairwise exon connections, and estimates the fragment size and start distributions non-parametrically, which improves estimation precision.

Abbreviations: casper

Synonyms: casper - Characterization of Alternative Splicing based on Paired-End Reads, Characterization of Alternative Splicing based on Paired-End Reads

Resource Type: software resource

Keywords: bio.tools

Availability: GNU General Public License, v2 or greater

Resource Name: casper

Resource ID: SCR_006613

Alternate IDs: biotools:casper, OMICS_01270

Alternate URLs: <https://bio.tools/casper>

Record Creation Time: 20220129T080237+0000

Record Last Update: 20260801T190313+0000

Ratings and Alerts

No rating or validation information has been found for casper.

No alerts have been found for casper.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 149 mentions in open access literature.

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

Song M, et al. (2025) Benchmarking copy number aberrations inference tools using single-cell multi-omics datasets. Briefings in bioinformatics, 26(2).

Banks PW, et al. (2025) Multiple Mini Interviews vs Traditional Interviews: Investigating Racial and Socioeconomic Differences in Interview Processes. Advances in medical education and practice, 16, 157.

Luu QQ, et al. (2025) Therapeutic Potential of Arginine-Loaded Red Blood Cell Nanovesicles Targeting Obese Asthma. Mediators of inflammation, 2025, 8248722.

Xu H, et al. (2025) Medicaid Payments and Racial and Ethnic Disparities in Alzheimer Disease Special Care Units. JAMA network open, 8(8), e2525057.

Ortiz-Ortiz M, et al. (2025) Entomological survey to determine the role of cisterns in the production of Aedes aegypti in the U.S. Virgin Islands. PloS one, 20(9), e0329222.

Lee HK, et al. (2025) Predictors of Treatment Response to Fecal Microbiota Transplantation in Irritable Bowel Syndrome: A Pilot Study. Journal of neurogastroenterology and motility, 31(4), 462.

Mukuria C, et al. (2025) Mapping functions for the PHQ-9 and GAD-7 to generate EQ-5D-3L for economic evaluation. The European journal of health economics : HEPAC : health economics in prevention and care, 26(1), 63.

Hong SH, et al. (2025) Age- and Sex-Specific Gut Microbiota Signatures Associated with Dementia-Related Brain Pathologies: An LEfSe-Based Metagenomic Study. Brain sciences, 15(6).

Zheng Y, et al. (2025) Total whole-arm chromosome losses predict malignancy in human cancer. Proceedings of the National Academy of Sciences of the United States of America, 122(18), e2505385122.

Keesee E, et al. (2025) Rural hospital closures and nursing home outcomes. The Journal of rural health : official journal of the American Rural Health Association and the National Rural Health Care Association, 41(2), e70026.

Schmid KT, et al. (2025) Benchmarking scRNA-seq copy number variation callers. *Nature communications*, 16(1), 8777.

Zhang L, et al. (2025) Murine cell lines with defined mutations model different histological subtypes of epithelial ovarian cancer. *Disease models & mechanisms*, 18(7).

Kim K, et al. (2025) Robust phylogenetic tree-based microbiome association test using repeatedly measured data for composition bias. *BMC bioinformatics*, 26(1), 75.

Kober AKMH, et al. (2025) Exopolysaccharides from camel milk-derived *Limosilactobacillus reuteri* C66: Structural characterization, bioactive and rheological properties for food applications. *Food chemistry: X*, 25, 102164.

Jara-Servin A, et al. (2025) Microbial Communities in Agave Fermentations Vary by Local Biogeographic Regions. *Environmental microbiology reports*, 17(1), e70057.

Shanel G, et al. (2025) Transcriptional programs of cell identity and p53-induced stress responses are associated with distinctive features of spatial genome organization. *Nucleic acids research*, 53(13).

Daida YG, et al. (2025) Differences in Coronary Heart Disease and Stroke Incidence Among Single-Race and Multiracial Asian and Pacific Islander Subgroups in Hawaii and California: A Retrospective Cohort Study. *Journal of the American Heart Association*, 14(9), e039076.

Yalcin K, et al. (2025) Subgroup of meningiomas involving FOS and FOSB gene fusions. *Nature communications*, 16(1), 10532.

Magyari B, et al. (2024) Single-Center Experience with the Balloon-Expandable Myval Transcatheter Aortic Valve System in Patients with Bicuspid Anatomy: Procedural and 30-Day Follow-Up. *Journal of clinical medicine*, 13(2).

Thirimanne HN, et al. (2024) Meningioma transcriptomic landscape demonstrates novel subtypes with regional associated biology and patient outcome. *Cell genomics*, 4(6), 100566.