

# Resource Summary Report

Generated by [RRID](#) on Sep 11, 2026

## BBSeq

RRID:SCR\_011877

Type: Tool

### Proper Citation

BBSeq (RRID:SCR\_011877)

### Resource Information

**URL:** [http://www.bios.unc.edu/research/genomic\\_software/BBSeq/](http://www.bios.unc.edu/research/genomic_software/BBSeq/)

**Proper Citation:** BBSeq (RRID:SCR\_011877)

**Description:** A Powerful and Flexible Approach to the Analysis of RNA Sequence Count Data.

**Abbreviations:** BBSeq

**Resource Type:** software resource

**Keywords:** bio.tools

**Resource Name:** BBSeq

**Resource ID:** SCR\_011877

**Alternate IDs:** biotools:bbseq, OMICS\_01300

**Alternate URLs:** <https://bio.tools/bbseq>

**Record Creation Time:** 20220129T080307+0000

**Record Last Update:** 20260905T132714+0000

### Ratings and Alerts

No rating or validation information has been found for BBSeq.

No alerts have been found for BBSeq.

### Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Mate-Kole EM, et al. (2024) Machine learning-enhanced stochastic uncertainty and sensitivity analysis of the ICRP human respiratory tract model for an inhaled radionuclide. Journal of radiological protection : official journal of the Society for Radiological Protection, 44(4).

Sun S, et al. (2017) Differential expression analysis for RNAseq using Poisson mixed models. Nucleic acids research, 45(11), e106.

Fang Z, et al. (2012) Statistical methods for identifying differentially expressed genes in RNA-Seq experiments. Cell & bioscience, 2(1), 26.