

Resource Summary Report

Generated by [NIF](#) on Oct 21, 2026

[seqbias](#)

RRID:SCR_006832

Type: Tool

Proper Citation

seqbias (RRID:SCR_006832)

Resource Information

URL: <http://bioconductor.org/packages/2.12/bioc/html/seqbias.html>

Proper Citation: seqbias (RRID:SCR_006832)

Description: Software package that implements a model of per-position sequencing bias in high-throughput sequencing data using a simple Bayesian network, the structure and parameters of which are trained on a set of aligned reads and a reference genome sequence.

Abbreviations: seqbias

Synonyms: seqbias - Estimation of per-position bias in high-throughput sequencing data

Resource Type: software resource

Keywords: bio.tools

Availability: GNU Lesser General Public License

Resource Name: seqbias

Resource ID: SCR_006832

Alternate IDs: OMICS_01237, biotools:seqbias, BioTools:seqbias

Alternate URLs: <https://bio.tools/seqbias>, <https://bio.tools/seqbias>, <https://bio.tools/seqbias>

Record Creation Time: 20220129T080238+0000

Record Last Update: 20261017T202643+0000

Ratings and Alerts

No rating or validation information has been found for seqbias.

No alerts have been found for seqbias.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.