

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## [dna-bison](#)

RRID:SCR\_005913

Type: Tool

### Proper Citation

dna-bison (RRID:SCR\_005913)

### Resource Information

**URL:** <http://sourceforge.net/projects/dna-bison/>

**Proper Citation:** dna-bison (RRID:SCR\_005913)

**Description:** Allows users with access to a computer cluster to rapidly align whole-genome bisulfite sequencing or RRBS reads.

**Abbreviations:** dna-bison

**Resource Type:** software resource

**Resource Name:** dna-bison

**Resource ID:** SCR\_005913

**Alternate IDs:** OMICS\_00582

**Record Creation Time:** 20220129T080233+0000

**Record Last Update:** 20260905T132543+0000

### Ratings and Alerts

No rating or validation information has been found for dna-bison.

No alerts have been found for dna-bison.

### 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 [Kravitz](#) .

Ryan DP, et al. (2014) Bison: bisulfite alignment on nodes of a cluster. BMC bioinformatics, 15(1), 337.