

Resource Summary Report

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

GNUMAP-BS

RRID:SCR_005995

Type: Tool

Proper Citation

GNUMAP-BS (RRID:SCR_005995)

Resource Information

URL: <http://dna.cs.byu.edu/gnumap/>

Proper Citation: GNUMAP-BS (RRID:SCR_005995)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 3rd,2023. A probabilistic algorithm that addresses the computational problems associated with aligning bisulfite sequencing data to a reference genome.

Abbreviations: GNUMAP-BS

Resource Type: software resource

Defining Citation: [PMID:24261665](#)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: GNUMAP-BS

Resource ID: SCR_005995

Alternate IDs: OMICS_00583

Record Creation Time: 20220129T080233+0000

Record Last Update: 20260905T132545+0000

Ratings and Alerts

No rating or validation information has been found for GNUMAP-BS.

No alerts have been found for GNUMAP-BS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Blango MG, et al. (2016) Identification of the long, edited dsRNAome of LPS-stimulated immune cells. Genome research, 26(6), 852.

Iorizzo M, et al. (2012) De novo assembly of the carrot mitochondrial genome using next generation sequencing of whole genomic DNA provides first evidence of DNA transfer into an angiosperm plastid genome. BMC plant biology, 12, 61.