

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 16, 2026

GBSA

RRID:SCR_003413

Type: Tool

Proper Citation

GBSA (RRID:SCR_003413)

Resource Information

URL: <http://ctrad-csi.nus.edu.sg/gbsa/>

Proper Citation: GBSA (RRID:SCR_003413)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on October 28,2025. Software for analyzing whole-genome bisulfite sequencing data.

Abbreviations: GBSA

Synonyms: Genome-Wide Bisulfite Sequencing Analyser Software, GBSA - Genome-Wide Bisulfite Sequencing Analyser Software

Resource Type: software resource

Defining Citation: [PMID:23268441](#)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: GBSA

Resource ID: SCR_003413

Alternate IDs: OMICS_00600

Record Creation Time: 20220129T080218+0000

Record Last Update: 20260912T195557+0000

Ratings and Alerts

No rating or validation information has been found for GBSA.

No alerts have been found for GBSA.

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 [SPARC SAWG](#) .

Yamamoto T, et al. (2023) DNA Methylation Aberrations in Dimethylarsinic Acid-Induced Bladder Carcinogenesis. Cancers, 15(21).

Corbet M, et al. (2021) Suppression of inflammatory arthritis by the parasitic worm product ES-62 is associated with epigenetic changes in synovial fibroblasts. PLoS pathogens, 17(11), e1010069.