Resource Summary Report

Generated by <u>NIF</u> on May 13, 2025

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: 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

Funding:

Availability: GNU General Public License, v3, Acknowledgement requested

Resource Name: GBSA

Resource ID: SCR_003413

Alternate IDs: OMICS_00600

Record Creation Time: 20220129T080218+0000

Record Last Update: 20250420T014142+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: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

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.