

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

loadBAM2ge_db

RRID:SCR_015951

Type: Tool

Proper Citation

loadBAM2ge_db (RRID:SCR_015951)

Resource Information

URL: http://cell-innovation.nig.ac.jp/maser/AllPipelines/P000001138_en.html

Proper Citation: loadBAM2ge_db (RRID:SCR_015951)

Description: Software pipeline that visualizes mapping results (in BAM format) on Genome Explorer.

Synonyms: loadBAM2ge_db (PE)

Resource Type: software resource, software application, data visualization software, data processing software

Keywords: bam, map, mapping, genome, pipeline, maser, visualization, explorer

Availability: Public

Resource Name: loadBAM2ge_db

Resource ID: SCR_015951

License: Free for academic use

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260808T190242+0000

Ratings and Alerts

No rating or validation information has been found for loadBAM2ge_db.

No alerts have been found for loadBAM2ge_db.

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 [SPARC SAWG](#) .

Kinjo S, et al. (2018) Maser: one-stop platform for NGS big data from analysis to visualization. Database : the journal of biological databases and curation, 2018.