

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

loadGtfToGe_db

RRID:SCR_015998

Type: Tool

Proper Citation

loadGtfToGe_db (RRID:SCR_015998)

Resource Information

URL: https://cell-innovation.nig.ac.jp/maser/Tools/visualization_top_en.html

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Description: One stop platform for NGS big data from analysis to visualization. There are about 400 analysis pipelines integrated on Maser. List of all analysis pipelines, including descriptions and approximate execution times, can be found on page for 'All pipelines' in the User Guide. loadGtfToGe_db software loads GTF files to a database for Genome Explorer. It allows the user to browse the results through the GE.

Synonyms: Management and Analysis System for Enormous Reads: loadGtfToGe_db

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

Keywords: analysis, pipeline, maser, genome, explorer, upload, database, browse

Funding: Ministry of Education ;
Culture ;
Sports ;
Science and Technology ;
Japan

Availability: Restricted

Resource Name: loadGtfToGe_db

Resource ID: SCR_015998

Alternate URLs: http://cell-innovation.nig.ac.jp/maser_cgi/cip-pl_list_violin_en.cgi

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260808T190047+0000

Ratings and Alerts

No rating or validation information has been found for loadGtfToGe_db.

No alerts have been found for loadGtfToGe_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.