

Resource Summary Report

Generated by [RRID](#) on Aug 14, 2026

loadGffToGe_db for custom genome

RRID:SCR_015997

Type: Tool

Proper Citation

loadGffToGe_db for custom genome (RRID:SCR_015997)

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. loadGffToGe_db for custom genome software loads GFF files of custom genomes 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: loadGffToGe_db for custom genome

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

Keywords: Next Generation Sequencers, Next Generation Sequencers data, Next Generation Sequencers data analysis and visualization,

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

Availability: Restricted

Resource Name: loadGffToGe_db for custom genome

Resource ID: SCR_015997

Alternate URLs: https://cell-innovation.nig.ac.jp/maser/index_en.html

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260814T042932+0000

Ratings and Alerts

No rating or validation information has been found for loadGffToGe_db for custom genome.

No alerts have been found for loadGffToGe_db for custom genome.

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 [RRID](#) .

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.