

# Resource Summary Report

Generated by [kravitz2](#) on Sep 20, 2026

## regist custom genome

RRID:SCR\_015999

Type: Tool

### Proper Citation

regist custom genome (RRID:SCR\_015999)

### Resource Information

**URL:** [https://cell-innovation.nig.ac.jp/maser/Tools/visualization\\_top\\_en.html](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.. Regist custom genome software registers custom genomes to Genome Explorer (IN: FASTA).

**Synonyms:** Management and Analysis System for Enormous Reads: regist custom genome

**Resource Type:** software resource, source code

**Keywords:** analysis, pipeline, maser, genome, explorer, fasta, custom, register

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

**Availability:** Restricted

**Resource Name:** regist custom genome

**Resource ID:** SCR\_015999

**Alternate URLs:** [http://cell-innovation.nig.ac.jp/maser\\_cgi/cip-pl\\_list\\_violin\\_en.cgi](http://cell-innovation.nig.ac.jp/maser_cgi/cip-pl_list_violin_en.cgi)

**Record Creation Time:** 20220129T080328+0000

**Record Last Update:** 20260919T195927+0000

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## Ratings and Alerts

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

No alerts have been found for regist custom genome.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

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.