

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 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

Proper Citation: regist custom genome (RRID:SCR_015999)

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

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260808T190639+0000

Ratings and Alerts

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

No alerts have been found for regist 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 [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.