

# Resource Summary Report

Generated by [NIF](#) on Sep 11, 2026

## GenoMiner

RRID:SCR\_001202

Type: Tool

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### Proper Citation

GenoMiner (RRID:SCR\_001202)

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### Resource Information

**URL:** <http://astridbio.com/genominer-genome-analyzer/>

**Proper Citation:** GenoMiner (RRID:SCR\_001202)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022. A next generation sequencing data analysis computer for biologists with or without IT background. It has an easy to-use graphical interface to analyze sequencing data in with only 15 clicks. A range of standard, add-on and custom applications help analyze and visualize data generated by Next Generation Sequencing machines. These are installed on each GenoMiner by default: \* Reference assembly \* De novo assembly \* ChiP-Seq \* BLAST \* Hybrid de novo assembly \* Hybrid reference assembly Add-on applications: \* Quality assesment \* RNA-Seq \* Copy Number Variation (CNV) \* Multiple Sequence Alignment \* miRNA-Seq \* Variant Calling

**Abbreviations:** GenoMiner

**Synonyms:** GenoMiner: Genome Analyzer

**Resource Type:** software resource

**Defining Citation:** [PMID:16267081](#)

**Keywords:** next-generation sequencing, reference assembly, de novo assembly, chip-seq, blast, hybrid de novo assembly, hybrid reference assembly, genome, computer, hardware, instrument, equipment

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** GenoMiner

**Resource ID:** SCR\_001202

**Alternate IDs:** OMICS\_02149

**Old URLs:** <http://www.astridbio.com/genominer.html>

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20260905T132431+0000

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## Ratings and Alerts

No rating or validation information has been found for GenoMiner.

No alerts have been found for GenoMiner.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.