

Resource Summary Report

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GenoMiner

RRID:SCR_001202

Type: Tool

Proper Citation

GenoMiner (RRID:SCR_001202)

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: 20260912T195521+0000

Ratings and Alerts

No rating or validation information has been found for GenoMiner.

No alerts have been found for GenoMiner.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.