Resource Summary Report

Generated by NIF on May 15, 2025

ATRHUNTER

RRID:SCR_006480

Type: Tool

Proper Citation

ATRHUNTER (RRID:SCR_006480)

Resource Information

URL: http://bioinfo.cs.technion.ac.il/atrhunter/

Proper Citation: ATRHUNTER (RRID:SCR_006480)

Description: Software that finds and displays approximate tandem repeats in DNA

sequences.

Abbreviations: ATRHUNTER

Resource Type: software resource, service resource, analysis service resource, production

service resource, data analysis service

Defining Citation: PMID:16201913

Keywords: bio.tools

Funding:

Resource Name: ATRHUNTER

Resource ID: SCR_006480

Alternate IDs: biotools:atrhunter, OMICS_00102

Alternate URLs: https://bio.tools/atrhunter

Record Creation Time: 20220129T080236+0000

Record Last Update: 20250514T061404+0000

Ratings and Alerts

No rating or validation information has been found for ATRHUNTER.

No alerts have been found for ATRHUNTER.

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 NIF.

Alves SIA, et al. (2024) What are microsatellites and how to choose the best tool: a user-friendly review of SSR and 74 SSR mining tools. Frontiers in genetics, 15, 1474611.