

Resource Summary Report

Generated by SPARC SAWG on Sep 15, 2026

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: analysis service resource, data analysis service, production service resource, service resource, software resource

Defining Citation: [PMID:16201913](#)

Keywords: bio.tools

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: 20260912T200012+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 [SPARC SAWG](#) .

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.