

# Resource Summary Report

Generated by [kravitz2](#) on Sep 5, 2026

## ATRHUNTER

RRID:SCR\_006480

Type: Tool

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### Proper Citation

ATRHUNTER (RRID:SCR\_006480)

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### 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:** 20260903T235802+0000

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### Ratings and Alerts

No rating or validation information has been found for ATRHUNTER.

No alerts have been found for ATRHUNTER.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

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.