

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## TRhist

RRID:SCR\_010769

Type: Tool

### Proper Citation

TRhist (RRID:SCR\_010769)

### Resource Information

**URL:** <http://trhist.gi.k.u-tokyo.ac.jp/>

**Proper Citation:** TRhist (RRID:SCR\_010769)

**Description:** A software tool for finding significant tandem repeats using short reads.

**Abbreviations:** TRhist

**Synonyms:** Trhist - a tandem repeat profiler in personal genomes

**Resource Type:** software resource

**Defining Citation:** [PMID:24215022](#)

**Resource Name:** TRhist

**Resource ID:** SCR\_010769

**Alternate IDs:** OMICS\_00117

**Record Creation Time:** 20220129T080300+0000

**Record Last Update:** 20260905T132646+0000

### Ratings and Alerts

No rating or validation information has been found for TRhist.

No alerts have been found for TRhist.

### 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 [Kravitz](#) .

Doi K, et al. (2014) Rapid detection of expanded short tandem repeats in personal genomics using hybrid sequencing. Bioinformatics (Oxford, England), 30(6), 815.