

# Resource Summary Report

Generated by [NIF](#) on Aug 26, 2026

## TE-locate

RRID:SCR\_012063

Type: Tool

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### Proper Citation

TE-locate (RRID:SCR\_012063)

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### Resource Information

**URL:** <http://sourceforge.net/projects/te-locate/>

**Proper Citation:** TE-locate (RRID:SCR\_012063)

**Description:** A software tool to locate all copies of sequences in a reference sequence using read-pairs.

**Resource Type:** software resource

**Defining Citation:** [PMID:24832231](#)

**Keywords:** java, perl

**Resource Name:** TE-locate

**Resource ID:** SCR\_012063

**Alternate IDs:** OMICS\_03507

**Record Creation Time:** 20220129T080308+0000

**Record Last Update:** 20260815T182445+0000

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### Ratings and Alerts

No rating or validation information has been found for TE-locate.

No alerts have been found for TE-locate.

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### 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](#) .

Ewing AD, et al. (2015) Transposable element detection from whole genome sequence data. Mobile DNA, 6, 24.