Resource Summary Report

Generated by RRID on May 13, 2025

Transposon Insertion Finder

RRID:SCR 001159

Type: Tool

Proper Citation

Transposon Insertion Finder (RRID:SCR_001159)

Resource Information

URL: https://github.com/akiomiyao/tif

Proper Citation: Transposon Insertion Finder (RRID:SCR_001159)

Description: A search program to detect insertions of transposable element from short

reads of next generation sequencer.

Abbreviations: TIF

Synonyms: Transposon Insertion Finder

Resource Type: software resource

Defining Citation: PMID:24629057

Keywords: perl, unix/linux

Funding:

Resource Name: Transposon Insertion Finder

Resource ID: SCR_001159

Alternate IDs: OMICS_03508

Record Creation Time: 20220129T080205+0000

Record Last Update: 20250420T014022+0000

Ratings and Alerts

No rating or validation information has been found for Transposon Insertion Finder.

No alerts have been found for Transposon Insertion Finder.

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 RRID.

Ding X, et al. (2020) Genome sequence of the agarwood tree Aquilaria sinensis (Lour.) Spreng: the first chromosome-level draft genome in the Thymelaeceae family. GigaScience, 9(3).