Resource Summary Report

Generated by T1D on Sep 5, 2026

MindTheGap

RRID:SCR_024115
Type: Tool

Proper Citation

MindTheGap (RRID:SCR_024115)

Resource Information

URL: https://github.com/GATB/MindTheGap

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Description: Software tool to perform detection and assembly of DNA insertion variants in NGS read datasets with respect to reference genome.Used to call insertions of any size, whether they are novel or duplicated, homozygous or heterozygous in the donor genome.

Synonyms: mindthegap

Resource Type: data analysis software, data processing software, software application, software resource

Keywords: perform detection and assembly of DNA insertion variants, NGS read datasets, reference genome, call insertions, donor genome,

Availability: Free, Available for download, Freely available,

Resource Name: MindTheGap

Resource ID: SCR_024115

Alternate IDs: OMICS_05472

Alternate URLs: https://sources.debian.org/src/mindthegap/

License: AGPL-3.0 license

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260903T235919+0000

Ratings and Alerts

No rating or validation information has been found for MindTheGap.

No alerts have been found for MindTheGap.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at T1D .

Schwarz JM, et al. (2021) Novel sequencing technologies and bioinformatic tools for deciphering the non-coding genome. Medizinische Genetik : Mitteilungsblatt des Berufsverbandes Medizinische Genetik e.V, 33(2), 133.

Godfroid M, et al. (2020) Insertion and deletion evolution reflects antibiotics selection pressure in a Mycobacterium tuberculosis outbreak. PLoS pathogens, 16(9), e1008357.

Daval S, et al. (2019) Computational analysis of the Plasmodiophora brassicae genome: mitochondrial sequence description and metabolic pathway database design. Genomics, 111(6), 1629.