

# Resource Summary Report

Generated by [T1D](#) on Sep 20, 2026

## [MapAI](#)

RRID:SCR\_004938

Type: Tool

### Proper Citation

MapAI (RRID:SCR\_004938)

### Resource Information

**URL:** <http://www.bioinf.boku.ac.at/pub/MapAI/>

**Proper Citation:** MapAI (RRID:SCR\_004938)

**Description:** A software tool for RNA-Seq expression profiling that builds on the established programs Bowtie and Cufflinks. Allowing an incorporation of "gene models" already at the alignment stage almost doubles the number of transcripts that can be measured reliably.

**Abbreviations:** MapAI

**Resource Type:** software resource

**Defining Citation:** [PMID:22485116](#)

**Keywords:** rna?seq

**Availability:** GNU General Public License

**Resource Name:** MapAI

**Resource ID:** SCR\_004938

**Alternate IDs:** OMICS\_01261

**Record Creation Time:** 20220129T080227+0000

**Record Last Update:** 20260919T195044+0000

### Ratings and Alerts

No rating or validation information has been found for MapAI.

No alerts have been found for MapAI.

## 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 [T1D](#) .

Polednik B, et al. (2021) COVID-19 lockdown and particle exposure of road users. Journal of transport & health, 22, 101233.