

Resource Summary Report

Generated by [RRID](#) on Apr 8, 2025

[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

Funding:

Availability: GNU General Public License

Resource Name: MapAI

Resource ID: SCR_004938

Alternate IDs: OMICS_01261

Record Creation Time: 20220129T080227+0000

Record Last Update: 20250214T183023+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 [RRID](#).

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