Resource Summary Report

Generated by NIF on May 21, 2025

MapAl

RRID:SCR_004938

Type: Tool

Proper Citation

MapAI (RRID:SCR_004938)

Resource Information

URL: http://www.bioinf.boku.ac.at/pub/MapAl/

Proper Citation: MapAl (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: MapAl

Resource Type: software resource

Defining Citation: PMID:22485116

Keywords: rna?seq

Funding:

Availability: GNU General Public License

Resource Name: MapAl

Resource ID: SCR_004938

Alternate IDs: OMICS_01261

Record Creation Time: 20220129T080227+0000

Record Last Update: 20250519T203340+0000

Ratings and Alerts

No rating or validation information has been found for MapAl.

No alerts have been found for MapAl.

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.

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