### **Resource Summary Report**

Generated by dkNET on Apr 28, 2025

# **MapAl**

RRID:SCR\_004938

Type: Tool

### **Proper Citation**

MapAl (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: 20250420T014241+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 dkNET.

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