# **Resource Summary Report**

Generated by dkNET on Apr 28, 2025

# **MotifLab**

RRID:SCR\_012649

Type: Tool

## **Proper Citation**

MotifLab (RRID:SCR\_012649)

#### **Resource Information**

URL: http://tare.medisin.ntnu.no/motiflab/

**Proper Citation:** MotifLab (RRID:SCR\_012649)

**Description:** Software for a general workbench for analyzing regulatory sequence regions

and discovering transcription factor binding sites and cis-regulatory modules.

**Abbreviations:** MotifLab

**Resource Type:** software resource

**Defining Citation:** PMID:23323883

Funding: Research Council of Norway

Availability: Acknowledgement requested, Free, Public

Resource Name: MotifLab

Resource ID: SCR\_012649

Alternate IDs: OMICS\_00486

**Record Creation Time:** 20220129T080311+0000

Record Last Update: 20250420T014616+0000

## Ratings and Alerts

No rating or validation information has been found for MotifLab.

No alerts have been found for MotifLab.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

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

Negrini F, et al. (2020) Genomic structure and transcript analysis of the Rapid Alkalinization Factor (RALF) gene family during host-pathogen crosstalk in Fragaria vesca and Fragaria x ananassa strawberry. PloS one, 15(3), e0226448.

Karpinski P, et al. (2018) Genome-wide analysis of gene expression after one year of venom immunotherapy. Immunology letters, 204, 23.