

# Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 10, 2026

## 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:** 20260808T190001+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](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [SPARC SAWG](#) .

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.