

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

MICSA

RRID:SCR_010860

Type: Tool

Proper Citation

MICSA (RRID:SCR_010860)

Resource Information

URL: <http://bioinfo-out.curie.fr/projects/micsa/>

Proper Citation: MICSA (RRID:SCR_010860)

Description: A software package for the identification of transcription factor binding sites in ChIP-Seq data, developed by Computational Systems Biology of Cancer group at the Bioinformatics Laboratory of Institut Curie (Paris).

Abbreviations: MICSA

Synonyms: MICSA: Motif Identification for ChIP-Seq Analysis, Motif Identification for ChIP-Seq Analysis

Resource Type: software resource

Keywords: bio.tools

Resource Name: MICSA

Resource ID: SCR_010860

Alternate IDs: biotools:micsa, OMICS_00447

Alternate URLs: <https://bio.tools/micsa>

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260725T190706+0000

Ratings and Alerts

No rating or validation information has been found for MICSA.

No alerts have been found for MICSA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.