

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

## MicroMUMMIE

RRID:SCR\_010847

Type: Tool

### Proper Citation

MicroMUMMIE (RRID:SCR\_010847)

### Resource Information

**URL:** <http://www.genome.duke.edu/labs/ohler/research/MUMMIE/mir.html>

**Proper Citation:** MicroMUMMIE (RRID:SCR\_010847)

**Description:** Software for a specific model, implemented within the MUMMIE framework, for predicting micro-RNA binding sites using PAR-CLIP data.

**Abbreviations:** MicroMUMMIE

**Resource Type:** software resource

**Defining Citation:** [PMID:23708386](#)

**Resource Name:** MicroMUMMIE

**Resource ID:** SCR\_010847

**Alternate IDs:** OMICS\_00401

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260725T190703+0000

### Ratings and Alerts

No rating or validation information has been found for MicroMUMMIE.

No alerts have been found for MicroMUMMIE.

### 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 [nidm-terms](#) .

Wessels HH, et al. (2019) Global identification of functional microRNA-mRNA interactions in Drosophila. Nature communications, 10(1), 1626.