

# Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 16, 2026

## MicroRazerS

RRID:SCR\_013316

Type: Tool

### Proper Citation

MicroRazerS (RRID:SCR\_013316)

### Resource Information

**URL:** <http://www.seqan.de/projects/microrazers/>

**Proper Citation:** MicroRazerS (RRID:SCR\_013316)

**Description:** A software tool optimized for mapping short RNAs onto a reference genome.

**Abbreviations:** MicroRazerS

**Resource Type:** software resource

**Defining Citation:** [PMID:19880369](#)

**Resource Name:** MicroRazerS

**Resource ID:** SCR\_013316

**Alternate IDs:** OMICS\_00371

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20260912T195802+0000

### Ratings and Alerts

No rating or validation information has been found for MicroRazerS.

No alerts have been found for MicroRazerS.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

### Usage and Citation Metrics

We found 5 mentions in open access literature.

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

King SE, et al. (2025) Prenatal maternal stress in rats alters the epigenetic and transcriptomic landscape of the maternal-fetal interface across four generations. Communications biology, 8(1), 38.

Ambeskovic M, et al. (2020) Ancestral stress programs sex-specific biological aging trajectories and non-communicable disease risk. Aging, 12(4), 3828.

Ambeskovic M, et al. (2019) Ancestral Stress Alters Lifetime Mental Health Trajectories and Cortical Neuromorphology via Epigenetic Regulation. Scientific reports, 9(1), 6389.

Ziemann M, et al. (2016) Evaluation of microRNA alignment techniques. RNA (New York, N.Y.), 22(8), 1120.

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.