

# Resource Summary Report

Generated by [T1D](#) on Sep 19, 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:** 20260919T195241+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 [T1D](#) .

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.