

Resource Summary Report

Generated by [NIF](#) on Aug 10, 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: 20260808T190010+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 [NIF](#) .

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.