

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## Single Read Paired Read Indel Substitution Minimizer

RRID:SCR\_018023

Type: Tool

### Proper Citation

Single Read Paired Read Indel Substitution Minimizer (RRID:SCR\_018023)

### Resource Information

**URL:** <https://github.com/ncbi/SRPRISM/>

**Proper Citation:** Single Read Paired Read Indel Substitution Minimizer (RRID:SCR\_018023)

**Description:** Software tool as single read paired read indel substitution minimizer.

**Abbreviations:** SRPRISM

**Synonyms:** Single Read Paired Read Indel Substitution Minimizer

**Resource Type:** software resource, software application, data processing software

**Keywords:** Single read, paired indel, substitution minimizer, next generation sequencing, data, indel

**Availability:** Free, Available for download, Freely available

**Resource Name:** Single Read Paired Read Indel Substitution Minimizer

**Resource ID:** SCR\_018023

**Alternate URLs:** <https://github.com/ncbi/SRPRISM/tree/master/srprism>

**Record Creation Time:** 20220129T080338+0000

**Record Last Update:** 20260731T043024+0000

### Ratings and Alerts

No rating or validation information has been found for Single Read Paired Read Indel Substitution Minimizer.

No alerts have been found for Single Read Paired Read Indel Substitution Minimizer.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Mario-Vasquez JE, et al. (2024) Finding Candida auris in public metagenomic repositories. PloS one, 19(1), e0291406.

Morgulis A, et al. (2020) SRPRISM (Single Read Paired Read Indel Substitution Minimizer): an efficient aligner for assemblies with explicit guarantees. GigaScience, 9(4).

Steinberg KM, et al. (2014) Single haplotype assembly of the human genome from a hydatidiform mole. Genome research, 24(12), 2066.