

# Resource Summary Report

Generated by [kravitz2](#) on Jul 30, 2026

## Sequence Read Format

RRID:SCR\_000132

Type: Tool

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### Proper Citation

Sequence Read Format (RRID:SCR\_000132)

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### Resource Information

**URL:** <http://srf.sourceforge.net/>

**Proper Citation:** Sequence Read Format (RRID:SCR\_000132)

**Description:** A generic format for DNA sequence data. The primary motivation for creating SRF has been to enable a single format capable of storing data generated by any DNA sequencing technology.

**Abbreviations:** SRF

**Synonyms:** Sequence Read Format (SRF)

**Resource Type:** data or information resource, interchange format, narrative resource, standard specification

**Keywords:** dna sequence, dna sequencing, interchange format

**Availability:** Public, A C++ implementation of Sequence Read Format is available

**Resource Name:** Sequence Read Format

**Resource ID:** SCR\_000132

**Alternate IDs:** OMICS\_05130

**Record Creation Time:** 20220129T080159+0000

**Record Last Update:** 20260729T043957+0000

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### Ratings and Alerts

No rating or validation information has been found for Sequence Read Format.

No alerts have been found for Sequence Read Format.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Cochrane G, et al. (2009) Petabyte-scale innovations at the European Nucleotide Archive. Nucleic acids research, 37(Database issue), D19.