

Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

Sequence Read Format

RRID:SCR_000132

Type: Tool

Proper Citation

Sequence Read Format (RRID:SCR_000132)

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: 20260912T195503+0000

Ratings and Alerts

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

No alerts have been found for Sequence Read Format.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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