

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

Bio-SCF

RRID:SCR_024068

Type: Tool

Proper Citation

Bio-SCF (RRID:SCR_024068)

Resource Information

URL: <https://metacpan.org/dist/Bio-SCF>

Proper Citation: Bio-SCF (RRID:SCR_024068)

Description: Software Perl extension for reading and writting SCF sequence files.This module provides Perl interface to SCF DNA sequencing files. It has both tied hash and an object-oriented interfaces. It provides the ability to read fields from SCF files and limited ability to modify them and write them back.

Synonyms: Bio-SCF - Perl extension for reading and writting SCF sequence files, libbio-scf-perl

Resource Type: software resource, software application

Keywords: Perl, Perl extension, reading and writting SCF sequence files,

Availability: Free, Available for download, Freely available,

Resource Name: Bio-SCF

Resource ID: SCR_024068

Alternate URLs: <https://sources.debian.org/src/libbio-scf-perl/>

License: perl_5

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260808T190539+0000

Ratings and Alerts

No rating or validation information has been found for Bio-SCF.

No alerts have been found for Bio-SCF.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.