

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 10, 2026

Sequin

RRID:SCR_016581

Type: Tool

Proper Citation

Sequin (RRID:SCR_016581)

Resource Information

URL: <https://www.ncbi.nlm.nih.gov/projects/Sequin/>

Proper Citation: Sequin (RRID:SCR_016581)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on November 12,2024. Software tool for DNA sequence submission. Used for submitting and updating entries to the GenBank or EMBL sequence databases.

Resource Type: standalone software, software application, software resource

Keywords: DNA, sequence, submission, GeneBank, EMBL, database

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Sequin

Resource ID: SCR_016581

Record Creation Time: 20220129T080331+0000

Record Last Update: 20260810T040646+0000

Ratings and Alerts

No rating or validation information has been found for Sequin.

No alerts have been found for Sequin.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Wang Y, et al. (2022) Comparative Analysis of Bacillariophyceae Chloroplast Genomes Uncovers Extensive Genome Rearrangements Associated with Speciation. International journal of environmental research and public health, 19(16).

Xu Q, et al. (2021) Comparative Analysis of Chloroplast Genomes of Seven Chaetoceros Species Revealed Variation Hotspots and Speciation Time. Frontiers in microbiology, 12, 742554.

Kundu S, et al. (2018) Complete mitochondrial genome of Black Soft-shell Turtle (*Nilssononia nigricans*) and comparative analysis with other Trionychidae. Scientific reports, 8(1), 17378.