

# Resource Summary Report

Generated by [ASWG](#) on Aug 4, 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:** software resource, software application, standalone software

**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:** 20260804T043651+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 2 mentions in open access literature.

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

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.