

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

seq-annot

RRID:SCR_018731

Type: Tool

Proper Citation

seq-annot (RRID:SCR_018731)

Resource Information

URL: <https://github.com/Brazelton-Lab/seq-annot>

Proper Citation: seq-annot (RRID:SCR_018731)

Description: Software Python package for annotating and counting genomic features in genomes and metagenomes. Software tools to facilitate annotation and comparison of genomes and metagenomes.

Resource Type: software application, software resource, software toolkit, standalone software

Keywords: Annotating, counting, comparison, genomic feature, genome, metagenome, metagenomics, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: seq-annot

Resource ID: SCR_018731

Alternate IDs: biotools:seq-annot

Alternate URLs: <https://bio.tools/seq-annot>

License: GNU Lesser General Public License v3.0

Record Creation Time: 20220129T080341+0000

Record Last Update: 20260919T195610+0000

Ratings and Alerts

No rating or validation information has been found for seq-annot.

No alerts have been found for seq-annot.

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 [nidm-terms](#) .

Thornton CN, et al. (2020) Localized effect of treated wastewater effluent on the resistome of an urban watershed. GigaScience, 9(11).