

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

Clinotator

RRID:SCR_016054

Type: Tool

Proper Citation

Clinotator (RRID:SCR_016054)

Resource Information

URL: <https://github.com/rbutleriii/Clinotator>

Proper Citation: Clinotator (RRID:SCR_016054)

Description: Software that performs clinical interpretation of ambiguous ClinVar annotations. This software takes batches of variants as input and queries NCBI eutilities to generate scoring metrics.

Synonyms: clinotator.py

Resource Type: software application, software resource

Keywords: clinical, age, weight, score, metric, vcf, python, nbc, annotation, variant, scoring, bio.tools

Availability: Free

Resource Name: Clinotator

Resource ID: SCR_016054

Alternate IDs: biotools:clinotator

Alternate URLs: <https://bio.tools/clinotator>

License: <https://github.com/rbutleriii/Clinotator/blob/master/LICENSE>

License URLs: <https://github.com/rbutleriii/Clinotator/blob/master/LICENSE>

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260912T200253+0000

Ratings and Alerts

No rating or validation information has been found for Clinotator.

No alerts have been found for Clinotator.

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 [SPARC SAWG](#) .

Schwarzerova J, et al. (2024) A perspective on genetic and polygenic risk scores-advances and limitations and overview of associated tools. Briefings in bioinformatics, 25(3).

Butler Iii RR, et al. (2018) Clinotator: analyzing ClinVar variation reports to prioritize reclassification efforts. F1000Research, 7, 462.