Resource Summary Report

Generated by <u>NIF</u> on Apr 26, 2025

VADR

RRID:SCR_024377 Type: Tool

Proper Citation

VADR (RRID:SCR_024377)

Resource Information

URL: https://github.com/nawrockie/vadr

Proper Citation: VADR (RRID:SCR_024377)

Description: Software suite of tools for classifying and analyzing sequences homologous to set of reference models of viral genomes or gene families.

Synonyms: ushahidi

Resource Type: data analysis software, software application, software resource, data processing software

Defining Citation: PMID:35441162

Keywords: classifying and analyzing sequences, sequences homologous to set of reference models, viral genomes reference models, gene families reference models,

Funding:

Availability: Free, Available for download, Freely available,

Resource Name: VADR

Resource ID: SCR_024377

Old URLs: https://sources.debian.org/src/vadr/

Record Creation Time: 20230830T050217+0000

Record Last Update: 20250426T061009+0000

Ratings and Alerts

No rating or validation information has been found for VADR.

No alerts have been found for VADR.

Data and Source Information

Source: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We have not found any literature mentions for this resource.