Resource Summary Report

Generated by SPARC SAWG on Jul 29, 2026

VAGUE

RRID:SCR_005607
Type: Tool

Proper Citation

VAGUE (RRID:SCR_005607)

Resource Information

URL: http://www.vicbioinformatics.com/software.vague.shtml

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Description: An open-source de novo genome assembly software tool, which is run from the Unix command line, providing a multi-platform graphical front-end for the Velvet de novo assembler. VAGUE is implemented in JRuby and targets the Java Virtual Machine.

Abbreviations: VAGUE

Synonyms: Velvet Assembler Graphical Front End

Resource Type: software resource

Defining Citation: PMID:23162059

Keywords: command line, assembler

Availability: GNU General Public License, v2, Acknowledgement requested

Resource Name: VAGUE

Resource ID: SCR_005607

Alternate IDs: OMICS_00897

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260725T190609+0000

Ratings and Alerts

No rating or validation information has been found for VAGUE.

No alerts have been found for VAGUE.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at SPARC SAWG .

Wilk M, et al. (2024) Pain catastrophizing in rheumatic diseases: prevalence, origin, and implications. Rheumatology international, 44(6), 985.

Zhao Z, et al. (2019) Comparative genomics reveal pathogenicity-related loci in Pseudomonas syringae pv. actinidiae biovar 3. Molecular plant pathology, 20(7), 923.

Guerrero A, et al. (2019) Whole-genome comparison between reference sequences and oyster Vibrio vulnificus C-genotype strains. PloS one, 14(7), e0220385.

Guerrero A, et al. (2017) Genetic Analysis of Vibrio parahaemolyticus O3:K6 Strains That Have Been Isolated in Mexico Since 1998. PloS one, 12(1), e0169722.

Lee W, et al. (2010) Celecoxib does not attenuate the antiplatelet effects of aspirin and clopidogrel in healthy volunteers. Korean circulation journal, 40(7), 321.