Resource Summary Report

Generated by <u>dkNET</u> on Apr 29, 2025

<u>bamova</u>

RRID:SCR_000510 Type: Tool

Proper Citation

bamova (RRID:SCR_000510)

Resource Information

URL: http://www.uwyo.edu/buerkle/software/bamova/

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Description: Software that implements a Bayesian Analysis of Molecular Variance and different likelihood models for three different types of molecular data (including two models for high throughput sequence data).

Abbreviations: bamova

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

Defining Citation: PMID:21212231

Keywords: mac os x, high throughput sequence

Funding: NSF 701757

Availability: Free, Available for download, Freely available,

Resource Name: bamova

Resource ID: SCR_000510

Alternate IDs: OMICS_01532

License: GNU General Public License, v3 or later

Record Creation Time: 20220129T080202+0000

Ratings and Alerts

No rating or validation information has been found for bamova.

No alerts have been found for bamova.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.