Mauve
RRID:SCR_012852
Type: Tool

Proper Citation
Mauve (RRID:SCR_012852)

Resource Information

URL: http://gel.ahabs.wisc.edu/mauve/

Description: A system for efficiently constructing multiple genome alignments in the presence of large-scale evolutionary events such as rearrangement and inversion.

Resource Name: Mauve
Proper Citation: Mauve (RRID:SCR_012852)
Resource Type: Resource, software resource
Resource ID: SCR_012852
Parent Organization: University of Wisconsin-Madison; Wisconsin; USA
Website Status: Last checked down
Alternate IDs: OMICS_00940
Abbreviations: Mauve
Mentions Count: 627

Ratings and Alerts

No rating or validation information has been found for Mauve.
No alerts have been found for Mauve.
Usage and Citation Metrics

We found 627 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](https://www.fdi-lab.com).


Nascimento FX, et al. (2020) Plant growth-promoting activities and genomic analysis of the stress-resistant STB1, a bacterium of agricultural and biotechnological interest. Biotechnology reports (Amsterdam, Netherlands), 25, e00406.


Kawalek A, et al. (2020) Genome sequence of Pseudomonas aeruginosa PAO1161, a PAO1 derivative with the ICEPae1161 integrative and conjugative element. BMC genomics, 21(1), 14.


Zwickl NF, et al. (2020) Comparative genome characterization of the periodontal pathogen Tannerella forsythia. BMC genomics, 21(1), 150.


Makke G, et al. (2020) Whole-Genome-Sequence-Based Characterization of Extensively Drug-Resistant Acinetobacter baumannii Hospital Outbreak. mSphere, 5(1).


