Ariba
RRID:SCR_015976
Type: Tool

Proper Citation
Ariba (RRID:SCR_015976)

Resource Information

URL: https://github.com/sanger-pathogens/ariba

Proper Citation: Ariba (RRID:SCR_015976)

Description: Analysis software that identifies antibiotic resistance genes by running local assemblies. It can also be used for MLST calling.

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

References: PMID:29177089

Keywords: software, analysis, tool, sequence, antibiotic, resistance, assembly, local, mlst

Funding Agency: Biotechnology and Biological Sciences Research Council, Wellcome Trust

Availability: Free, Available for download, Freely available

Website Status: Last checked up

Resource Name: Ariba

Resource ID: SCR_015976

Ratings and Alerts
No rating or validation information has been found for Ariba.
No alerts have been found for Ariba.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 67 mentions in open access literature.

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

Madden DE, et al. (2021) Taking the next-gen step: Comprehensive antimicrobial resistance detection from Burkholderia pseudomallei. EBioMedicine, 63, 103152.


Duong VT, et al. (2020) Genomic Serotyping, Clinical Manifestations, and Antimicrobial Resistance of Nontyphoidal Gastroenteritis in Hospitalized Children in Ho Chi Minh City,


Constantinides B, et al. (2020) Genomic surveillance of and spp. in hospital sink drains and patients. Microbial genomics, 6(7).


Magalhães B, et al. (2020) Combining Standard Molecular Typing and Whole Genome Sequencing to Investigate Epidemiology in Intensive Care Units. Frontiers in public health, 8, 3.
