Ariba

RRID:SCR_015976
Type: Tool

Proper Citation

Ariba (RRID:SCR_015976)

Resource Information

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

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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

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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 91 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch.org.


Haim MS, et al. (2021) Comparative genomics of ST5 and ST30 methicillin-resistant sequential isolates recovered from paediatric patients with cystic fibrosis. Microbial genomics, 7(3).


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


Montilla-Escudero EA, et al. (2021) Draft Genome Sequences of Corynebacterium diphtheriae Clinical Isolates from Colombia. Microbiology resource announcements, 10(29), e0033521.


Di Gregorio S, et al. (2021) Genomic Epidemiology of CC30 Methicillin-Resistant Staphylococcus aureus Strains from Argentina Reveals Four Major Clades with Distinctive Genetic Features. mSphere, 6(2).

McGuinness SL, et al. (2021) Clinical and Molecular Epidemiology of an Emerging Panton-Valentine Leukocidin-Positive ST5 Methicillin-Resistant Staphylococcus aureus Clone in Northern Australia. mSphere, 6(1).