

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

Kaptive

RRID:SCR_024046

Type: Tool

Proper Citation

Kaptive (RRID:SCR_024046)

Resource Information

URL: <https://github.com/klebgenomics/Kaptive>

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Description: Software tool to report information about surface polysaccharide loci for *Klebsiella pneumoniae* species complex and *Acinetobacter baumannii* genome assemblies.

Synonyms: kaptive

Resource Type: software application, software resource

Defining Citation: [PMID:28348840](#)

Keywords: report information, surface polysaccharide loci, genome assemblies,

Availability: Free, Available for download, Freely available,

Resource Name: Kaptive

Resource ID: SCR_024046

Alternate IDs: OMICS_27932

Alternate URLs: <https://sources.debian.org/src/kaptive/>

License: GPL-3.0 license

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260727T040818+0000

Ratings and Alerts

No rating or validation information has been found for Kaptive.

No alerts have been found for Kaptive.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#).

Stanton TD, et al. (2025) Fast and accurate in silico antigen typing with Kaptive 3. Microbial genomics, 11(6).

Lean SS, et al. (2025) The Road Less Traveled: Exploring the Genomic Characteristics and Antimicrobial Resistance Potential of *Acinetobacter baumannii* From the Indigenous Orang Asli Community in Peninsular Malaysia. MicrobiologyOpen, 14(5), e70073.

Boeckaerts D, et al. (2024) Prediction of *Klebsiella* phage-host specificity at the strain level. Nature communications, 15(1), 4355.

Shpirt AM, et al. (2024) An *Acinetobacter baumannii* nasal carriage isolate recovered from an asymptomatic patient in Vietnam is extensively antibiotic resistant and produces a rare K71 type capsule. Microbiology spectrum, 12(12), e0183824.

Feng Y, et al. (2024) Population genomics uncovers global distribution, antimicrobial resistance, and virulence genes of the opportunistic pathogen *Klebsiella aerogenes*. Cell reports, 43(8), 114602.

Slarve MJ, et al. (2024) Clinical assays rapidly predict bacterial susceptibility to monoclonal antibody therapy. JCI insight, 9(2).

Amadesi S, et al. (2024) Complete Genome Sequence of a *Klebsiella pneumoniae* Strain Carrying Novel Variant blaKPC-203, Cross-Resistant to Ceftazidime/Avibactam and Cefiderocol, but Susceptible to Carbapenems, Isolated in Italy, 2023. Pathogens (Basel, Switzerland), 13(6).

Yang Y, et al. (2024) Large-scale genomic survey with deep learning-based method reveals strain-level phage specificity determinants. GigaScience, 13.

Koncz M, et al. (2024) Genomic surveillance as a scalable framework for precision phage therapy against antibiotic-resistant pathogens. Cell, 187(21), 5901.