

Takehiro Tomita

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8078978/publications.pdf>

Version: 2024-02-01

10
papers

1,595
citations

1162367

8
h-index

1372195

10
g-index

11
all docs

11
docs citations

11
times ranked

2983
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | SRST2: Rapid genomic surveillance for public health and hospital microbiology labs. <i>Genome Medicine</i> , 2014, 6, 90. | 3.6 | 953 |
| 2 | Prospective Whole-Genome Sequencing Enhances National Surveillance of <i>Listeria monocytogenes</i> . <i>Journal of Clinical Microbiology</i> , 2016, 54, 333-342. | 1.8 | 239 |
| 3 | Intercontinental dissemination of azithromycin-resistant shigellosis through sexual transmission: a cross-sectional study. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 913-921. | 4.6 | 204 |
| 4 | Translating genomics into practice for real-time surveillance and response to carbapenemase-producing Enterobacteriaceae: evidence from a complex multi-institutional KPC outbreak. <i>PeerJ</i> , 2018, 6, e4210. | 0.9 | 66 |
| 5 | Emergence of endemic MLST non-typeable vancomycin-resistant <i>Enterococcus faecium</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 3367-3371. | 1.3 | 44 |
| 6 | Comparative Genomics Shows That <i>Mycobacterium ulcerans</i> Migration and Expansion Preceded the Rise of Buruli Ulcer in Southeastern Australia. <i>Applied and Environmental Microbiology</i> , 2018, 84, . | 1.4 | 32 |
| 7 | Comprehensive antibiotic-linked mutation assessment by resistance mutation sequencing (RM-seq). <i>Genome Medicine</i> , 2018, 10, 63. | 3.6 | 26 |
| 8 | A nanomechanical study of the effects of colistin on the <i>Klebsiella pneumoniae</i> AJ218 capsule. <i>European Biophysics Journal</i> , 2017, 46, 351-361. | 1.2 | 12 |
| 9 | Complete microbial genomes for public health in Australia and the Southwest Pacific. <i>Microbial Genomics</i> , 2020, 6, . | 1.0 | 10 |
| 10 | A Supervised Statistical Learning Approach for Accurate <i>Legionella pneumophila</i> Source Attribution during Outbreaks. <i>Applied and Environmental Microbiology</i> , 2017, 83, . | 1.4 | 8 |