Luiz Irber

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9274839/publications.pdf

Version: 2024-02-01

| | | 1307594 | 1372567 | |
|----------|----------------|--------------|----------------|--|
| 10 | 1,055 | 7 | 10 | |
| papers | citations | h-index | g-index | |
| | | | | |
| | | | | |
| | | | | |
| 18 | 18 | 18 | 1968 | |
| all docs | docs citations | times ranked | citing authors | |
| | | | | |

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Critical Assessment of Metagenome Interpretation: the second round of challenges. Nature Methods, 2022, 19, 429-440. | 19.0 | 133 |
| 2 | Streamlining data-intensive biology with workflow systems. GigaScience, 2021, 10, . | 6.4 | 32 |
| 3 | Context-aware genomic surveillance reveals hidden transmission of a carbapenemase-producing Klebsiella pneumoniae. Microbial Genomics, $2021, 7, .$ | 2.0 | 9 |
| 4 | Large-scale sequence comparisons with sourmash. F1000Research, 2019, 8, 1006. | 1.6 | 130 |
| 5 | khmer release v2.1: software for biological sequence analysis. Journal of Open Source Software, 2017, 2, 272. | 4.6 | 4 |
| 6 | Haplotype-Phased Synthetic Long Reads from Short-Read Sequencing. PLoS ONE, 2016, 11, e0147229. | 2.5 | 29 |
| 7 | sourmash: a library for MinHash sketching of DNA. Journal of Open Source Software, 2016, 1, 27. | 4.6 | 230 |
| 8 | The khmer software package: enabling efficient nucleotide sequence analysis. F1000Research, 2015, 4, 900. | 1.6 | 362 |
| 9 | Climate Simulation and Change in the Brazilian Climate Model. Journal of Climate, 2013, 26, 6716-6732. | 3.2 | 61 |
| 10 | An objective reference system for studying rings in the ocean. Computers and Geosciences, 2013, 61, 43-49. | 4.2 | 1 |