## Julio Guerra

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4474446/publications.pdf

Version: 2024-02-01

| 18<br>papers | 176<br>citations | 7<br>h-index | 1125743<br>13<br>g-index |
|--------------|------------------|--------------|--------------------------|
| 20           | 20               | 20           | 285                      |
| all docs     | docs citations   | times ranked | citing authors           |

| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 1  | A first insight on the population structure of Mycobacterium tuberculosis complex as studied by spoligotyping and MIRU-VNTRs in Bogot $	ilde{A}_i$ , Colombia. Infection, Genetics and Evolution, 2012, 12, 657-663.               | 2.3 | 42        |
| 2  | Comparative genomic analysis of Mycobacterium tuberculosis Beijing-like strains revealed specific genetic variations associated with virulence and drug resistance. Infection, Genetics and Evolution, 2017, 54, 314-323.          | 2.3 | 30        |
| 3  | Virulence and Immune Response Induced by Mycobacterium avium Complex Strains in a Model of Progressive Pulmonary Tuberculosis and Subcutaneous Infection in BALB/c Mice. Infection and Immunity, 2013, 81, 4001-4012.              | 2.2 | 18        |
| 4  | First case of multidrug-resistant tuberculosis caused by a rare "Beijing-like―genotype of Mycobacterium tuberculosis in Bogotá, Colombia. Infection, Genetics and Evolution, 2010, 10, 678-681.                                    | 2.3 | 16        |
| 5  | Circulation of <i>M. tuberculosis</i> Beijing genotype in Latin America and the Caribbean. Pathogens and Global Health, 2019, 113, 336-351.  | 2.3 | 13        |
| 6  | Active and latent tuberculosis among inmates in La Esperanza prison in Guaduas, Colombia. PLoS ONE, 2019, 14, e0209895.  | 2.5 | 12        |
| 7  | Complete Genome Sequence of the Clinical Beijing-Like Strain Mycobacterium tuberculosis 323 Using the PacBio Real-Time Sequencing Platform. Genome Announcements, $2015, 3, .$   | 0.8 | 10        |
| 8  | Profiling the immune response to <i>Mycobacterium tuberculosis</i> Beijing family infection: a perspective from the transcriptome. Virulence, 2021, 12, 1689-1704.   | 4.4 | 9         |
| 9  | First approach to the population structure of Mycobacterium tuberculosis complex in the indigenous population in Puerto Nariño-Amazonas, Colombia. PLoS ONE, 2021, 16, e0245084.   | 2.5 | 5         |
| 10 | Population structure of multidrug-resistant Mycobacterium tuberculosis clinical isolates in Colombia. Tuberculosis, 2020, 125, 102011.   | 1.9 | 4         |
| 11 | Characterization of clinical isolates of Mycobacterium tuberculosis from indigenous peoples of Colombia. Biomedica, 2019, 39, 78-92.   | 0.7 | 3         |
| 12 | Heterogeneous fitness landscape cues, pknG high expression, and phthiocerol dimycocerosate low production of Mycobacterium tuberculosis ATCC25618 rpoB S450L in enriched broth. Tuberculosis, 2022, 132, 102156.                   | 1.9 | 3         |
| 13 | Close Related Drug-Resistance Beijing Isolates of Mycobacterium tuberculosis Reveal a Different<br>Transcriptomic Signature in a Murine Disease Progression Model. International Journal of Molecular<br>Sciences, 2022, 23, 5157. | 4.1 | 3         |
| 14 | Genetic diversity of Mycobacterium tuberculosis clinical isolates from HIV-TB patients from two public hospitals at $Bogot\tilde{A}_i$ , $Colombia$ . Infection, Genetics and Evolution, 2020, 77, 104059.                         | 2.3 | O         |
| 15 | Title is missing!. , 2021, 16, e0245084.   |     | 0         |
| 16 | Title is missing!. , 2021, 16, e0245084.   |     | 0         |
| 17 | Title is missing!. , 2021, 16, e0245084.   |     | O         |
| 18 | Title is missing!. , 2021, 16, e0245084.   |     | 0         |