

MarÃ-a-JosÃ© Iglesias

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8266435/publications.pdf>

Version: 2024-02-01

10
papers

255
citations

1307594

7
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

370
citing authors

#	ARTICLE	IF	CITATIONS
1	Incidence and risk factors for childhood brain tumors in the ILE DE France. International Journal of Cancer, 1994, 59, 776-782.	5.1	124
2	Data-driven model for the assessment of <i>Mycobacterium tuberculosis</i> transmission in evolving demographic structures. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E3238-E3245.	7.1	36
3	Conspicuous multidrug-resistant <i>Mycobacterium tuberculosis</i> cluster strains do not trespass country borders in Latin America and Spain. Infection, Genetics and Evolution, 2012, 12, 711-717.	2.3	30
4	Long-term molecular surveillance of multidrug-resistant tuberculosis in Spain. Infection, Genetics and Evolution, 2012, 12, 701-710.	2.3	18
5	Investigation of a rapidly spreading tuberculosis outbreak using whole-genome sequencing. Infection, Genetics and Evolution, 2020, 81, 104184.	2.3	13
6	Single nucleotide polymorphism (SNP) analysis used for the phylogeny of the <i>Mycobacterium tuberculosis</i> complex based on a pyrosequencing assay. BMC Microbiology, 2014, 14, 21.	3.3	11
7	The value of the continuous genotyping of multi-drug resistant tuberculosis over 20 years in Spain. Scientific Reports, 2020, 10, 20433.	3.3	9
8	Editorial Commentary: Nonspecific Beneficial Effects of BCG Vaccination in High-income Countries, Should We Extend Recommendation of BCG Vaccination?. Clinical Infectious Diseases, 2015, 60, 1620-1621.	5.8	8
9	<i>Mycobacterium tuberculosis</i> genotypes and predominant clones among the multidrug-resistant isolates in Spain 1998–2005. Infection, Genetics and Evolution, 2017, 55, 117-126.	2.3	3
10	The MtZ Strain: Molecular Characteristics and Outbreak Investigation of the Most Successful <i>Mycobacterium tuberculosis</i> Strain in Aragon Using Whole-Genome Sequencing. Frontiers in Cellular and Infection Microbiology, 2022, 12, .	3.9	3