Sara MonzÃ³n

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Monkeypox outbreak in Madrid (Spain): Clinical and virological aspects. Journal of Infection, 2022, 85, 412-417.	1.7	109
2	Emergence of NDM-producing Klebsiella pneumoniae and Escherichia coli in Spain: phylogeny, resistome, virulence and plasmids encoding blaNDM-like genes as determined by WGS. Journal of Antimicrobial Chemotherapy, 2019, 74, 3489-3496.	1.3	60
3	Insight into the Significance of Aspergillus fumigatus cyp51A Polymorphisms. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	57
4	A Founder Effect Led Early SARS-CoV-2 Transmission in Spain. Journal of Virology, 2021, 95, .	1.5	55
5	Genome-Wide Comparative Analysis of Aspergillus fumigatus Strains: The Reference Genome as a Matter of Concern. Genes, 2018, 9, 363.	1.0	51
6	Liver organoids reproduce alpha-1 antitrypsin deficiency-related liver disease. Hepatology International, 2020, 14, 127-137.	1.9	44
7	Familial retinoblastoma due to intronic LINE-1 insertion causes aberrant and noncanonical mRNA splicing of the RB1 gene. Journal of Human Genetics, 2016, 61, 463-466.	1.1	33
8	EWS-FL11-mediated suppression of the RAS-antagonist Sprouty 1 (SPRY1) confers aggressiveness to Ewing sarcoma. Oncogene, 2017, 36, 766-776.	2.6	29
9	SpainUDP: The Spanish Undiagnosed Rare Diseases Program. International Journal of Environmental Research and Public Health, 2018, 15, 1746.	1.2	19
10	Frequency of low-level and high-level mosaicism in sporadic retinoblastoma: genotype–phenotype relationships. Journal of Human Genetics, 2020, 65, 165-174.	1.1	16
11	Adaptation of the emerging pathogenic yeast <i>Candida auris</i> to high caspofungin concentrations correlates with cell wall changes. Virulence, 2021, 12, 1400-1417.	1.8	15
12	MicroRNA Profile of HCV Spontaneous Clarified Individuals, Denotes Previous HCV Infection. Journal of Clinical Medicine, 2019, 8, 849.	1.0	11
13	Downregulation of A20 Expression Increases the Immune Response and Apoptosis and Reduces Virus Production in Cells Infected by the Human Respiratory Syncytial Virus. Vaccines, 2020, 8, 100.	2.1	11
14	Saezia sanguinis gen. nov., sp. nov., a Betaproteobacteria member of order Burkholderiales, isolated from human blood. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2016-2025.	0.8	10
15	Genomic Background and Phylogeny of cfiA-Positive Bacteroides fragilis Strains Resistant to Meropenem-EDTA. Antibiotics, 2021, 10, 304.	1.5	8
16	Proficiency Testing of Metagenomics-Based Detection of Food-Borne Pathogens Using a Complex Artificial Sequencing Dataset. Frontiers in Microbiology, 2020, 11, 575377.	1.5	7
17	De novo small deletion affecting transcription start site of short isoform of <scp><i>AUTS2</i></scp> gene in a patient with syndromic neurodevelopmental defects. American Journal of Medical Genetics, Part A, 2021, 185, 877-883.	0.7	5
18	Molecular Characterization and Antimicrobial Susceptibilities of Nocardia Species Isolated from the Soil; A Comparison with Species Isolated from Humans. Microorganisms, 2020, 8, 900.	1.6	4

#	Article	IF	CITATIONS
19	First Draft Genome Sequence of a Clinical Strain of Nocardia cerradoensis. Genome Announcements, 2017, 5, .	0.8	2
20	First Insight into the Genome Sequences of Two Linezolid-Resistant Nocardia farcinica Strains Isolated from Patients with Cystic Fibrosis. Genome Announcements, 2017, 5, .	0.8	2