

Rosario Morales-Espinosa

List of Publications by Year in descending order

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29
papers

700
citations

567281

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33
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docs citations

33
times ranked

1231
citing authors

#	ARTICLE	IF	CITATIONS
1	Diversity in the composition of the accessory genome of Mexican <i>Pseudomonas aeruginosa</i> strains. <i>Genes and Genomics</i> , 2021, , 1.	1.4	0
2	Diversity of Potentially Pathogenic <i>Escherichia coli</i> O104 and O9 Serogroups Isolated before 2011 from Fecal Samples from Children from Different Geographic Regions. <i>Microorganisms</i> , 2021, 9, 2227.	3.6	3
3	Characterization of commensal <i>Escherichia coli</i> isolates from slaughtered sheep in Mexico. <i>Journal of Infection in Developing Countries</i> , 2021, 15, 1755-1760.	1.2	1
4	The receptor binding domain of SARS-CoV-2 spike protein is the result of an ancestral recombination between the bat-CoV RaTG13 and the pangolin-CoV MP789. <i>BMC Research Notes</i> , 2020, 13, 398.	1.4	31
5	High expression of <i>Helicobacter pylori</i> VapD in both the intracellular environment and biopsies from gastric patients with severity. <i>PLoS ONE</i> , 2020, 15, e0230220.	2.5	11
6	Title is missing!. , 2020, 15, e0230220.		0
7	Title is missing!. , 2020, 15, e0230220.		0
8	Title is missing!. , 2020, 15, e0230220.		0
9	Title is missing!. , 2020, 15, e0230220.		0
10	The Genomic Basis of Intrinsic and Acquired Antibiotic Resistance in the Genus <i>Serratia</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 828.	3.5	53
11	Complete Genome Sequences of Two <i>Pseudomonas aeruginosa</i> Strains Isolated from Children with Bacteremia. <i>Genome Announcements</i> , 2017, 5, .	0.8	4
12	Complete Genome Sequences of Four Extensively Drug-Resistant <i>Pseudomonas aeruginosa</i> Strains, Isolated from Adults with Ventilator-Associated Pneumonia at a Tertiary Referral Hospital in Mexico City. <i>Genome Announcements</i> , 2017, 5, .	0.8	5
13	Integrated view of <i>Vibrio cholerae</i> in the Americas. <i>Science</i> , 2017, 358, 789-793.	12.6	128
14	Fingerprint Analysis and Identification of Strains ST309 as a Potential High Risk Clone in a <i>Pseudomonas aeruginosa</i> Population Isolated from Children with Bacteremia in Mexico City. <i>Frontiers in Microbiology</i> , 2017, 8, 313.	3.5	26
15	<i>Pseudomonas aeruginosa</i> ATCC 9027 is a non-virulent strain suitable for mono-rhamnolipids production. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 9995-10004.	3.6	47
16	Complete Genome Sequence of <i>Serratia marcescens</i> SmUNAM836, a Nonpigmented Multidrug-Resistant Strain Isolated from a Mexican Patient with Obstructive Pulmonary Disease. <i>Genome Announcements</i> , 2016, 4, .	0.8	17
17	UPEC strain characterization isolated from Mexican patients with recurrent urinary infections. <i>Journal of Infection in Developing Countries</i> , 2016, 10, 317-328.	1.2	18
18	Occurrence in Mexico, 1998–2008, of <i>Vibrio cholerae</i> CTX ⁺ El Tor carrying an additional truncated CTX prophage. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 9917-9922.	7.1	20

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19	<i>Pseudomonas aeruginosa</i> clinical and environmental isolates constitute a single population with high phenotypic diversity. <i>BMC Genomics</i> , 2014, 15, 318.	2.8	85
20	<i>E. coli</i> outbreak in a neonate intensive care unit in a general hospital in Mexico City. <i>Folia Microbiologica</i> , 2013, 58, 229-234.	2.3	1
21	Genetic Characterization of Atypical <i>Citrobacter freundii</i> . <i>PLoS ONE</i> , 2013, 8, e74120.	2.5	12
22	Genetic and Phenotypic Characterization of a <i>Pseudomonas aeruginosa</i> Population with High Frequency of Genomic Islands. <i>PLoS ONE</i> , 2012, 7, e37459.	2.5	20
23	<i>Vibrio cholerae</i> Classical Biotype Strains Reveal Distinct Signatures in Mexico. <i>Journal of Clinical Microbiology</i> , 2012, 50, 2212-2216.	3.9	32
24	New enterovirulent <i>Escherichia coli</i> serogroup 64474 showing antigenic and genotypic relationships to <i>Shigella boydii</i> 16. <i>Journal of Medical Microbiology</i> , 2010, 59, 453-461.	1.8	10
25	Cholera between 1991 and 1997 in Mexico Was Associated with Infection by Classical, El Tor, and El Tor Variants of <i>Vibrio cholerae</i> . <i>Journal of Clinical Microbiology</i> , 2010, 48, 3666-3674.	3.9	34
26	<i>Helicobacter pylori</i> in the oral cavity is associated with gastroesophageal disease. <i>Oral Microbiology and Immunology</i> , 2009, 24, 464-468.	2.8	27
27	Lewis Antigen Expression by <i>Helicobacter pylori</i> Strains Colonizing Different Regions of the Stomach of Individual Patients. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2783-2785.	3.9	6
28	Bleeding peptic ulcers and presence of <i>Helicobacter pylori</i> by various tests: a case-control study. <i>European Journal of Gastroenterology and Hepatology</i> , 2002, 14, 1113-1118.	1.6	23
29	Colonization of Mexican Patients by Multiple <i>Helicobacter pylori</i> Strains with Different <i>vacA</i> and <i>cagA</i> Genotypes. <i>Journal of Clinical Microbiology</i> , 1999, 37, 3001-3004.	3.9	86