

Deyanira PÃ©rez-Morales

List of Publications by Year in descending order

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27
papers

884
citations

623188

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525886

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27
all docs

27
docs citations

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times ranked

1109
citing authors

#	ARTICLE	IF	CITATIONS
1	Regulatory Evolution of the <i>phoH</i> Ancestral Gene in <i>Salmonella enterica</i> Serovar Typhimurium. <i>Journal of Bacteriology</i> , 2022, 204, e0058521.	1.0	2
2	(p)ppGpp-Dependent Regulation of the Nucleotide Hydrolase PpnN Confers Complement Resistance in <i>Salmonella enterica</i> Serovar Typhimurium. <i>Infection and Immunity</i> , 2021, 89, .	1.0	2
3	Cross-kingdom metabolic manipulation promotes <i>Salmonella</i> replication inside macrophages. <i>Nature Communications</i> , 2021, 12, 1862.	5.8	2
4	An incoherent feedforward loop formed by SirA/BarA, HilE and HilD is involved in controlling the growth cost of virulence factor expression by <i>Salmonella</i> Typhimurium. <i>PLoS Pathogens</i> , 2021, 17, e1009630.	2.1	12
5	Genomic Analysis Reveals the Genetic Determinants Associated With Antibiotic Resistance in the Zoonotic Pathogen <i>Campylobacter</i> spp. Distributed Globally. <i>Frontiers in Microbiology</i> , 2020, 11, 513070.	1.5	14
6	The <i>Salmonella</i> Typhimurium InvF-SicA complex is necessary for the transcription of <i>sopB</i> in the absence of the repressor H-NS. <i>PLoS ONE</i> , 2020, 15, e0240617.	1.1	9
7	HilD induces expression of a novel <i>Salmonella</i> Typhimurium invasion factor, YobH, through a regulatory cascade involving SprB. <i>Scientific Reports</i> , 2019, 9, 12725.	1.6	12
8	SlyA and HilD Counteract H-NS-Mediated Repression on the <i>ssrAB</i> Virulence Operon of <i>Salmonella enterica</i> Serovar Typhimurium and Thus Promote Its Activation by OmpR. <i>Journal of Bacteriology</i> , 2019, 201, .	1.0	23
9	HilD and PhoP independently regulate the expression of <i>grhD1</i> , a novel gene required for <i>Salmonella</i> Typhimurium invasion of host cells. <i>Scientific Reports</i> , 2018, 8, 4841.	1.6	9
10	The Hcp-like protein HilE inhibits homodimerization and DNA binding of the virulence-associated transcriptional regulator HilD in <i>Salmonella</i> . <i>Journal of Biological Chemistry</i> , 2018, 293, 6578-6592.	1.6	14
11	Regulatory Evolution Drives Evasion of Host Inflammasomes by <i>Salmonella</i> Typhimurium. <i>Cell Reports</i> , 2018, 25, 825-832.e5.	2.9	22
12	Ultrastructural and physiological changes induced by different stress conditions on the human parasite <i>Trypanosoma cruzi</i> . <i>Cell Stress and Chaperones</i> , 2017, 22, 15-27.	1.2	10
13	The transcriptional regulator SsrB is involved in a molecular switch controlling virulence lifestyles of <i>Salmonella</i> . <i>PLoS Pathogens</i> , 2017, 13, e1006497.	2.1	50
14	In silico clustering of <i>Salmonella</i> global gene expression data reveals novel genes co-regulated with the SPI-1 virulence genes through HilD. <i>Scientific Reports</i> , 2016, 6, 37858.	1.6	19
15	The global regulatory system Csr senses glucose through the phosphoenolpyruvate: carbohydrate phosphotransferase system. <i>Molecular Microbiology</i> , 2016, 99, 623-626.	1.2	8
16	A multi-drug resistant <i>Salmonella</i> Typhimurium ST213 human-invasive strain (33676) containing the <i>bla</i> CMY-2 gene on an IncF plasmid is attenuated for virulence in BALB/c mice. <i>BMC Microbiology</i> , 2016, 16, 18.	1.3	13
17	The two-component system CpxR/A represses the expression of <i>Salmonella</i> virulence genes by affecting the stability of the transcriptional regulator HilD. <i>Frontiers in Microbiology</i> , 2015, 6, 807.	1.5	40
18	The role of small heat shock proteins in parasites. <i>Cell Stress and Chaperones</i> , 2015, 20, 767-780.	1.2	53

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19	Analytical Validation of Quantitative Real-Time PCR Methods for Quantification of <i>Trypanosoma cruzi</i> DNA in Blood Samples from Chagas Disease Patients. <i>Journal of Molecular Diagnostics</i> , 2015, 17, 605-615.	1.2	153
20	HilD Induces Expression of <i>Salmonella</i> Pathogenicity Island 2 Genes by Displacing the Global Negative Regulator H-NS from <i>ssrAB</i> . <i>Journal of Bacteriology</i> , 2014, 196, 3746-3755.	1.0	35
21	In Silico Identification and Experimental Characterization of Regulatory Elements Controlling the Expression of the <i>Salmonella</i> <i>csrB</i> and <i>csrC</i> Genes. <i>Journal of Bacteriology</i> , 2014, 196, 325-336.	1.0	34
22	Seroprevalence and major antigens recognized by sera from <i>Trypanosoma cruzi</i> -infected dogs from Jalisco, MÃ©xico. <i>Revista Argentina De Microbiologia</i> , 2014, 46, 85-90.	0.4	8
23	Proteomic Analysis of <i>Trypanosoma cruzi</i> Epimastigotes Subjected to Heat Shock. <i>Journal of Biomedicine and Biotechnology</i> , 2012, 2012, 1-9.	3.0	23
24	Changes in cysts' nuclear chromatin resulting after experimental manipulation of <i>Taenia crassiceps</i> mice infections: Biological implications. <i>Experimental Parasitology</i> , 2012, 130, 423-429.	0.5	2
25	Integration of a complex regulatory cascade involving the SirA/BarA and Csr global regulatory systems that controls expression of the <i>Salmonella</i> SPI-1 and SPI-2 virulence regulons through HilD. <i>Molecular Microbiology</i> , 2011, 80, 1637-1656.	1.2	138
26	<i>Trypanosoma cruzi</i> SHSP16: Characterization of an α -crystallin small heat shock protein. <i>Experimental Parasitology</i> , 2009, 123, 182-189.	0.5	26
27	HilD-mediated transcriptional cross-talk between SPI-1 and SPI-2. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 14591-14596.	3.3	151