RocÃ-o Canals

List of Publications by Year in descending order

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Version: 2024-02-01

623734 752698 1,074 26 14 20 citations g-index h-index papers 30 30 30 1604 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	An Infection-Relevant Transcriptomic Compendium for Salmonella enterica Serovar Typhimurium. Cell Host and Microbe, 2013, 14, 683-695.	11.0	427
2	Salmonella Persistence in Tomatoes Requires a Distinct Set of Metabolic Functions Identified by Transposon Insertion Sequencing. Applied and Environmental Microbiology, 2017, 83, .	3.1	78
3	Role of a single noncoding nucleotide in the evolution of an epidemic African clade of <i>Salmonella</i> . Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E2614-E2623.	7.1	75
4	Characterization of the Prophage Repertoire of African Salmonella Typhimurium ST313 Reveals High Levels of Spontaneous Induction of Novel Phage BTP1. Frontiers in Microbiology, 2017, 8, 235.	3.5	73
5	Stepwise evolution of Salmonella Typhimurium ST313 causing bloodstream infection in Africa. Nature Microbiology, 2021, 6, 327-338.	13.3	68
6	Adding function to the genome of African Salmonella Typhimurium ST313 strain D23580. PLoS Biology, 2019, 17, e3000059.	5.6	62
7	A <scp>BTP</scp> 1 prophage gene present in invasive nonâ€typhoidal <scp><i>S</i>S>i>scp><i>almonella</i> determines composition and length of the <scp>O</scp>â€antigen of the lipopolysaccharide. Molecular Microbiology, 2015, 96, 263-275.</scp>	2.5	57
8	Structural Studies of the Oâ€Chain Polysaccharide from <i>Plesiomonas shigelloides</i> Strain 302–73 (Serotype O1). European Journal of Organic Chemistry, 2008, 2008, 3149-3155.	2.4	26
9	High-throughput comparison of gene fitness among related bacteria. BMC Genomics, 2012, 13, 212.	2.8	26
10	A window into lysogeny: revealing temperate phage biology with transcriptomics. Microbial Genomics, 2020, 6, .	2.0	25
11	The fitness landscape of the African Salmonella Typhimurium ST313 strain D23580 reveals unique properties of the pBT1 plasmid. PLoS Pathogens, 2019, 15, e1007948.	4.7	20
12	Evasion of MAIT cell recognition by the African <i>Salmonella</i> Typhimurium ST313 pathovar that causes invasive disease. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 20717-20728.	7.1	20
13	The use of chicken and insect infection models to assess the virulence of African Salmonella Typhimurium ST313. PLoS Neglected Tropical Diseases, 2019, 13, e0007540.	3.0	19
14	Molecular and Chemical Analysis of the Lipopolysaccharide from Aeromonas hydrophila Strain AH-1 (Serotype O11). Marine Drugs, 2015, 13, 2233-2249.	4.6	18
15	Functional Genomics of the Aeromonas salmonicida Lipopolysaccharide O-Antigen and A-Layer from Typical and Atypical Strains. Marine Drugs, 2015, 13, 3791-3808.	4.6	16
16	The diversity, evolution and ecology of Salmonella in venomous snakes. PLoS Neglected Tropical Diseases, 2019, 13, e0007169.	3.0	16
17	Genetic variation in the MacAB-TolC efflux pump influences pathogenesis of invasive Salmonella isolates from Africa. PLoS Pathogens, 2020, 16, e1008763.	4.7	15
18	Hha has a defined regulatory role that is not dependent upon H-NS or StpA. Frontiers in Microbiology, 2015, 6, 773.	3.5	12

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19	The FUN of identifying gene function in bacterial pathogens; insights from Salmonella functional genomics. Current Opinion in Microbiology, 2013, 16, 643-651.	5.1	8
20	A Cronobacter turicensis O1 Antigen-Specific Monoclonal Antibody Inhibits Bacterial Motility and Entry into Epithelial Cells. Infection and Immunity, 2015, 83, 876-887.	2.2	8
21	Genomics of Salmonella Species. , 2011, , 171-235.		1
22	Global gene expression profiling of a virulent Klebsiella pneumoniae strain during pulmonary infection. Access Microbiology, 2019, 1 , .	0.5	0
23	Identification of genes that contribute to fitness of African and Global clades of Salmonella Enteritidis during infection of macrophages. Access Microbiology, 2020, 2, .	0.5	0
24	Title is missing!. , 2019, 15, e1007948.		0
25	Title is missing!. , 2019, 15, e1007948.		0
26	Title is missing!. , 2019, 15, e1007948.		0