

Salvador Almagro-Moreno

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2931880/publications.pdf>

Version: 2024-02-01

31
papers

1,328
citations

471509

17
h-index

477307

29
g-index

34
all docs

34
docs citations

34
times ranked

1704
citing authors

#	ARTICLE	IF	CITATIONS
1	Sialic Acid Catabolism Confers a Competitive Advantage to Pathogenic <i>Vibrio cholerae</i> in the Mouse Intestine. <i>Infection and Immunity</i> , 2009, 77, 3807-3816.	2.2	173
2	Insights into the evolution of sialic acid catabolism among bacteria. <i>BMC Evolutionary Biology</i> , 2009, 9, 118.	3.2	163
3	Intestinal Colonization Dynamics of <i>Vibrio cholerae</i> . <i>PLoS Pathogens</i> , 2015, 11, e1004787.	4.7	140
4	Genomic islands are dynamic, ancient integrative elements in bacterial evolution. <i>Trends in Microbiology</i> , 2009, 17, 47-53.	7.7	137
5	The genomic code: inferring Vibrionaceae niche specialization. <i>Nature Reviews Microbiology</i> , 2006, 4, 697-704.	28.6	115
6	Bacterial catabolism of nonulosonic (sialic) acid and fitness in the gut. <i>Gut Microbes</i> , 2010, 1, 45-50.	9.8	54
7	Cholera: Environmental Reservoirs and Impact on Disease Transmission. <i>Microbiology Spectrum</i> , 2013, 1, .	3.0	53
8	Environmental Role of Pathogenic Traits in <i>Vibrio cholerae</i> . <i>Journal of Bacteriology</i> , 2018, 200, e00795-17.	2.2	51
9	Proteolysis of Virulence Regulator ToxR Is Associated with Entry of <i>Vibrio cholerae</i> into a Dormant State. <i>PLoS Genetics</i> , 2015, 11, e1005145.	3.5	49
10	Origins of pandemic <i>Vibrio cholerae</i> from environmental gene pools. <i>Nature Microbiology</i> , 2017, 2, 16240.	13.3	42
11	Bile salts and alkaline p _H reciprocally modulate the interaction between the periplasmic domains of <i>V</i> and <i>T</i> ox _R and <i>T</i> ox _S . <i>Molecular Microbiology</i> , 2017, 105, 258-272.	2.5	41
12	Evolutionary Model of Cluster Divergence of the Emergent Marine Pathogen <i>Vibrio vulnificus</i> : From Genotype to Ecotype. <i>MBio</i> , 2019, 10, .	4.1	41
13	Ecology and Genetic Structure of a Northern Temperate <i>Vibrio cholerae</i> Population Related to Toxicogenic Isolates. <i>Applied and Environmental Microbiology</i> , 2011, 77, 7568-7575.	3.1	32
14	Role of <i>ToxS</i> in the proteolytic cascade of virulence regulator <i>ToxR</i> in <i>V</i> . <i>Molecular Microbiology</i> , 2015, 98, 963-976.	2.5	32
15	Catabolism of mucus components influences motility of <i>Vibrio cholerae</i> in the presence of environmental reservoirs. <i>PLoS ONE</i> , 2018, 13, e0201383.	2.5	28
16	Excision dynamics of <i>Vibrio</i> pathogenicity island-2 from <i>Vibrio cholerae</i> : role of a recombination directionality factor VefA. <i>BMC Microbiology</i> , 2010, 10, 306.	3.3	27
17	Host-Like Carbohydrates Promote Bloodstream Survival of <i>Vibrio vulnificus</i> <i>In Vivo</i> . <i>Infection and Immunity</i> , 2015, 83, 3126-3136.	2.2	19
18	Molecular mechanisms and drivers of pathogen emergence. <i>Trends in Microbiology</i> , 2022, 30, 898-911.	7.7	19

#	ARTICLE	IF	CITATIONS
19	Dichotomy in the evolution of pathogenicity island and bacteriophage encoded integrases from pathogenic <i>Escherichia coli</i> strains. <i>Infection, Genetics and Evolution</i> , 2011, 11, 423-436.	2.3	17
20	Direct transmission via households informs models of disease and intervention dynamics in cholera. <i>PLoS ONE</i> , 2020, 15, e0229837.	2.5	14
21	Cholera dynamics: lessons from an epidemic. <i>Journal of Medical Microbiology</i> , 2021, 70, .	1.8	14
22	Ecological diversification reveals routes of pathogen emergence in endemic <i>Vibrio vulnificus</i> populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	14
23	Dysbiosis in marine aquaculture revealed through microbiome analysis: reverse ecology for environmental sustainability. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	2.7	10
24	Synergistic role of abiotic factors driving viable but non-culturable <i>Vibrio cholerae</i> . <i>Environmental Microbiology Reports</i> , 2020, 12, 454-465.	2.4	10
25	Hepatitis C virus modelled as an indirectly transmitted infection highlights the centrality of injection drug equipment in disease dynamics. <i>Journal of the Royal Society Interface</i> , 2019, 16, 20190334.	3.4	9
26	Cholera: Environmental Reservoirs and Impact on Disease Transmission. , 0, , 149-165.		9
27	An atomic force microscopy method for the detection of binding forces between bacteria and a lipid bilayer containing higher order gangliosides. <i>Journal of Microbiological Methods</i> , 2011, 84, 352-354.	1.6	5
28	JMM Profile: <i>Vibrio cholerae</i> : an opportunist of human crises. <i>Journal of Medical Microbiology</i> , 2021, 70, .	1.8	5
29	Improved Method for Transformation of <i>Vibrio vulnificus</i> by Electroporation. <i>Current Protocols in Microbiology</i> , 2020, 58, e106.	6.5	2
30	How Genomics Has Shaped Our Understanding of the Evolution and Emergence of Pathogenic <i>Vibrio cholerae</i> . , 0, , 85-99.		2
31	Thanks, but no thanks: Cholera pathogen keeps incoming DNA at bay. <i>Cell Host and Microbe</i> , 2022, 30, 877-879.	11.0	1