

Simon Le Hello

List of Publications by Year in descending order

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76
papers

3,434
citations

159585

30
h-index

155660

55
g-index

79
all docs

79
docs citations

79
times ranked

4638
citing authors

#	ARTICLE	IF	CITATIONS
1	Antimicrobial resistance genes, virulence markers and prophage sequences in <i>Salmonella enterica</i> serovar Enteritidis isolated in Tunisia using whole genome sequencing. <i>Current Research in Microbial Sciences</i> , 2022, 3, 100151.	2.3	1
2	Intrauterine infection caused by nontyphoidal <i>Salmonella</i> : a literature review. <i>Journal of Maternal-Fetal and Neonatal Medicine</i> , 2021, 34, 4000-4004.	1.5	6
3	Comparison of conventional molecular and whole-genome sequencing methods for subtyping <i>Salmonella enterica</i> serovar Enteritidis strains from Tunisia. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 597-606.	2.9	8
4	Whole Genome Analysis Detects the Emergence of a Single <i>Salmonella enterica</i> Serovar Chester Clone in Japan's Kanto Region. <i>Frontiers in Microbiology</i> , 2021, 12, 705679.	3.5	1
5	Detection of Colistin Resistance in <i>Salmonella enterica</i> Using MALDIxin Test on the Routine MALDI Biotyper Sirius Mass Spectrometer. <i>Frontiers in Microbiology</i> , 2020, 11, 1141.	3.5	12
6	Genetic diversity of clinical <i>Salmonella enterica</i> serovar Typhimurium in a university hospital of south Tunisia, 2000–2013. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104436.	2.3	3
7	Reptiles in Guadeloupe (French West Indies) are a reservoir of major human <i>Salmonella enterica</i> serovars. <i>PLoS ONE</i> , 2019, 14, e0220145.	2.5	10
8	A Simple and Robust Statistical Method to Define Genetic Relatedness of Samples Related to Outbreaks at the Genomic Scale – Application to Retrospective <i>Salmonella</i> Foodborne Outbreak Investigations. <i>Frontiers in Microbiology</i> , 2019, 10, 2413.	3.5	17
9	An international outbreak of <i>Salmonella enterica</i> serotype Enteritidis linked to eggs from Poland: a microbiological and epidemiological study. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 778-786.	9.1	81
10	Emergence of phylogenetically diverse and fluoroquinolone resistant <i>Salmonella</i> Enteritidis as a cause of invasive nontyphoidal <i>Salmonella</i> disease in Ghana. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007485.	3.0	30
11	Serotype distribution and antimicrobial resistance of human <i>Salmonella enterica</i> in Bangui, Central African Republic, from 2004 to 2013. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007917.	3.0	17
12	Evaluation of temocillin for phenotypic carbapenemase screening of <i>Escherichia coli</i> and <i>Salmonella enterica</i> isolates in relation to the presence of genes encoding ESBLs and carbapenemase production. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 639-644.	3.0	5
13	CTX-M-55-type ESBL-producing <i>Salmonella enterica</i> are emerging among retail meats in Phnom Penh, Cambodia. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 342-348.	3.0	58
14	Global phylogenomics of multidrug-resistant <i>Salmonella enterica</i> serotype Kentucky ST198. <i>Microbial Genomics</i> , 2019, 5, .	2.0	69
15	Disentangling a complex nationwide <i>Salmonella</i> Dublin outbreak associated with raw-milk cheese consumption, France, 2015 to 2016. <i>Eurosurveillance</i> , 2019, 24, .	7.0	41
16	Early transmissible ampicillin resistance in zoonotic <i>Salmonella enterica</i> serotype Typhimurium in the late 1950s: a retrospective, whole-genome sequencing study. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 207-214.	9.1	33
17	In-Host Adaptation of <i>Salmonella enterica</i> Serotype Dublin during Prosthetic Hip Joint Infection. <i>Emerging Infectious Diseases</i> , 2018, 24, 2360-2363.	4.3	6
18	Occurrence of Bacterial Pathogens and Human Noroviruses in Shellfish-Harvesting Areas and Their Catchments in France. <i>Frontiers in Microbiology</i> , 2018, 9, 2443.	3.5	38

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19	Ongoing nationwide outbreak of Salmonella Agona associated with internationally distributed infant milk products, France, December 2017. <i>Eurosurveillance</i> , 2018, 23, .	7.0	59
20	In-Host Adaptation of <i>Salmonella enterica</i> Serotype Dublin during Prosthetic Hip Joint Infection. <i>Emerging Infectious Diseases</i> , 2018, 24, 2360-2363.	4.3	0
21	Global and regional dissemination and evolution of <i>Burkholderia pseudomallei</i> . <i>Nature Microbiology</i> , 2017, 2, 16263.	13.3	124
22	MLVA for <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Dublin: Development of a Method Suitable for Inter-Laboratory Surveillance and Application in the Context of a Raw Milk Cheese Outbreak in France in 2012. <i>Frontiers in Microbiology</i> , 2017, 08, 295.	3.5	17
23	The invasome of <i>Salmonella</i> Dublin as revealed by whole genome sequencing. <i>BMC Infectious Diseases</i> , 2017, 17, 544.	2.9	45
24	The clinical and microbiological characteristics of enteric fever in Cambodia, 2008-2015. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005964.	3.0	40
25	Multinational outbreak of travel-related <i>Salmonella</i> Chester infections in Europe, summers 2014 and 2015. <i>Eurosurveillance</i> , 2017, 22, .	7.0	31
26	Multi-laboratory validation study of multilocus variable-number tandem repeat analysis (MLVA) for <i>Salmonella enterica</i> serovar Enteritidis, 2015. <i>Eurosurveillance</i> , 2017, 22, .	7.0	12
27	Phylogenetic structure of European <i>Salmonella</i> Enteritidis outbreak correlates with national and international egg distribution network. <i>Microbial Genomics</i> , 2016, 2, e000070.	2.0	67
28	Genome-scale rates of evolutionary change in bacteria. <i>Microbial Genomics</i> , 2016, 2, e000094.	2.0	224
29	Molecular Surveillance Identifies Multiple Transmissions of Typhoid in West Africa. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004781.	3.0	46
30	Genomic analysis of <i>Salmonella enterica</i> serotype Paratyphi A during an outbreak in Cambodia, 2013–2015. <i>Microbial Genomics</i> , 2016, 2, e000092.	2.0	24
31	Outbreak of <i>Salmonella</i> Enteritidis linked to the consumption of frozen beefburgers received from a food bank and originating from Poland: northern France, December 2014 to April 2015. <i>Eurosurveillance</i> , 2016, 21, .	7.0	7
32	Global Genomic Epidemiology of <i>Salmonella enterica</i> Serovar Typhimurium DT104. <i>Applied and Environmental Microbiology</i> , 2016, 82, 2516-2526.	3.1	105
33	Distinct <i>Salmonella</i> Enteritidis lineages associated with enterocolitis in high-income settings and invasive disease in low-income settings. <i>Nature Genetics</i> , 2016, 48, 1211-1217.	21.4	191
34	Whatâ€™s in a Name? Species-Wide Whole-Genome Sequencing Resolves Invasive and Noninvasive Lineages of <i>Salmonella enterica</i> Serotype Paratyphi B. <i>MBio</i> , 2016, 7, .	4.1	29
35	Global phylogeography and evolutionary history of <i>Shigella dysenteriae</i> type 1. <i>Nature Microbiology</i> , 2016, 1, 16027.	13.3	65
36	A Phylogenetic and Phenotypic Analysis of <i>Salmonella enterica</i> Serovar Weltevreden, an Emerging Agent of Diarrheal Disease in Tropical Regions. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004446.	3.0	59

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37	Investigating Salmonella Eko from Various Sources in Nigeria by Whole Genome Sequencing to Identify the Source of Human Infections. PLoS ONE, 2016, 11, e0156212.	2.5	8
38	Community Incidence of Campylobacteriosis and Nontyphoidal Salmonellosis, France, 2008–2013. Foodborne Pathogens and Disease, 2015, 12, 664-669.	1.8	23
39	Carbapenemase-producing <i>Salmonella enterica</i> serotype Kentucky ST198, North Africa: Table 1. Journal of Antimicrobial Chemotherapy, 2015, 70, dkv276.	3.0	10
40	High-Throughput CRISPR Typing of Mycobacterium tuberculosis Complex and Salmonella enterica Serotype Typhimurium. Methods in Molecular Biology, 2015, 1311, 91-109.	0.9	15
41	Multidrug-Resistant <i>Salmonella enterica</i> Serotype Typhi, Gulf of Guinea Region, Africa. Emerging Infectious Diseases, 2015, 21, 655-659.	4.3	16
42	Salmonella enterica Serotype Enteritidis in French Polynesia, South Pacific, 2008–2013. Emerging Infectious Diseases, 2015, 21, 1045-1048.	4.3	6
43	Enteric Bacterial Pathogens in Children with Diarrhea in Niger: Diversity and Antimicrobial Resistance. PLoS ONE, 2015, 10, e0120275.	2.5	66
44	Lack of efflux mediated quinolone resistance in Salmonella enterica serovars Typhi and Paratyphi A. Frontiers in Microbiology, 2014, 5, 12.	3.5	15
45	CRISPR Is an Optimal Target for the Design of Specific PCR Assays for Salmonella enterica Serotypes Typhi and Paratyphi A. PLoS Neglected Tropical Diseases, 2014, 8, e2671.	3.0	25
46	Invasive Salmonella enterica Serotype Typhimurium Infections, Democratic Republic of the Congo, 2007–2011. Emerging Infectious Diseases, 2014, 20, 701-704.	4.3	20
47	Genomic Definition of Hypervirulent and Multidrug-Resistant <i>Klebsiella pneumoniae</i> Clonal Groups. Emerging Infectious Diseases, 2014, 20, 1812-1820.	4.3	409
48	Repeated isolation of Salmonella enterica Gerverdhan, a very rare serovar, from Danish poultry surveillance samples. Veterinary Microbiology, 2014, 174, 596-599.	1.9	2
49	Targeting relaxase genes for classification of the predominant plasmids in Enterobacteriaceae. International Journal of Medical Microbiology, 2014, 304, 236-242.	3.6	43
50	Prevalence and characterization of extended-spectrum β -lactamase-producing clinical Salmonella enterica isolates in Dakar, Senegal, from 1999 to 2009. Clinical Microbiology and Infection, 2014, 20, O109-O116.	6.0	46
51	Salmonella serotypes in reptiles and humans, French Guiana. Veterinary Microbiology, 2014, 170, 167-171.	1.9	24
52	The French human Salmonella surveillance system: evaluation of timeliness of laboratory reporting and factors associated with delays, 2007 to 2011. Eurosurveillance, 2014, 19, .	7.0	21
53	Highly drug-resistant Salmonella enterica serotype Kentucky ST198-X1: a microbiological study. Lancet Infectious Diseases, The, 2013, 13, 672-679.	9.1	149
54	Genetic lineages of Salmonella enterica serovar Kentucky spreading in pet reptiles. Veterinary Microbiology, 2013, 166, 686-689.	1.9	24

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55	ramR mutations affecting fluoroquinolone susceptibility in epidemic multidrug-resistant <i>Salmonella enterica</i> serovar Kentucky ST198. <i>Frontiers in Microbiology</i> , 2013, 4, 213.	3.5	26
56	The global establishment of a highly-fluoroquinolone resistant <i>Salmonella enterica</i> serotype Kentucky ST198 strain. <i>Frontiers in Microbiology</i> , 2013, 4, 395.	3.5	114
57	Unusual increase in reported cases of Paratyphoid A fever among travellers returning from Cambodia, January to September 2013. <i>Eurosurveillance</i> , 2013, 18, .	7.0	12
58	Use of multilocus variable-number tandem repeat analysis (MLVA) in eight European countries, 2012. <i>Eurosurveillance</i> , 2013, 18, 20385.	7.0	63
59	Characterization of Isolates of <i>Salmonella enterica</i> Serovar Stanley, a Serovar Endemic to Asia and Associated with Travel. <i>Journal of Clinical Microbiology</i> , 2012, 50, 709-720.	3.9	32
60	Early Strains of Multidrug-Resistant <i>Salmonella enterica</i> Serovar Kentucky Sequence Type 198 from Southeast Asia Harbor <i>Salmonella</i> Genomic Island 1-J Variants with a Novel Insertion Sequence. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 5096-5102.	3.2	36
61	CRISPR Typing and Subtyping for Improved Laboratory Surveillance of <i>Salmonella</i> Infections. <i>PLoS ONE</i> , 2012, 7, e36995.	2.5	198
62	Foodborne Outbreak and Nonmotile <i>Salmonella enterica</i> Variant, France. <i>Emerging Infectious Diseases</i> , 2012, 18, 132-134.	4.3	15
63	International Spread of an Epidemic Population of <i>Salmonella enterica</i> Serotype Kentucky ST198 Resistant to Ciprofloxacin. <i>Journal of Infectious Diseases</i> , 2011, 204, 675-684.	4.0	226
64	Clinical and Microbial Characteristics of Invasive <i>Streptococcus pyogenes</i> Disease in New Caledonia, a Region in Oceania with a High Incidence of Acute Rheumatic Fever. <i>Journal of Clinical Microbiology</i> , 2010, 48, 1993-1993.	3.9	2
65	Invasive Serotype 1 <i>Streptococcus pneumoniae</i> Outbreaks in the South Pacific from 2000 to 2007. <i>Journal of Clinical Microbiology</i> , 2010, 48, 2968-2971.	3.9	21
66	Clinical and Microbial Characteristics of Invasive <i>Streptococcus pyogenes</i> Disease in New Caledonia, a Region in Oceania with a High Incidence of Acute Rheumatic Fever. <i>Journal of Clinical Microbiology</i> , 2010, 48, 526-530.	3.9	35
67	Serological reactivity and bacterial genotypes in <i>Chlamydia trachomatis</i> urogenital infections in Guadeloupe, French West Indies. <i>Sexually Transmitted Infections</i> , 2010, 86, 101-105.	1.9	11
68	Risk factors for carbapenem-resistant <i>Acinetobacter baumannii</i> infections at a tertiary care hospital in New Caledonia, South Pacific. <i>Scandinavian Journal of Infectious Diseases</i> , 2010, 42, 821-826.	1.5	17
69	Ceftazidime-Resistant <i>Salmonella enterica</i> , Morocco. <i>Emerging Infectious Diseases</i> , 2009, 15, 1693b-1695.	4.3	15
70	High HIV Type 1 Prevalence and Wide Genetic Diversity with Dominance of Recombinant Strains But Low Level of Antiretroviral Drug-Resistance Mutations in Untreated Patients in Northeast Gabon, Central Africa. <i>AIDS Research and Human Retroviruses</i> , 2009, 25, 411-418.	1.1	16
71	High prevalence of hepatitis C virus infection and predominance of genotype 4 in rural gabon. <i>Journal of Medical Virology</i> , 2008, 80, 1581-1587.	5.0	33
72	Molecular epidemiology of carbapenem-resistant <i>Acinetobacter baumannii</i> in New Caledonia. <i>Clinical Microbiology and Infection</i> , 2008, 14, 977-981.	6.0	11

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73	Fluoroquinolone resistance in a clinical isolate of <i>Streptococcus pneumoniae</i> in the South Pacific. <i>International Journal of Antimicrobial Agents</i> , 2008, 32, 91-92.	2.5	1
74	Real-Time PCR Detection of <i>gyrA</i> and <i>parC</i> Mutations in <i>Streptococcus pneumoniae</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 4155-4158.	3.2	8
75	Melioidosis in New Caledonia. <i>Emerging Infectious Diseases</i> , 2005, 11, 1607-1609.	4.3	23
76	Seroepidemiology of <i>Helicobacter pylori</i> infection in Guadeloupe. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2002, 96, 517-519.	1.8	9