

# John R Wain

## List of Publications by Year in descending order

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168  
papers

15,904  
citations

20036

63  
h-index

21239

119  
g-index

179  
all docs

179  
docs citations

179  
times ranked

16544  
citing authors

#	ARTICLE	IF	CITATIONS
1	OUP accepted manuscript. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2022, , .	0.7	2
2	Microbiomes of Urine and the Prostate Are Linked to Human Prostate Cancer Risk Groups. European Urology Oncology, 2022, 5, 412-419.	2.6	21
3	The ecology and antimicrobial resistance of Staphylococci colonising neonates. Access Microbiology, 2022, 4, .	0.2	0
4	Genetic characterization of Salmonella Infantis from South Africa, 2004â€“2016. Access Microbiology, 2022, 4, .	0.2	1
5	Salmonella nomenclature in the genomic era: a time for change. Scientific Reports, 2021, 11, 7494.	1.6	30
6	Bactericidal activities and post-antibiotic effects of ofloxacin and ceftriaxone against drug-resistant Salmonella enterica serovar Typhi. Journal of Antimicrobial Chemotherapy, 2021, 76, 2606-2609.	1.3	1
7	Characterization of a pESI-like plasmid and analysis of multidrug-resistant Salmonella enterica Infantis isolates in England and Wales. Microbial Genomics, 2021, 7, .	1.0	14
8	Massively parallel transposon mutagenesis identifies temporally essential genes for biofilm formation in Escherichia coli. Microbial Genomics, 2021, 7, .	1.0	10
9	Emergence of Resistance to Fluoroquinolones and Third-Generation Cephalosporins in Salmonella Typhi in Lahore, Pakistan. Microorganisms, 2020, 8, 1336.	1.6	35
10	Identification and characterisation of enteroaggregative Escherichia coli subtypes associated with human disease. Scientific Reports, 2020, 10, 7475.	1.6	23
11	A whole-genome screen identifies Salmonella enterica serovar Typhi genes involved in fluoroquinolone susceptibility. Journal of Antimicrobial Chemotherapy, 2020, 75, 2516-2525.	1.3	11
12	Concomitant Bacteremia in Adults With Severe Falciparum Malaria. Clinical Infectious Diseases, 2020, 71, e465-e470.	2.9	22
13	Genetic Markers in S. Paratyphi C Reveal Primary Adaptation to Pigs. Microorganisms, 2020, 8, 657.	1.6	5
14	High-density transposon libraries utilising outward-oriented promoters identify mechanisms of action and resistance to antimicrobials. FEMS Microbiology Letters, 2020, 367, .	0.7	4
15	Nanopore metagenomics enables rapid clinical diagnosis of bacterial lower respiratory infection. Nature Biotechnology, 2019, 37, 783-792.	9.4	396
16	Safety and efficacy of 2% chlorhexidine gluconate aqueous versus 2% chlorhexidine gluconate in 70% isopropyl alcohol for skin disinfection prior to percutaneous central venous catheter insertion in preterm neonates: the ARCTIC randomised-controlled feasibility trial protocol. BMJ Open, 2019, 9, e028022.	0.8	12
17	Ceftriaxone-resistant Salmonella Typhi carries an IncI1-ST31 plasmid encoding CTX-M-15. Journal of Medical Microbiology, 2018, 67, 620-627.	0.7	25
18	Genomic Diversity in Salmonella enterica. , 2017, , 91-107.		3

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19	Optimisation of 16S rRNA gut microbiota profiling of extremely low birth weight infants. BMC Genomics, 2017, 18, 841.	1.2	47
20	Salmonella. , 2017, , 425-433.		8
21	Clonality, virulence and antimicrobial resistance of enteroaggregative Escherichia coli from Mirzapur, Bangladesh. Journal of Medical Microbiology, 2017, 66, 1429-1435.	0.7	19
22	Development and evaluation of a multiple-locus variable-number tandem-repeats analysis assay for subtyping Salmonella Typhi strains from sub-Saharan Africa. Journal of Medical Microbiology, 2017, 66, 937-945.	0.7	9
23	Microevolution of Monophasic <i>Salmonella</i> Typhimurium during Epidemic, United Kingdom, 2005–2010. Emerging Infectious Diseases, 2016, 22, 617-624.	2.0	158
24	What's in a Name? Species-Wide Whole-Genome Sequencing Resolves Invasive and Noninvasive Lineages of Salmonella enterica Serotype Paratyphi B. MBio, 2016, 7, .	1.8	29
25	Detection of mutations in gyrB using denaturing high performance liquid chromatography (DHPLC) among Salmonella enterica serovar Typhi and Paratyphi A. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2016, 110, 684-689.	0.7	4
26	A novel broadly applicable PCR-RFLP method for rapid identification and subtyping of H58 Salmonella Typhi. Journal of Microbiological Methods, 2016, 127, 219-223.	0.7	16
27	Diversity of STs, plasmids and ESBL genes among <i>Escherichia coli</i> from humans, animals and food in Germany, the Netherlands and the UK. Journal of Antimicrobial Chemotherapy, 2016, 71, 1178-1182.	1.3	110
28	Rapid Identification of Major Escherichia coli Sequence Types Causing Urinary Tract and Bloodstream Infections. Journal of Clinical Microbiology, 2015, 53, 160-166.	1.8	121
29	Whole-Genome Sequencing for National Surveillance of Shiga Toxin–Producing <i>Escherichia coli</i> O157. Clinical Infectious Diseases, 2015, 61, 305-312.	2.9	181
30	MinION nanopore sequencing identifies the position and structure of a bacterial antibiotic resistance island. Nature Biotechnology, 2015, 33, 296-300.	9.4	404
31	Typhoid fever. Lancet, The, 2015, 385, 1136-1145.	6.3	265
32	Applying phylogenomics to understand the emergence of Shiga-toxin-producing Escherichia coli O157:H7 strains causing severe human disease in the UK. Microbial Genomics, 2015, 1, e000029.	1.0	105
33	Epidemiological Evidence That Garden Birds Are a Source of Human Salmonellosis in England and Wales. PLoS ONE, 2014, 9, e88968.	1.1	67
34	Characterization of non-classical quinolone resistance in Salmonella enterica serovar Typhi: Report of a novel mutation in gyrB gene and diagnostic challenges. Biomolecular Detection and Quantification, 2014, 2, 30-34.	7.0	7
35	Evidence of Evolving Extraintestinal Enteroaggregative Escherichia coli ST38 Clone. Emerging Infectious Diseases, 2014, 20, 1935-1937.	2.0	51
36	Transient Darwinian selection in <i>Salmonella enterica</i> serovar Paratyphi A during 450 years of global spread of enteric fever. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 12199-12204.	3.3	122

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37	In vitro activity of rifaximin against clinical isolates of Escherichia coli and other enteropathogenic bacteria isolated from travellers returning to the UK. International Journal of Antimicrobial Agents, 2014, 43, 431-437.	1.1	31
38	Risk factors for the development of severe typhoid fever in Vietnam. BMC Infectious Diseases, 2014, 14, 73.	1.3	39
39	Salmonella enterica subspecies II infections in England and Wales – the use of multilocus sequence typing to assist serovar identification. Journal of Medical Microbiology, 2014, 63, 831-834.	0.7	20
40	Quantification of IgM molecular response by droplet digital PCR as a potential tool for the early diagnosis of sepsis. Critical Care, 2014, 18, 433.	2.5	12
41	Evaluating the Use of Multilocus Variable Number Tandem Repeat Analysis of Shiga Toxin-Producing Escherichia coli O157 as a Routine Public Health Tool in England. PLoS ONE, 2014, 9, e85901.	1.1	20
42	An Investigation of the Diversity of Strains of Enteraggregative Escherichia coli Isolated from Cases Associated with a Large Multi-Pathogen Foodborne Outbreak in the UK. PLoS ONE, 2014, 9, e98103.	1.1	41
43	Enteraggregative Escherichia coli Have Evolved Independently as Distinct Complexes within the E. coli Population with Varying Ability to Cause Disease. PLoS ONE, 2014, 9, e112967.	1.1	17
44	Universal extraction method for gastrointestinal pathogens. Journal of Medical Microbiology, 2013, 62, 1535-1539.	0.7	20
45	Immunological monitoring to prevent and treat sepsis. Critical Care, 2013, 17, 109.	2.5	21
46	Public Health Value of Next-Generation DNA Sequencing of Enterohemorrhagic Escherichia coli Isolates from an Outbreak. Journal of Clinical Microbiology, 2013, 51, 232-237.	1.8	79
47	Next-generation sequencing in clinical microbiology. Expert Review of Molecular Diagnostics, 2013, 13, 225-227.	1.5	14
48	Genomic Characterisation of Invasive Non-Typhoidal Salmonella enterica Subspecies enterica Serovar Bovismorbificans Isolates from Malawi. PLoS Neglected Tropical Diseases, 2013, 7, e2557.	1.3	24
49	Diversity among human non-typhoidal salmonellae isolates from Zimbabwe. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2013, 107, 487-492.	0.7	14
50	Harnessing the genome: development of a hierarchical typing scheme for meticillin-resistant Staphylococcus aureus. Journal of Medical Microbiology, 2013, 62, 36-45.	0.7	4
51	A comparison of dense transposon insertion libraries in the Salmonella serovars Typhi and Typhimurium. Nucleic Acids Research, 2013, 41, 4549-4564.	6.5	108
52	Revolutionising Bacteriology to Improve Treatment Outcomes and Antibiotic Stewardship. Infection and Chemotherapy, 2013, 45, 1.	1.0	38
53	Comparative Analysis of ESBL-Positive Escherichia coli Isolates from Animals and Humans from the UK, The Netherlands and Germany. PLoS ONE, 2013, 8, e75392.	1.1	106
54	Development and application of MLVA methods as a tool for inter-laboratory surveillance. Eurosurveillance, 2013, 18, 20565.	3.9	94

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55	Using next generation sequencing to tackle non-typhoidal Salmonella infections. <i>Journal of Infection in Developing Countries</i> , 2013, 7, 001-005.	0.5	25
56	Multilocus Sequence Typing as a Replacement for Serotyping in <i>Salmonella enterica</i> . <i>PLoS Pathogens</i> , 2012, 8, e1002776.	2.1	574
57	Population Genetic Structure of 4,12:a:â” Salmonella enterica Strains from Harbor Porpoises. <i>Applied and Environmental Microbiology</i> , 2012, 78, 8829-8833.	1.4	6
58	The TCA cycle is not required for selection or survival of multidrug-resistant <i>Salmonella</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2012, 67, 589-599.	1.3	14
59	Changes in Causes of Acute Gastroenteritis in the United Kingdom Over 15 Years: Microbiologic Findings From 2 Prospective, Population-Based Studies of Infectious Intestinal Disease. <i>Clinical Infectious Diseases</i> , 2012, 54, 1275-1286.	2.9	145
60	Invasive Salmonellosis in Humans. <i>EcoSal Plus</i> , 2012, 5, .	2.1	17
61	The decline of typhoid and the rise of non-typhoid salmonellae and fungal infections in a changing HIV landscape: bloodstream infection trends over 15 years in southern Vietnam. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2012, 106, 26-34.	0.7	60
62	Intracontinental spread of human invasive <i>Salmonella</i> Typhimurium pathovariants in sub-Saharan Africa. <i>Nature Genetics</i> , 2012, 44, 1215-1221.	9.4	370
63	Improved Multiplex PCR Strategy for Rapid Assignment of the Four Major <i>Escherichia coli</i> Phylogenetic Groups. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3108-3110.	1.8	145
64	Performance comparison of benchtop high-throughput sequencing platforms. <i>Nature Biotechnology</i> , 2012, 30, 434-439.	9.4	1,226
65	The Impact of JIDC. <i>Journal of Infection in Developing Countries</i> , 2012, 6, 542-542.	0.5	0
66	The Influence of Reduced Susceptibility to Fluoroquinolones in <i>Salmonella enterica</i> Serovar Typhi on the Clinical Response to Ofloxacin Therapy. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1163.	1.3	66
67	Rapid evolution of fluoroquinolone-resistant <i>Escherichia coli</i> in Nigeria is temporally associated with fluoroquinolone use. <i>BMC Infectious Diseases</i> , 2011, 11, 312.	1.3	47
68	Foreign Travel and Decreased Ciprofloxacin Susceptibility in <i>Salmonella enterica</i> Infections. <i>Emerging Infectious Diseases</i> , 2011, 17, 123-125.	2.0	20
69	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.	1.9	226
70	Emergence of a Globally Dominant IncHI1 Plasmid Type Associated with Multiple Drug Resistant Typhoid. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1245.	1.3	114
71	Enterohemorrhagic <i>E. coli</i> O104 from an outbreak of HUS in Germany 2011, could it happen again?. <i>Journal of Infection in Developing Countries</i> , 2011, 5, 425-436.	0.5	53
72	Multidrug resistant <i>Salmonella</i> Concord is a major cause of salmonellosis in children in Ethiopia. <i>Journal of Infection in Developing Countries</i> , 2011, 5, 023-033.	0.5	79

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73	Ten years experience of Salmonella infections in Cambridge, UK. <i>Journal of Infection</i> , 2010, 60, 21-25.	1.7	52
74	Multi-Locus Sequence Typing of Enteroaggregative Escherichia coli Isolates from Nigerian Children Uncovers Multiple Lineages. <i>PLoS ONE</i> , 2010, 5, e14093.	1.1	79
75	A Conserved Acetyl Esterase Domain Targets Diverse Bacteriophages to the Vi Capsular Receptor of <i>Salmonella enterica</i> Serovar Typhi. <i>Journal of Bacteriology</i> , 2010, 192, 5746-5754.	1.0	79
76	First Report of <i>Salmonella enterica</i> Serotype Paratyphi A Azithromycin Resistance Leading to Treatment Failure. <i>Journal of Clinical Microbiology</i> , 2010, 48, 4655-4657.	1.8	62
77	Diarrhoeagenic Escherichia coli from routine diagnostic faecal samples in England and Wales. <i>Journal of Medical Microbiology</i> , 2010, 59, 870-872.	0.7	5
78	A multiplex single nucleotide polymorphism typing assay for detecting mutations that result in decreased fluoroquinolone susceptibility in <i>Salmonella enterica</i> serovars Typhi and Paratyphi A. <i>Journal of Antimicrobial Chemotherapy</i> , 2010, 65, 1631-1641.	1.3	36
79	Emergence and Characterization of <i>Salmonella enterica</i> Serovar Typhimurium Phage Type DT191a. <i>Journal of Clinical Microbiology</i> , 2010, 48, 3375-3377.	1.8	9
80	RamA, a Member of the AraC/XylS Family, Influences Both Virulence and Efflux in <i>Salmonella enterica</i> Serovar Typhimurium. <i>Journal of Bacteriology</i> , 2010, 192, 1607-1616.	1.0	101
81	Suitable Disk Antimicrobial Susceptibility Breakpoints Defining <i>Salmonella enterica</i> Serovar Typhi Isolates with Reduced Susceptibility to Fluoroquinolones. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 5201-5208.	1.4	45
82	Comparison of two DNA microarrays for detection of plasmid-mediated antimicrobial resistance and virulence factor genes in clinical isolates of Enterobacteriaceae and non-Enterobacteriaceae. <i>International Journal of Antimicrobial Agents</i> , 2010, 35, 593-598.	1.1	13
83	Simultaneous assay of every <i>Salmonella</i> Typhi gene using one million transposon mutants. <i>Genome Research</i> , 2009, 19, 2308-2316.	2.4	544
84	Report of Neonatal Meningitis Due to <i>Salmonella enterica</i> Serotype Agona and Review of Breast Milk-Associated Neonatal <i>Salmonella</i> Infections. <i>Journal of Clinical Microbiology</i> , 2009, 47, 3045-3049.	1.8	22
85	The Global Consequence of Disruption of the AcrAB-TolC Efflux Pump in <i>Salmonella enterica</i> Includes Reduced Expression of SPI-1 and Other Attributes Required To Infect the Host. <i>Journal of Bacteriology</i> , 2009, 191, 4276-4285.	1.0	107
86	Exposure of Escherichia coli and Salmonella enterica serovar Typhimurium to triclosan induces a species-specific response, including drug detoxification. <i>Journal of Antimicrobial Chemotherapy</i> , 2009, 64, 973-985.	1.3	65
87	Nontyphoidal <i>Salmonella</i> Serovars Cause Different Degrees of Invasive Disease Globally. <i>Journal of Infectious Diseases</i> , 2009, 199, 602-603.	1.9	23
88	Detecting SNPs and estimating allele frequencies in clonal bacterial populations by sequencing pooled DNA. <i>Bioinformatics</i> , 2009, 25, 2074-2075.	1.8	40
89	Variation in <i>Salmonella enterica</i> Serovar Typhi IncHI1 Plasmids during the Global Spread of Resistant Typhoid Fever. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 716-727.	1.4	81
90	Pseudogene accumulation in the evolutionary histories of <i>Salmonella enterica</i> serovars Paratyphi A and Typhi. <i>BMC Genomics</i> , 2009, 10, 36.	1.2	161

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91	High-throughput sequencing provides insights into genome variation and evolution in Salmonella Typhi. <i>Nature Genetics</i> , 2008, 40, 987-993.	9.4	453
92	Post-genomic challenges for collaborative research in infectious diseases. <i>Nature Reviews Microbiology</i> , 2008, 6, 858-864.	13.6	13
93	Emergence of highly fluoroquinolone-resistant Salmonella enterica serovar Typhi in a community-based fever surveillance from Kolkata, India. <i>International Journal of Antimicrobial Agents</i> , 2008, 31, 387-389.	1.1	24
94	Characterization of the Genomes of a Diverse Collection of <i>Salmonella enterica</i> Serovar Typhimurium Definitive Phage Type 104. <i>Journal of Bacteriology</i> , 2008, 190, 8155-8162.	1.0	51
95	Molecular Characterization of the <i>Salmonella enterica</i> Serovar Typhi Vi-Typing Bacteriophage E1. <i>Journal of Bacteriology</i> , 2008, 190, 2580-2587.	1.0	41
96	a study of typhoid fever in five Asian countries: disease burden and implications for controls. <i>Bulletin of the World Health Organization</i> , 2008, 86, 260-268.	1.5	494
97	The laboratory diagnosis of enteric fever. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 421-5.	0.5	76
98	Typhoid fever in Ethiopia. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 448-53.	0.5	27
99	Specimens and culture media for the laboratory diagnosis of typhoid fever. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 469-74.	0.5	43
100	IncHI plasmids, a dynamic link between resistance and pathogenicity. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 272-8.	0.5	40
101	Antibodies to the Vi capsule of Salmonella Typhi in the serum of typhoid patients and healthy control subjects from a typhoid endemic region. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 308-12.	0.5	20
102	Drug susceptibility testing using molecular techniques can enhance tuberculosis diagnosis. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 40-5.	0.5	8
103	Multidrug-Resistant Salmonella enterica Serovar Paratyphi A Harbors IncHI1 Plasmids Similar to Those Found in Serovar Typhi. <i>Journal of Bacteriology</i> , 2007, 189, 4257-4264.	1.0	80
104	Serodiagnosis of Salmonella enterica serovar Typhi and S. enterica serovars Paratyphi A, B and C human infections. <i>Journal of Medical Microbiology</i> , 2007, 56, 1161-1166.	0.7	27
105	Antimicrobial Drug Resistance of <i>Salmonella enterica</i> Serovar Typhi in Asia and Molecular Mechanism of Reduced Susceptibility to the Fluoroquinolones. <i>Antimicrobial Agents and Chemotherapy</i> , 2007, 51, 4315-4323.	1.4	203
106	Prophage Sequences Defining Hot Spots of Genome Variation in Salmonella enterica Serovar Typhimurium Can Be Used To Discriminate between Field Isolates. <i>Journal of Clinical Microbiology</i> , 2007, 45, 2590-2598.	1.8	59
107	Randomized Controlled Comparison of Ofloxacin, Azithromycin, and an Ofloxacin-Azithromycin Combination for Treatment of Multidrug-Resistant and Nalidixic Acid-Resistant Typhoid Fever. <i>Antimicrobial Agents and Chemotherapy</i> , 2007, 51, 819-825.	1.4	146
108	An H-NS-like Stealth Protein Aids Horizontal DNA Transmission in Bacteria. <i>Science</i> , 2007, 315, 251-252.	6.0	204

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109	Multilocus sequence typing analysis of <i>Shigella flexneri</i> isolates collected in Asian countries. <i>Journal of Medical Microbiology</i> , 2007, 56, 1460-1466.	0.7	33
110	Enteroaggregative <i>Escherichia coli</i> Related to Uropathogenic Clonal Group A. <i>Emerging Infectious Diseases</i> , 2007, 13, 757-760.	2.0	33
111	Cases of typhoid fever imported into England, Scotland and Wales (2000-2003). <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2007, 101, 398-404.	0.7	67
112	A deletion defining a common Asian lineage of <i>Mycobacterium tuberculosis</i> associates with immune subversion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 15594-15598.	3.3	100
113	Evaluation of new-generation serologic tests for the diagnosis of typhoid fever: data from a community-based surveillance in Calcutta, India. <i>Diagnostic Microbiology and Infectious Disease</i> , 2006, 56, 359-365.	0.8	68
114	Antibiotic resistance in <i>Salmonella</i> infections. , 2006, , 25-56.		3
115	Molecular Analysis of Fluoroquinolone-resistant <i>Salmonella</i> Paratyphi A Isolate, India. <i>Emerging Infectious Diseases</i> , 2006, 12, 489-491.	2.0	17
116	The acquisition of full fluoroquinolone resistance in <i>Salmonella</i> Typhi by accumulation of point mutations in the topoisomerase targets. <i>Journal of Antimicrobial Chemotherapy</i> , 2006, 58, 733-740.	1.3	61
117	Deep-Seated Resistance in Relapsed Paratyphoid Fever. <i>Clinical Infectious Diseases</i> , 2006, 42, e92-e94.	2.9	4
118	Isolation of <i>Salmonella enterica</i> subspecies <i>enterica</i> serovar Paratyphi B dT+, or <i>Salmonella</i> Java, from Indonesia and alteration of the d-tartrate fermentation phenotype by disrupting the ORF STM 3356. <i>Journal of Medical Microbiology</i> , 2006, 55, 1661-1665.	0.7	12
119	Fluoroquinolone resistance in <i>Salmonella</i> Typhi. <i>BMJ: British Medical Journal</i> , 2006, 333, 353.3-354.	2.4	24
120	Molecular Analysis of Fluoroquinolone-resistant <i>Salmonella</i> Paratyphi A Isolate, India. <i>Emerging Infectious Diseases</i> , 2006, 12, 489-491.	2.0	13
121	<i>Salmonella</i> Paratyphi A Rates, Asia. <i>Emerging Infectious Diseases</i> , 2005, 11, 1764-1766.	2.0	173
122	Comparative trial of short-course ofloxacin for uncomplicated typhoid fever in Vietnamese children. <i>Annals of Tropical Paediatrics</i> , 2005, 25, 17-22.	1.0	15
123	Effect of Deletion or Overexpression of the 19-Kilodalton Lipoprotein Rv3763 on the Innate Response to <i>Mycobacterium tuberculosis</i> . <i>Infection and Immunity</i> , 2005, 73, 6831-6837.	1.0	37
124	Host Susceptibility and Clinical Outcomes in Toll-Like Receptor 5-Deficient Patients with Typhoid Fever in Vietnam. <i>Journal of Infectious Diseases</i> , 2005, 191, 1068-1071.	1.9	61
125	Use of Paired Serum Samples for Serodiagnosis of Typhoid Fever. <i>Journal of Clinical Microbiology</i> , 2005, 43, 4889-4890.	1.8	39
126	Infection Biology of a Novel $\beta$ -Crystallin of <i>Mycobacterium tuberculosis</i> : Acr2. <i>Journal of Immunology</i> , 2005, 174, 4237-4243.	0.4	64



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127	Vi Antigen Expression in Salmonella enterica Serovar Typhi Clinical Isolates from Pakistan. Journal of Clinical Microbiology, 2005, 43, 1158-1165.	1.8	327
128	Detection of Vi-Negative Salmonella enterica Serovar Typhi in the Peripheral Blood of Patients with Typhoid Fever in the Faisalabad Region of Pakistan. Journal of Clinical Microbiology, 2005, 43, 4418-4425.	1.8	333
129	Typhoid and paratyphoid fever. Lancet, The, 2005, 366, 1603-1604.	6.3	11
130	The emergence of multidrug resistance to antimicrobial agents for the treatment of typhoid fever. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2004, 98, 423-430.	0.7	72
131	A Clinical, Microbiological, and Pathological Study of Intestinal Perforation Associated with Typhoid Fever. Clinical Infectious Diseases, 2004, 39, 61-67.	2.9	79
132	The emergence of antibiotic resistance in typhoid fever. Travel Medicine and Infectious Disease, 2004, 2, 67-74.	1.5	58
133	The Role of Prophage-like Elements in the Diversity of Salmonella enterica Serovars. Journal of Molecular Biology, 2004, 339, 279-300.	2.0	111
134	Double blind comparison of ibuprofen and paracetamol for adjunctive treatment of uncomplicated typhoid fever. Pediatric Infectious Disease Journal, 2004, 23, 226-230.	1.1	26
135	Molecular Analysis of <i>incHI1</i> Antimicrobial Resistance Plasmids from Salmonella Serovar Typhi Strains Associated with Typhoid Fever. Antimicrobial Agents and Chemotherapy, 2003, 47, 2732-2739.	1.4	88
136	Composition, Acquisition, and Distribution of the Vi Exopolysaccharide-Encoding Salmonella enterica Pathogenicity Island SPI-7. Journal of Bacteriology, 2003, 185, 5055-5065.	1.0	142
137	Emergence in Vietnam of <i>Streptococcus pneumoniae</i> Resistant to Multiple Antimicrobial Agents as a Result of Dissemination of the Multiresistant Spain 23F -1 Clone. Antimicrobial Agents and Chemotherapy, 2002, 46, 3512-3517.	1.4	28
138	Cytokine Release by Lipopolysaccharide- $\alpha$ -Stimulated Whole Blood from Patients with Typhoid Fever. Journal of Infectious Diseases, 2002, 186, 240-245.	1.9	22
139	Unlocking the genome of the human typhoid bacillus. Lancet Infectious Diseases, The, 2002, 2, 163-170.	4.6	39
140	Characterisation and distribution of a cryptic Salmonella typhi plasmid pHCM2. Plasmid, 2002, 47, 159-171.	0.4	36
141	Salmonella typhi, the causative agent of typhoid fever, is approximately 50,000 years old. Infection, Genetics and Evolution, 2002, 2, 39-45.	1.0	328
142	Genes of the Class II and Class III Major Histocompatibility Complex Are Associated with Typhoid Fever in Vietnam. Journal of Infectious Diseases, 2001, 183, 261-268.	1.9	95
143	The Escherichia coli gene pool. Current Opinion in Microbiology, 2001, 4, 90-94.	2.3	23
144	The molecular mechanisms of severe typhoid fever. Trends in Microbiology, 2001, 9, 316-320.	3.5	109

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145	Typhoid fever: pathogenesis and disease. <i>Current Opinion in Infectious Diseases</i> , 2001, 14, 573-578.	1.3	140
146	Risk factors for typhoid fever in the Mekong delta, southern Viet Nam: a case-control study. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2001, 95, 19-23.	0.7	62
147	Complete genome sequence of a multiple drug resistant <i>Salmonella enterica</i> serovar Typhi CT18. <i>Nature</i> , 2001, 413, 848-852.	13.7	1,192
148	Serology of Typhoid Fever in an Area of Endemicity and Its Relevance to Diagnosis. <i>Journal of Clinical Microbiology</i> , 2001, 39, 1002-1007.	1.8	119
149	Typhoid Fever and Genetic Polymorphisms at the Natural Resistance-associated Macrophage Protein 1. <i>Journal of Infectious Diseases</i> , 2001, 183, 1156-1160.	1.9	39
150	<i>Yersinia pestis</i> pFra Shows Biovar-Specific Differences and Recent Common Ancestry with a <i>Salmonella enterica</i> Serovar Typhi Plasmid. <i>Journal of Bacteriology</i> , 2001, 183, 2586-2594.	1.0	56
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