

James R Johnson

List of Publications by Year in descending order

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313
papers

24,179
citations

5248

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9839

141
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317
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docs citations

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times ranked

12994
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#	ARTICLE	IF	CITATIONS
1	Extended Virulence Genotypes of <i>Escherichia coli</i> Strains from Patients with Urosepsis in Relation to Phylogeny and Host Compromise. <i>Journal of Infectious Diseases</i> , 2000, 181, 261-272.	1.9	1,091
2	Guidelines for Antimicrobial Treatment of Uncomplicated Acute Bacterial Cystitis and Acute Pyelonephritis in Women. <i>Clinical Infectious Diseases</i> , 1999, 29, 745-758.	2.9	1,005
3	Organised Genome Dynamics in the <i>Escherichia coli</i> Species Results in Highly Diverse Adaptive Paths. <i>PLoS Genetics</i> , 2009, 5, e1000344.	1.5	1,005
4	Intercontinental emergence of <i>Escherichia coli</i> clone O25:H4-ST131 producing CTX-M-15. <i>Journal of Antimicrobial Chemotherapy</i> , 2007, 61, 273-281.	1.3	737
5	Medical and economic impact of extraintestinal infections due to <i>Escherichia coli</i> : focus on an increasingly important endemic problem. <i>Microbes and Infection</i> , 2003, 5, 449-456.	1.0	649
6	Widespread Distribution of Urinary Tract Infections Caused by a Multidrug-Resistant <i>Escherichia coli</i> Clonal Group. <i>New England Journal of Medicine</i> , 2001, 345, 1007-1013.	13.9	470
7	<i>Escherichia coli</i> Sequence Type ST131 as the Major Cause of Serious Multidrug-Resistant <i>E. coli</i> Infections in the United States. <i>Clinical Infectious Diseases</i> , 2010, 51, 286-294.	2.9	457
8	The Epidemic of Extended-Spectrum- β -Lactamase-Producing <i>Escherichia coli</i> ST131 Is Driven by a Single Highly Pathogenic Subclone, <i>H30-Rx</i> . <i>MBio</i> , 2013, 4, e00377-13.	1.8	380
9	Phylogenetic Distribution of Extraintestinal Virulence-Associated Traits in <i>Escherichia coli</i> . <i>Journal of Infectious Diseases</i> , 2001, 183, 78-88.	1.9	356
10	The Genome Sequence of Avian Pathogenic <i>Escherichia coli</i> Strain O1:K1:H7 Shares Strong Similarities with Human Extraintestinal Pathogenic <i>E. coli</i> Genomes. <i>Journal of Bacteriology</i> , 2007, 189, 3228-3236.	1.0	342
11	Extraintestinal pathogenic <i>Escherichia coli</i> : "The other bad <i>E. coli</i> ". <i>Translational Research</i> , 2002, 139, 155-162.	2.4	326
12	Isolation and Molecular Characterization of Nalidixic Acid-Resistant Extraintestinal Pathogenic <i>Escherichia coli</i> from Retail Chicken Products. <i>Antimicrobial Agents and Chemotherapy</i> , 2003, 47, 2161-2168.	1.4	301
13	Evolutionary History of the Global Emergence of the <i>Escherichia coli</i> Epidemic Clone ST131. <i>MBio</i> , 2016, 7, e02162.	1.8	289
14	Systematic Review: Antimicrobial Urinary Catheters To Prevent Catheter-Associated Urinary Tract Infection in Hospitalized Patients. <i>Annals of Internal Medicine</i> , 2006, 144, 116.	2.0	276
15	Comparison of Extraintestinal Pathogenic <i>Escherichia coli</i> Strains from Human and Avian Sources Reveals a Mixed Subset Representing Potential Zoonotic Pathogens. <i>Applied and Environmental Microbiology</i> , 2008, 74, 7043-7050.	1.4	256
16	Food-Borne Origins of <i>Escherichia coli</i> Causing Extraintestinal Infections. <i>Clinical Infectious Diseases</i> , 2012, 55, 712-719.	2.9	255
17	Epidemic Clonal Groups of <i>Escherichia coli</i> as a Cause of Antimicrobial-Resistant Urinary Tract Infections in Canada, 2002 to 2004. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2733-2739.	1.4	249
18	Abrupt Emergence of a Single Dominant Multidrug-Resistant Strain of <i>Escherichia coli</i> . <i>Journal of Infectious Diseases</i> , 2013, 207, 919-928.	1.9	247

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19	Extended-Spectrum β -Lactamase-Producing <i>Escherichia coli</i> From Retail Chicken Meat and Humans: Comparison of Strains, Plasmids, Resistance Genes, and Virulence Factors. <i>Clinical Infectious Diseases</i> , 2013, 56, 478-487.	2.9	233
20	Phylogenetic Origin and Virulence Genotype in Relation to Resistance to Fluoroquinolones and/or Extended-Spectrum Cephalosporins and Cephamycins among <i>Escherichia coli</i> Isolates from Animals and Humans. <i>Journal of Infectious Diseases</i> , 2003, 188, 759-768.	1.9	227
21	<i>Escherichia coli</i> Isolates That Carry <i>vat</i> , <i>fyuA</i> , <i>chuA</i> , and <i>yfcV</i> Efficiently Colonize the Urinary Tract. <i>Infection and Immunity</i> , 2012, 80, 4115-4122.	1.0	226
22	Loop-Mediated Isothermal Amplification Assay for Rapid Detection of Common Strains of <i>Escherichia coli</i> . <i>Journal of Clinical Microbiology</i> , 2008, 46, 2800-2804.	1.8	225
23	Antimicrobial-Resistant and Extraintestinal Pathogenic <i>Escherichia coli</i> in Retail Foods. <i>Journal of Infectious Diseases</i> , 2005, 191, 1040-1049.	1.9	223
24	Molecular epidemiology of extraintestinal pathogenic (uropathogenic) <i>Escherichia coli</i> . <i>International Journal of Medical Microbiology</i> , 2005, 295, 383-404.	1.5	218
25	A New Clone Sweeps Clean: the Enigmatic Emergence of <i>Escherichia coli</i> Sequence Type 131. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 4997-5004.	1.4	207
26	Antimicrobial Drug-Resistant <i>Escherichia coli</i> from Humans and Poultry Products, Minnesota and Wisconsin, 2002-2004. <i>Emerging Infectious Diseases</i> , 2007, 13, 838-846.	2.0	190
27	<i>Escherichia coli</i> ST131- <i>H</i> 22 as a Foodborne Uropathogen. <i>MBio</i> , 2018, 9, .	1.8	184
28	<i>In Silico</i> Genotyping of <i>Escherichia coli</i> Isolates for Extraintestinal Virulence Genes by Use of Whole-Genome Sequencing Data. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	1.8	179
29	High-Resolution Two-Locus Clonal Typing of Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>Applied and Environmental Microbiology</i> , 2012, 78, 1353-1360.	1.4	172
30	Relationship between <i>Escherichia coli</i> Strains Causing Acute Cystitis in Women and the Fecal <i>E. coli</i> Population of the Host. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2529-2534.	1.8	169
31	<i>Escherichia coli</i> Sequence Type 131 (ST131) Subclone H30 as an Emergent Multidrug-Resistant Pathogen Among US Veterans. <i>Clinical Infectious Diseases</i> , 2013, 57, 1256-1265.	2.9	167
32	Molecular Epidemiology and Phylogenetic Distribution of the <i>Escherichia coli</i> <i>pks</i> Genomic Island. <i>Journal of Clinical Microbiology</i> , 2008, 46, 3906-3911.	1.8	157
33	Microbial virulence determinants and the pathogenesis of urinary tract infection. <i>Infectious Disease Clinics of North America</i> , 2003, 17, 261-278.	1.9	156
34	Zoonotic Potential of <i>Escherichia coli</i> Isolates from Retail Chicken Meat Products and Eggs. <i>Applied and Environmental Microbiology</i> , 2015, 81, 1177-1187.	1.4	156
35	Extended Virulence Genotypes and Phylogenetic Background of <i>Escherichia coli</i> Isolates from Patients with Cystitis, Pyelonephritis, or Prostatitis. <i>Journal of Infectious Diseases</i> , 2005, 191, 46-50.	1.9	151
36	Molecular Epidemiological and Phylogenetic Associations of Two Novel Putative Virulence Genes, <i>iha</i> and <i>iroN</i> <i>E. coli</i> , among <i>Escherichia coli</i> Isolates from Patients with Urosepsis. <i>Infection and Immunity</i> , 2000, 68, 3040-3047.	1.0	150

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37	Phylogenetic Distribution of Virulence-Associated Genes among <i>Escherichia coli</i> Isolates Associated with Neonatal Bacterial Meningitis in The Netherlands. <i>Journal of Infectious Diseases</i> , 2002, 185, 774-784.	1.9	150
38	Experimental Mouse Lethality of <i>Escherichia coli</i> Isolates, in Relation to Accessory Traits, Phylogenetic Group, and Ecological Source. <i>Journal of Infectious Diseases</i> , 2006, 194, 1141-1150.	1.9	146
39	Acquisition of Avian Pathogenic <i>Escherichia coli</i> Plasmids by a Commensal <i>E. coli</i> Isolate Enhances Its Abilities To Kill Chicken Embryos, Grow in Human Urine, and Colonize the Murine Kidney. <i>Infection and Immunity</i> , 2006, 74, 6287-6292.	1.0	145
40	The genetic structure of <i>Escherichia coli</i> populations in primary and secondary habitats. <i>Microbiology (United Kingdom)</i> , 2002, 148, 1513-1522.	0.7	142
41	Similarity between Human and Chicken <i>Escherichia coli</i> Isolates in Relation to Ciprofloxacin Resistance Status. <i>Journal of Infectious Diseases</i> , 2006, 194, 71-78.	1.9	138
42	<i>Escherichia coli</i> Colonization Patterns among Human Household Members and Pets, with Attention to Acute Urinary Tract Infection. <i>Journal of Infectious Diseases</i> , 2008, 197, 218-224.	1.9	134
43	Rates of Mutation and Host Transmission for an <i>Escherichia coli</i> Clone over 3 Years. <i>PLoS ONE</i> , 2011, 6, e26907.	1.1	132
44	Quinolone, fluoroquinolone and trimethoprim/sulfamethoxazole resistance in relation to virulence determinants and phylogenetic background among uropathogenic <i>Escherichia coli</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2006, 57, 204-211.	1.3	131
45	Changes in Colonic Bile Acid Composition following Fecal Microbiota Transplantation Are Sufficient to Control <i>Clostridium difficile</i> Germination and Growth. <i>PLoS ONE</i> , 2016, 11, e0147210.	1.1	130
46	Multiple-Host Sharing, Long-Term Persistence, and Virulence of <i>Escherichia coli</i> Clones from Human and Animal Household Members. <i>Journal of Clinical Microbiology</i> , 2008, 46, 4078-4082.	1.8	127
47	Associations Between Multidrug Resistance, Plasmid Content, and Virulence Potential Among Extraintestinal Pathogenic and Commensal <i>Escherichia coli</i> from Humans and Poultry. <i>Foodborne Pathogens and Disease</i> , 2012, 9, 37-46.	0.8	126
48	Fimbrial Profiles Predict Virulence of Uropathogenic <i>Escherichia coli</i> Strains: Contribution of Ygi and Yad Fimbriae. <i>Infection and Immunity</i> , 2011, 79, 4753-4763.	1.0	121
49	<i>Escherichia coli</i> Sequence Type 131 Is a Dominant, Antimicrobial-Resistant Clonal Group Associated with Healthcare and Elderly Hosts. <i>Infection Control and Hospital Epidemiology</i> , 2013, 34, 361-369.	1.0	121
50	Identification of a New Iron-Regulated Virulence Gene, <i>ireA</i> , in an Extraintestinal Pathogenic Isolate of <i>Escherichia coli</i> . <i>Infection and Immunity</i> , 2001, 69, 6209-6216.	1.0	118
51	IroN Functions as a Siderophore Receptor and Is a Urovirulence Factor in an Extraintestinal Pathogenic Isolate of <i>Escherichia coli</i> . <i>Infection and Immunity</i> , 2002, 70, 7156-7160.	1.0	118
52	CTX-M-27- and CTX-M-14-producing, ciprofloxacin-resistant <i>Escherichia coli</i> of the H30 subclonal group within ST131 drive a Japanese regional ESBL epidemic. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 1639-1649.	1.3	118
53	Epidemiology of <i>Escherichia coli</i> Bacteremia: A Systematic Literature Review. <i>Clinical Infectious Diseases</i> , 2021, 72, 1211-1219.	2.9	116
54	Determination of <i>Escherichia coli</i> O types by allele-specific polymerase chain reaction: application to the O types involved in human septicemia. <i>Diagnostic Microbiology and Infectious Disease</i> , 2007, 57, 129-136.	0.8	115

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55	Evaluation of <i>Escherichia coli</i> isolates from healthy chickens to determine their potential risk to poultry and human health. <i>PLoS ONE</i> , 2017, 12, e0180599.	1.1	113
56	Commonality among Fluoroquinolone-Resistant Sequence Type ST131 Extraintestinal <i>Escherichia coli</i> Isolates from Humans and Companion Animals in Australia. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 3782-3787.	1.4	112
57	Molecular Epidemiology of <i>Escherichia coli</i> Sequence Type 131 and Its H30 and H30-Rx Subclones among Extended-Spectrum-β-Lactamase-Positive and -Negative <i>E. coli</i> Clinical Isolates from the Chicago Region, 2007 to 2010. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 6385-6388.	1.4	112
58	Extraintestinal Pathogenic <i>Escherichia coli</i> as a Cause of Invasive Nonurinary Infections. <i>Journal of Clinical Microbiology</i> , 2003, 41, 5798-5802.	1.8	111
59	Molecular Epidemiological Analysis of <i>Escherichia coli</i> Sequence Type ST131 (O25:H4) and CTX-M-15 among Extended-Spectrum-β-Lactamase-Producing <i>E. coli</i> from the United States, 2000 to 2009. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 2364-2370.	1.4	107
60	Rapid and Specific Detection, Molecular Epidemiology, and Experimental Virulence of the O16 Subgroup within <i>Escherichia coli</i> Sequence Type 131. <i>Journal of Clinical Microbiology</i> , 2014, 52, 1358-1365.	1.8	107
61	Bacterial Characteristics in Relation to Clinical Source of <i>Escherichia coli</i> Isolates from Women with Acute Cystitis or Pyelonephritis and Uninfected Women. <i>Journal of Clinical Microbiology</i> , 2005, 43, 6064-6072.	1.8	106
62	Sharing of Virulent <i>Escherichia coli</i> Clones among Household Members of a Woman with Acute Cystitis. <i>Clinical Infectious Diseases</i> , 2006, 43, e101-e108.	2.9	106
63	Canine Feces as a Reservoir of Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>Infection and Immunity</i> , 2001, 69, 1306-1314.	1.0	105
64	The IrgA Homologue Adhesin Iha Is an <i>Escherichia coli</i> Virulence Factor in Murine Urinary Tract Infection. <i>Infection and Immunity</i> , 2005, 73, 965-971.	1.0	105
65	A disseminated multidrug-resistant clonal group of uropathogenic <i>Escherichia coli</i> in pyelonephritis. <i>Lancet</i> , The, 2002, 359, 2249-2251.	6.3	104
66	Intermingled <i>Klebsiella pneumoniae</i> Populations Between Retail Meats and Human Urinary Tract Infections. <i>Clinical Infectious Diseases</i> , 2015, 61, 892-899.	2.9	104
67	Virulence Factor Profiles and Phylogenetic Background of <i>Escherichia coli</i> Isolates from Veterans with Bacteremia and Uninfected Control Subjects. <i>Journal of Infectious Diseases</i> , 2004, 190, 2121-2128.	1.9	102
68	Sharing of <i>Escherichia coli</i> Sequence Type ST131 and Other Multidrug-Resistant and Urovirulent <i>E. coli</i> Strains among Dogs and Cats within a Household. <i>Journal of Clinical Microbiology</i> , 2009, 47, 3721-3725.	1.8	102
69	Multidrug-resistant extraintestinal pathogenic <i>Escherichia coli</i> of sequence type ST131 in animals and foods. <i>Veterinary Microbiology</i> , 2011, 153, 99-108.	0.8	102
70	Virulence of <i>Escherichia coli</i> Clinical Isolates in a Murine Sepsis Model in Relation to Sequence Type ST131 Status, Fluoroquinolone Resistance, and Virulence Genotype. <i>Infection and Immunity</i> , 2012, 80, 1554-1562.	1.0	101
71	Host Characteristics and Bacterial Traits Predict Experimental Virulence for <i>Escherichia coli</i> Bloodstream Isolates From Patients With Urosepsis. <i>Open Forum Infectious Diseases</i> , 2015, 2, ofv083.	0.4	100
72	Ongoing Horizontal and Vertical Transmission of Virulence Genes and papA Alleles among <i>Escherichia coli</i> Blood Isolates from Patients with Diverse-Source Bacteremia. <i>Infection and Immunity</i> , 2001, 69, 5363-5374.	1.0	99

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73	Quinolone-Resistant Uropathogenic <i>Escherichia coli</i> Strains from Phylogenetic Group B2 Have Fewer Virulence Factors than Their Susceptible Counterparts. <i>Journal of Clinical Microbiology</i> , 2005, 43, 2962-2964.	1.8	99
74	Evidence of Commonality between Canine and Human Extraintestinal Pathogenic <i>Escherichia coli</i> Strains That Express papG Allele III. <i>Infection and Immunity</i> , 2000, 68, 3327-3336.	1.0	98
75	Separate F-Type Plasmids Have Shaped the Evolution of the <i>H</i> 30 Subclone of <i>Escherichia coli</i> Sequence Type 131. <i>MSphere</i> , 2016, 1, .	1.3	98
76	Activities of a Nitrofurazone-Containing Urinary Catheter and a Silver Hydrogel Catheter against Multidrug-Resistant Bacteria Characteristic of Catheter-Associated Urinary Tract Infection. <i>Antimicrobial Agents and Chemotherapy</i> , 1999, 43, 2990-2995.	1.4	96
77	Modulation of Host Innate Immune Response in the Bladder by Uropathogenic <i>Escherichia coli</i> . <i>Infection and Immunity</i> , 2007, 75, 5353-5360.	1.0	96
78	Virulence Characteristics and Phylogenetic Background of Multidrug-Resistant and Antimicrobial-Susceptible Clinical Isolates of <i>Escherichia coli</i> from across the United States, 2000-2001. <i>Journal of Infectious Diseases</i> , 2004, 190, 1739-1744.	1.9	95
79	Comparison of <i>Escherichia coli</i> ST131 Pulsotypes, by Epidemiologic Traits, 1967-2009. <i>Emerging Infectious Diseases</i> , 2012, 18, 598-607.	2.0	93
80	Selection Footprint in the FimH Adhesin Shows Pathoadaptive Niche Differentiation in <i>Escherichia coli</i> . <i>Molecular Biology and Evolution</i> , 2004, 21, 1373-1383.	3.5	91
81	Identification of Urovirulence Traits in <i>Escherichia coli</i> by Comparison of Urinary and Rectal <i>E. coli</i> Isolates from Dogs with Urinary Tract Infection. <i>Journal of Clinical Microbiology</i> , 2003, 41, 337-345.	1.8	89
82	Epidemiological Correlates of Virulence Genotype and Phylogenetic Background among <i>Escherichia coli</i> Blood Isolates from Adults with Diverse Source Bacteremia. <i>Journal of Infectious Diseases</i> , 2002, 185, 1439-1447.	1.9	88
83	Analysis of the F Antigen-Specific papA Alleles of Extraintestinal Pathogenic <i>Escherichia coli</i> Using a Novel Multiplex PCR-Based Assay. <i>Infection and Immunity</i> , 2000, 68, 1587-1599.	1.0	87
84	Identification of two previously unrecognized genes (<i>guaA</i> and <i>argC</i>) important for uropathogenesis. <i>Molecular Microbiology</i> , 1996, 22, 217-229.	1.2	86
85	Virulence Factors of <i>Escherichia coli</i> Isolates That Produce CTX-M-Type Extended-Spectrum β -Lactamases. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 4667-4670.	1.4	85
86	Distribution and Characteristics of <i>Escherichia coli</i> Clonal Group A1. <i>Emerging Infectious Diseases</i> , 2005, 11, 141-145.	2.0	84
87	Contamination of Retail Foods, Particularly Turkey, from Community Markets (Minnesota, 1999-2000) with Antimicrobial-Resistant and Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>Foodborne Pathogens and Disease</i> , 2005, 2, 38-49.	0.8	84
88	Uropathogenic <i>Escherichia coli</i> Induces Chronic Pelvic Pain. <i>Infection and Immunity</i> , 2011, 79, 628-635.	1.0	83
89	Hepatitis Due to Herpes Simplex Virus in Marrow-Transplant Recipients. <i>Clinical Infectious Diseases</i> , 1992, 14, 38-45.	2.9	81
90	Rapid and Extensive Expansion in the United States of a New Multidrug-resistant <i>Escherichia coli</i> Clonal Group, Sequence Type 1193. <i>Clinical Infectious Diseases</i> , 2019, 68, 334-337.	2.9	81

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91	Improved Repetitive-Element PCR Fingerprinting for Resolving Pathogenic and Nonpathogenic Phylogenetic Groups within <i>Escherichia coli</i> . <i>Vaccine Journal</i> , 2000, 7, 265-273.	2.6	80
92	The Clonal Distribution and Diversity of Extraintestinal <i>Escherichia coli</i> Isolates Vary According to Patient Characteristics. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 5912-5917.	1.4	80
93	The Pandemic H30 Subclone of Sequence Type 131 (ST131) as the Leading Cause of Multidrug-Resistant <i>Escherichia coli</i> Infections in the United States (2011-2012). <i>Open Forum Infectious Diseases</i> , 2017, 4, ofx089.	0.4	79
94	Phylogenetic relationships among clonal groups of extraintestinal pathogenic <i>Escherichia coli</i> as assessed by multi-locus sequence analysis. <i>Microbes and Infection</i> , 2006, 8, 1702-1713.	1.0	78
95	Uropathogenic <i>Escherichia coli</i> Agents of Diverse Non-“Urinary Tract Extraintestinal Infections. <i>Journal of Infectious Diseases</i> , 2002, 186, 859-864.	1.9	77
96	Enteraggregative <i>Escherichia coli</i> O78:H10, the Cause of an Outbreak of Urinary Tract Infection. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3703-3711.	1.8	77
97	Prevalence and Characteristics of the Epidemic Multiresistant <i>Escherichia coli</i> ST131 Clonal Group among Extended-Spectrum Beta-Lactamase-Producing <i>E. coli</i> Isolates in Copenhagen, Denmark. <i>Journal of Clinical Microbiology</i> , 2013, 51, 1779-1785.	1.8	77
98	Clonal analysis reveals high rate of structural mutations in fimbrial adhesins of extraintestinal pathogenic <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2006, 59, 975-988.	1.2	76
99	Four Main Virotypes among Extended-Spectrum-β-Lactamase-Producing Isolates of <i>Escherichia coli</i> O25b:H4-B2-ST131: Bacterial, Epidemiological, and Clinical Characteristics. <i>Journal of Clinical Microbiology</i> , 2013, 51, 3358-3367.	1.8	76
100	Virulence Genotype and Phylogenetic Origin in Relation to Antibiotic Resistance Profile among <i>Escherichia coli</i> Urine Sample Isolates from Israeli Women with Acute Uncomplicated Cystitis. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 26-31.	1.4	72
101	Reservoirs of Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>Microbiology Spectrum</i> , 2015, 3, .	1.2	71
102	<i>Escherichia coli</i> Pyomyositis: An Emerging Infectious Disease among Patients with Hematologic Malignancies. <i>Clinical Infectious Diseases</i> , 2010, 50, 374-380.	2.9	70
103	Carbapenemase-producing bacteria in companion animals: a public health concern on the horizon. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 1155-1157.	1.3	68
104	Intensity and Mechanisms of Fluoroquinolone Resistance within the <i>H</i> 30 and <i>H</i> 30Rx Subclones of <i>Escherichia coli</i> Sequence Type 131 Compared with Other Fluoroquinolone-Resistant <i>E. coli</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 4471-4480.	1.4	68
105	Virulence Genotypes and Phylogenetic Background of <i>Escherichia coli</i> Serogroup O6 Isolates from Humans, Dogs, and Cats. <i>Journal of Clinical Microbiology</i> , 2008, 46, 417-422.	1.8	67
106	Structure and urovirulence characteristics of the fecal <i>Escherichia coli</i> population among healthy women. <i>Microbes and Infection</i> , 2009, 11, 274-280.	1.0	67
107	Prominence of an O75 Clonal Group (Clonal Complex 14) among Non-ST131 Fluoroquinolone-Resistant <i>Escherichia coli</i> Causing Extraintestinal Infections in Humans and Dogs in Australia. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 3898-3904.	1.4	66
108	Ciprofloxacin-Resistant Gram-Negative Bacilli in the Fecal Microflora of Children. <i>Antimicrobial Agents and Chemotherapy</i> , 2006, 50, 3325-3329.	1.4	64

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109	Phylogenomic Analysis of Extraintestinal Pathogenic <i>Escherichia coli</i> Sequence Type 1193, an Emerging Multidrug-Resistant Clonal Group. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	1.4	64
110	Transmission of an Extended-Spectrum-Beta-Lactamase-Producing <i>Escherichia coli</i> (Sequence) Tj ETQq0 0 0 rgBT /Overlock 10 Tf Pyelonephritis. <i>Journal of Clinical Microbiology</i> , 2009, 47, 3780-3782.	1.8	63
111	Genetic Diversity and Virulence Profiles of <i>Escherichia coli</i> Isolates Causing Spontaneous Bacterial Peritonitis and Bacteremia in Patients with Cirrhosis. <i>Journal of Clinical Microbiology</i> , 2010, 48, 2709-2714.	1.8	63
112	<i>Escherichia coli</i> Serotype O15:K52:H1 as a Uropathogenic Clone. <i>Journal of Clinical Microbiology</i> , 2000, 38, 201-209.	1.8	63
113	Phylogenetic Background and Virulence Profiles of Fluoroquinolone-Resistant Clinical <i>Escherichia coli</i> Isolates from The Netherlands. <i>Journal of Infectious Diseases</i> , 2002, 186, 1852-1856.	1.9	62
114	Predictive Diagnostics for <i>Escherichia coli</i> Infections Based on the Clonal Association of Antimicrobial Resistance and Clinical Outcome. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2991-2999.	1.8	62
115	Household Clustering of <i>Escherichia coli</i> Sequence Type 131 Clinical and Fecal Isolates According to Whole Genome Sequence Analysis. <i>Open Forum Infectious Diseases</i> , 2016, 3, ofw129.	0.4	62
116	Clonal Origin, Virulence Factors, and Virulence. <i>Infection and Immunity</i> , 2000, 68, 424-425.	1.0	61
117	<i>Escherichia coli</i> Sequence Type 131 as a Prominent Cause of Antibiotic Resistance among Urinary <i>Escherichia coli</i> Isolates from Reproductive-Age Women. <i>Journal of Clinical Microbiology</i> , 2013, 51, 3270-3276.	1.8	61
118	Rapid and Specific Detection of <i>Escherichia coli</i> Clonal Group A by Gene-Specific PCR. <i>Journal of Clinical Microbiology</i> , 2004, 42, 2618-2622.	1.8	60
119	<i>Escherichia coli</i> Sequence Type ST131 as an Emerging Fluoroquinolone-Resistant Uropathogen among Renal Transplant Recipients. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 546-550.	1.4	60
120	Gut Colonization of Healthy Children and Their Mothers With Pathogenic Ciprofloxacin-Resistant <i>Escherichia coli</i> . <i>Journal of Infectious Diseases</i> , 2015, 212, 1862-1868.	1.9	60
121	Urinary Tract Infections. <i>Primary Care - Clinics in Office Practice</i> , 2008, 35, 345-367.	0.7	59
122	Phylogenetic diversity, antimicrobial susceptibility and virulence characteristics of phylogroup F <i>Escherichia coli</i> in Australia. <i>Microbiology (United Kingdom)</i> , 2016, 162, 1904-1912.	0.7	59
123	Molecular Epidemiology of Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>EcoSal Plus</i> , 2018, 8, .	2.1	57
124	<i>lha</i> from an <i>Escherichia coli</i> Urinary Tract Infection Outbreak Clonal Group A Strain Is Expressed In Vivo in the Mouse Urinary Tract and Functions as a Catecholate Siderophore Receptor. <i>Infection and Immunity</i> , 2006, 74, 3427-3436.	1.0	56
125	Clonal group distribution of fluoroquinolone-resistant <i>Escherichia coli</i> among humans and companion animals in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2010, 65, 1936-1938.	1.3	56
126	Rapid Emergence, Subsidence, and Molecular Detection of <i>Escherichia coli</i> Sequence Type 1193- <i>fimH64</i> , a New Disseminated Multidrug-Resistant Commensal and Extraintestinal Pathogen. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	1.8	56

#	ARTICLE	IF	CITATIONS
127	Human-associated fluoroquinolone-resistant <i>Escherichia coli</i> clonal lineages, including ST354, isolated from canine feces and extraintestinal infections in Australia. <i>Microbes and Infection</i> , 2015, 17, 266-274.	1.0	55
128	<i>papG</i> Alleles among <i>Escherichia coli</i> Strains Causing Urosepsis: Associations with Other Bacterial Characteristics and Host Compromise. <i>Infection and Immunity</i> , 1998, 66, 4568-4571.	1.0	55
129	Predictors and Molecular Epidemiology of Community-Onset Extended-Spectrum β -Lactamase-Producing <i>Escherichia coli</i> Infection in a Midwestern Community. <i>Infection Control and Hospital Epidemiology</i> , 2013, 34, 947-953.	1.0	54
130	Colonisation with ESBL-producing and carbapenemase-producing Enterobacteriaceae, vancomycin-resistant enterococci, and meticillin-resistant <i>Staphylococcus aureus</i> in a long-term care facility over one year. <i>BMC Infectious Diseases</i> , 2015, 15, 168.	1.3	54
131	Global Molecular Epidemiology of the O15:K52:H1 Extraintestinal Pathogenic <i>Escherichia coli</i> Clonal Group: Evidence of Distribution beyond Europe. <i>Journal of Clinical Microbiology</i> , 2002, 40, 1913-1923.	1.8	53
132	Phylogenetic and Pathotypic Comparison of Concurrent Urine and Rectal <i>Escherichia coli</i> Isolates from Men with Febrile Urinary Tract Infection. <i>Journal of Clinical Microbiology</i> , 2005, 43, 3895-3900.	1.8	51
133	Three-Decade Epidemiological Analysis of <i>Escherichia coli</i> O15:K52:H1. <i>Journal of Clinical Microbiology</i> , 2009, 47, 1857-1862.	1.8	51
134	Presence of Putative Repeat-in-Toxin Gene <i>tosA</i> in <i>Escherichia coli</i> Predicts Successful Colonization of the Urinary Tract. <i>MBio</i> , 2011, 2, e00066-11.	1.8	51
135	Occurrence of Antibiotic-Resistant Uropathogenic <i>Escherichia coli</i> Clonal Group A in Wastewater Effluents. <i>Applied and Environmental Microbiology</i> , 2007, 73, 4180-4184.	1.4	50
136	Role of Homologous Recombination in Adaptive Diversification of Extraintestinal <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2013, 195, 231-242.	1.0	50
137	Risk Factors for Trimethoprim-Sulfamethoxazole Resistance in Patients with Acute Uncomplicated Cystitis. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 846-851.	1.4	49
138	Clinical and Microbiological Determinants of Infection After Transrectal Prostate Biopsy. <i>Clinical Infectious Diseases</i> , 2015, 60, 979-987.	2.9	49
139	Genetic Structure and Antimicrobial Resistance of <i>Escherichia coli</i> and Cryptic Clades in Birds with Diverse Human Associations. <i>Applied and Environmental Microbiology</i> , 2015, 81, 5123-5133.	1.4	49
140	Molecular Analysis of <i>Escherichia coli</i> from Retail Meats (2002-2004) from the United States National Antimicrobial Resistance Monitoring System. <i>Clinical Infectious Diseases</i> , 2009, 49, 195-201.	2.9	48
141	Prolonged colonisation with <i>Escherichia coli</i> O25:ST131 versus other extended-spectrum beta-lactamase-producing <i>E. coli</i> in a long-term care facility with high endemic level of rectal colonisation, the Netherlands, 2013 to 2014. <i>Eurosurveillance</i> , 2016, 21, .	3.9	48
142	Effect of Inactivation of the Global Oxidative Stress Regulator <i>oxyR</i> on the Colonization Ability of <i>Escherichia coli</i> O1:K1:H7 in a Mouse Model of Ascending Urinary Tract Infection. <i>Infection and Immunity</i> , 2006, 74, 461-468.	1.0	47
143	Antimicrobial urinary catheters: a systematic review. <i>Expert Review of Medical Devices</i> , 2008, 5, 495-506.	1.4	47
144	Extensive Household Outbreak of Urinary Tract Infection and Intestinal Colonization due to Extended-Spectrum β -Lactamase-Producing <i>Escherichia coli</i> Sequence Type 131. <i>Clinical Infectious Diseases</i> , 2015, 61, e5-e12.	2.9	46

#	ARTICLE	IF	CITATIONS
145	Temporal Trends in Antimicrobial Resistance and Virulence-Associated Traits within the Escherichia coli Sequence Type 131 Clonal Group and Its <i>H</i> 30 and <i>H</i> 30-Rx Subclones, 1968 to 2012. Antimicrobial Agents and Chemotherapy, 2014, 58, 6886-6895.	1.4	45
146	Extraintestinal Pathogenic and Antimicrobial-Resistant Escherichia coli, Including Sequence Type 131 (ST131), from Retail Chicken Breasts in the United States in 2013. Applied and Environmental Microbiology, 2017, 83, .	1.4	45
147	Epidemic Emergence in the United States of Escherichia coli Sequence Type 131- <i>H</i> 30 (ST131)- Tj ETQq1 1 0.784314 <i>rg</i> BT /Over	1.4	45
148	Community Transmission in the United States of a CTX-M-15-Producing Sequence Type ST131 Escherichia coli Strain Resulting in Death. Journal of Clinical Microbiology, 2011, 49, 3406-3408.	1.8	44
149	<i>In Vitro</i> Comparison of Nitrofurazone- and Silver Alloy-Coated Foley Catheters for Contact-Dependent and Diffusible Inhibition of Urinary Tract Infection-Associated Microorganisms. Antimicrobial Agents and Chemotherapy, 2012, 56, 4969-4972.	1.4	44
150	Development of polymerase chain reaction-based assays for bacterial gene detection. Journal of Microbiological Methods, 2000, 41, 201-209.	0.7	43
151	Genomic Analysis of Multidrug-Resistant Escherichia coli from North Carolina Community Hospitals: Ongoing Circulation of CTX-M-Producing ST131- <i>H</i> 30Rx and ST131- <i>H</i> 30R1 Strains. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	43
152	WITHIN-HOUSEHOLD SHARING OF A FLUOROQUINOLONE-RESISTANT ESCHERICHIA COLI SEQUENCE TYPE ST131 STRAIN CAUSING PEDIATRIC OSTEOARTICULAR INFECTION. Pediatric Infectious Disease Journal, 2010, 29, 473-475.	1.1	42
153	Complete Genome Sequence of the Epidemic and Highly Virulent CTX-M-15-Producing <i>H</i> 30-Rx Subclone of Escherichia coli ST131. Genome Announcements, 2013, 1, .	0.8	42
154	Diagnostic Errors that Lead to Inappropriate Antimicrobial Use. Infection Control and Hospital Epidemiology, 2015, 36, 949-956.	1.0	42
155	Analysis of mutational patterns in quinolone resistance-determining regions of C _{yr} A and ParC of clinical isolates. International Journal of Antimicrobial Agents, 2019, 53, 318-324.	1.1	42
156	Horizontally acquired papGII-containing pathogenicity islands underlie the emergence of invasive uropathogenic Escherichia coli lineages. Nature Communications, 2020, 11, 5968.	5.8	42
157	Virulence Factors Predict <i>Escherichia coli</i> Colonization Patterns among Human and Animal Household Members. Annals of Internal Medicine, 2004, 140, 848.	2.0	41
158	Virulence genotypes and phylogenetic background of fluoroquinolone-resistant and susceptible Escherichia coli urine isolates from dogs with urinary tract infection. Veterinary Microbiology, 2009, 136, 108-114.	0.8	41
159	The Pandemic <i>H</i> 30 Subclone of <i>Escherichia coli</i> Sequence Type 131 Is Associated With Persistent Infections and Adverse Outcomes Independent From Its Multidrug Resistance and Associations With Compromised Hosts. Clinical Infectious Diseases, 2016, 62, 1529-1536.	2.9	41
160	Spontaneous Conversion to Quinolone and Fluoroquinolone Resistance among Wild-Type Escherichia coli Isolates in Relation to Phylogenetic Background and Virulence Genotype. Antimicrobial Agents and Chemotherapy, 2005, 49, 4739-4744.	1.4	40
161	Multiplex PCR-Based Reverse Line Blot Assay for Simultaneous Detection of 22 Virulence Genes in Uropathogenic Escherichia coli. Applied and Environmental Microbiology, 2012, 78, 1198-1202.	1.4	40
162	Internet survey of Foley catheter practices and knowledge among Minnesota nurses. American Journal of Infection Control, 2010, 38, 31-37.	1.1	39

#	ARTICLE	IF	CITATIONS
163	Accessory Traits and Phylogenetic Background Predict <i>Escherichia coli</i> Extraintestinal Virulence Better Than Does Ecological Source. <i>Journal of Infectious Diseases</i> , 2019, 219, 121-132.	1.9	38
164	Companion Animals Are Spillover Hosts of the Multidrug-Resistant Human Extraintestinal <i>Escherichia coli</i> Pandemic Clones ST131 and ST1193. <i>Frontiers in Microbiology</i> , 2020, 11, 1968.	1.5	38
165	Host-Pathogen Relationships among <i>Escherichia coli</i> Isolates Recovered from Men with Febrile Urinary Tract Infection. <i>Clinical Infectious Diseases</i> , 2005, 40, 813-822.	2.9	37
166	Global Distribution and Epidemiologic Associations of <i>Escherichia coli</i> Clonal Group A, 1998–2007. <i>Emerging Infectious Diseases</i> , 2011, 17, 2001-9.	2.0	36
167	A Widely Used In Vitro Biofilm Assay Has Questionable Clinical Significance for Enterococcal Endocarditis. <i>PLoS ONE</i> , 2014, 9, e107282.	1.1	36
168	Fimoperon variation in the emergence of Enterohemorrhagic <i>Escherichia coli</i> : an evolutionary and functional analysis. <i>FEMS Microbiology Letters</i> , 2007, 273, 58-63.	0.7	35
169	Genotypic and Phenotypic Characterization of <i>Escherichia coli</i> Isolates From Children With Urinary Tract Infection and From Healthy Carriers. <i>Pediatric Infectious Disease Journal</i> , 2013, 32, 543-548.	1.1	35
170	Rapid and Specific Detection of the <i>Escherichia coli</i> Sequence Type 648 Complex within Phylogroup F. <i>Journal of Clinical Microbiology</i> , 2017, 55, 1116-1121.	1.8	35
171	Alpha-Toxin Contributes to Biofilm Formation among <i>Staphylococcus aureus</i> Wound Isolates. <i>Toxins</i> , 2018, 10, 157.	1.5	35
172	Identification of CTX-M β -lactamases in <i>Escherichia coli</i> from hospitalized patients and residents of long-term care facilities. <i>Diagnostic Microbiology and Infectious Disease</i> , 2010, 66, 402-406.	0.8	34
173	Long-Term Care Facilities Are Reservoirs for Antimicrobial-Resistant Sequence Type 131 <i>Escherichia coli</i> . <i>Open Forum Infectious Diseases</i> , 2015, 2, ofv011.	0.4	34
174	The Uropathogenic <i>Escherichia coli</i> Subclone Sequence Type 131-H30 Is Responsible for Most Antibiotic Prescription Errors at an Urgent Care Clinic. <i>Clinical Infectious Diseases</i> , 2019, 68, 781-787.	2.9	34
175	Enteraggregative <i>Escherichia coli</i> Related to Uropathogenic Clonal Group A. <i>Emerging Infectious Diseases</i> , 2007, 13, 757-760.	2.0	33
176	A Module Located at a Chromosomal Integration Hot Spot Is Responsible for the Multidrug Resistance of a Reference Strain from <i>Escherichia coli</i> Clonal Group A. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2283-2288.	1.4	33
177	Environmental Contamination in Households of Patients with Recurrent <i>Clostridium difficile</i> Infection. <i>Applied and Environmental Microbiology</i> , 2016, 82, 2686-2692.	1.4	33
178	Extensive Genetic Commonality among Wildlife, Wastewater, Community, and Nosocomial Isolates of <i>Escherichia coli</i> Sequence Type 131 (H30R1 and H30Rx Subclones) That Carry bla _{CTX-M-27} or bla _{CTX-M-15} . <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	1.4	33
179	Detection of the <i>Escherichia coli</i> Group 2 Polysaccharide Capsule Synthesis Gene kpsM by a Rapid and Specific PCR-Based Assay. <i>Journal of Clinical Microbiology</i> , 2004, 42, 1773-1776.	1.8	32
180	Molecular Analysis of Antimicrobial-Susceptible and -Resistant <i>Escherichia coli</i> from Retail Meats and Human Stool and Clinical Specimens in a Rural Community Setting. <i>Foodborne Pathogens and Disease</i> , 2009, 6, 285-295.	0.8	32

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181	Prevalence of Avian-Pathogenic <i>Escherichia coli</i> Strain O1 Genomic Islands among Extraintestinal and Commensal <i>E. coli</i> Isolates. <i>Journal of Bacteriology</i> , 2012, 194, 2846-2853.	1.0	32
182	Fine-Scale Structure Analysis Shows Epidemic Patterns of Clonal Complex 95, a Cosmopolitan <i>Escherichia coli</i> Lineage Responsible for Extraintestinal Infection. <i>MSphere</i> , 2017, 2, .	1.3	32
183	Survey of US wastewater for carbapenem-resistant <i>Enterobacteriaceae</i> . <i>Journal of Water and Health</i> , 2019, 17, 219-226.	1.1	32
184	Haemolytic <i>Escherichia coli</i> isolated from dogs with diarrhea have characteristics of both uropathogenic and necrotoxicogenic strains. <i>Veterinary Microbiology</i> , 2002, 85, 361-377.	0.8	31
185	Geographical distribution of antimicrobial resistance among <i>Escherichia coli</i> causing acute uncomplicated pyelonephritis in the United States. <i>FEMS Immunology and Medical Microbiology</i> , 2004, 42, 213-218.	2.7	31
186	Extraintestinal isolates of <i>Escherichia coli</i> : identification and prospects for vaccine development. <i>Expert Review of Vaccines</i> , 2006, 5, 45-54.	2.0	31
187	In vivo correlates of molecularly inferred virulence among extraintestinal pathogenic <i>Escherichia coli</i> (ExPEC) in the wax moth <i>Galleria mellonella</i> model system. <i>Virulence</i> , 2014, 5, 388-393.	1.8	31
188	Caspase-Activated Cell-Penetrating Peptides Reveal Temporal Coupling Between Endosomal Release and Apoptosis in an RGC-5 Cell Model. <i>Bioconjugate Chemistry</i> , 2012, 23, 1783-1793.	1.8	30
189	A killed, genetically engineered derivative of a wild-type extraintestinal pathogenic <i>E. coli</i> strain is a vaccine candidate. <i>Vaccine</i> , 2007, 25, 3859-3870.	1.7	29
190	Prevalence of ST131 Among Fluoroquinolone-resistant <i>Escherichia coli</i> Obtained From Rectal Swabs Before Transrectal Prostate Biopsy. <i>Urology</i> , 2013, 81, 548-556.	0.5	29
191	Prevalence and characteristics of <i>Escherichia coli</i> sequence type 131 and its H30 and H30Rx subclones: a multicenter study from Korea. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 84, 97-101.	0.8	29
192	Predictors of Antimicrobial-Resistant <i>Escherichia coli</i> in the Feces of Vegetarians and Newly Hospitalized Adults in Minnesota and Wisconsin. <i>Journal of Infectious Diseases</i> , 2008, 197, 430-434.	1.9	28
193	Enterococcal Aggregation Substance and Binding Substance Are Not Major Contributors to Urinary Tract Colonization by <i>Enterococcus faecalis</i> in a Mouse Model of Ascending Unobstructed Urinary Tract Infection. <i>Infection and Immunity</i> , 2004, 72, 2445-2448.	1.0	24
194	Molecular Epidemiology of Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>EcoSal Plus</i> , 2004, 1, .	2.1	24
195	Fluoroquinolone-resistant <i>Escherichia coli</i> , Indonesia. <i>Emerging Infectious Diseases</i> , 2005, 11, 1363-1369.	2.0	24
196	Single-Cell Resolution Imaging of Retinal Ganglion Cell Apoptosis In Vivo Using a Cell-Penetrating Caspase-Activatable Peptide Probe. <i>PLoS ONE</i> , 2014, 9, e88855.	1.1	24
197	A Novel 7-Single Nucleotide Polymorphism-Based Clonotyping Test Allows Rapid Prediction of Antimicrobial Susceptibility of Extraintestinal <i>Escherichia coli</i> Directly From Urine Specimens. <i>Open Forum Infectious Diseases</i> , 2016, 3, ofw002.	0.4	24
198	Despite Predominance of Uropathogenic/Extraintestinal Pathotypes Among Travel-acquired Extended-spectrum β -Lactamase-producing <i>Escherichia coli</i> , the Most Commonly Associated Clinical Manifestation Is Travelers' Diarrhea. <i>Clinical Infectious Diseases</i> , 2020, 70, 210-218.	2.9	24

#	ARTICLE	IF	CITATIONS
199	Activity of Cefiderocol, Ceftazidime-Avibactam, and Eravacycline against Carbapenem-Resistant <i>Escherichia coli</i> Isolates from the United States and International Sites in Relation to Clonal Background, Resistance Genes, Coresistance, and Region. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	1.4	24
200	Clinical and bacteriologic correlates of the papG alleles among <i>Escherichia coli</i> strains from children with acute cystitis. <i>Pediatric Infectious Disease Journal</i> , 1999, 18, 446-451.	1.1	23
201	Pigeon and Dove Eggwhite Protect Mice Against Renal Infection Due to P Fimbriated <i>Escherichia coli</i> . <i>American Journal of the Medical Sciences</i> , 1994, 307, 335-339.	0.4	22
202	Rapid and Specific Detection of the O15:K52:H1 Clonal Group of <i>Escherichia coli</i> by Gene-Specific PCR. <i>Journal of Clinical Microbiology</i> , 2004, 42, 3841-3843.	1.8	22
203	Clinical and molecular correlates of virulence in <i>Escherichia coli</i> causing bloodstream infection following transrectal ultrasound-guided (TRUS) prostate biopsy. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 2898-2906.	1.3	22
204	Safety and efficacy of a novel silver-impregnated urinary catheter system for preventing catheter-associated bacteriuria: A pilot randomized clinical trial. <i>American Journal of Infection Control</i> , 2015, 43, 260-265.	1.1	22
205	Emergence of Enteraggregative <i>Escherichia coli</i> within the ST131 Lineage as a Cause of Extraintestinal Infections. <i>MBio</i> , 2020, 11, .	1.8	22
206	Diversity of Hemagglutination Phenotypes among P-Fimbriated Wild-Type Strains of <i>Escherichia coli</i> in Relation to papG Allele Repertoire. <i>Vaccine Journal</i> , 1998, 5, 160-170.	2.6	21
207	Novel Molecular Variants of Allele I of the <i>Escherichia coli</i> P Fimbrial Adhesin GenepapG. <i>Infection and Immunity</i> , 2001, 69, 2318-2327.	1.0	20
208	Colonization with Extraintestinal Pathogenic <i>Escherichia coli</i> among Nursing Home Residents and Its Relationship to Fluoroquinolone Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2004, 48, 3618-3620.	1.4	20
209	Antimicrobial resistance of <i>Escherichia coli</i> strains isolated from urine of women with cystitis or pyelonephritis and feces of dogs and healthy humans. <i>Journal of the American Veterinary Medical Association</i> , 2004, 225, 368-373.	0.2	20
210	Drug-resistant <i>Escherichia coli</i> , Rural Idaho. <i>Emerging Infectious Diseases</i> , 2005, 11, 1614-1617.	2.0	20
211	Foodborne Illness Acquired in the United States. <i>Emerging Infectious Diseases</i> , 2011, 17, 1338-1339.	2.0	20
212	Post-Prostate Biopsy Infection with <i>Escherichia coli</i> ST131 Leading to Epididymo-Orchitis and Meningitis Caused by Gram-Negative Bacilli. <i>Journal of Clinical Microbiology</i> , 2012, 50, 4157-4159.	1.8	20
213	Clonal distribution and associated characteristics of <i>Escherichia coli</i> clinical and surveillance isolates from a military medical center. <i>Diagnostic Microbiology and Infectious Disease</i> , 2017, 87, 382-385.	0.8	20
214	Genomic analysis of phylogenetic group B2 extraintestinal pathogenic <i>E. coli</i> causing infections in dogs in Australia. <i>Veterinary Microbiology</i> , 2020, 248, 108783.	0.8	20
215	High-Frequency Secondary Mutations after Suicide-Driven Allelic Exchange Mutagenesis in Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2003, 185, 5301-5305.	1.0	19
216	Susceptibility to Alternative Oral Antimicrobial Agents in Relation to Sequence Type ST131 Status and Coresistance Phenotype among Recent <i>Escherichia coli</i> Isolates from U.S. Veterans. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 4856-4860.	1.4	19

#	ARTICLE	IF	CITATIONS
217	Bacterial clonal diagnostics as a tool for evidence-based empiric antibiotic selection. <i>PLoS ONE</i> , 2017, 12, e0174132.	1.1	19
218	Identification of CTX-M β -lactamases among <i>Escherichia coli</i> from the community in New York City. <i>Diagnostic Microbiology and Infectious Disease</i> , 2012, 72, 248-252.	0.8	17
219	Fluoroquinolone-resistant extraintestinal pathogenic <i>Escherichia coli</i> , including O25b-ST131, isolated from faeces of hospitalized dogs in an Australian veterinary referral centre. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 1025-1031.	1.3	17
220	Colonisation dynamics and virulence of two clonal groups of multidrug-resistant <i>Escherichia coli</i> isolated from dogs. <i>Microbes and Infection</i> , 2009, 11, 100-107.	1.0	16
221	Transrectal Prostate Biopsy-Associated Prophylaxis and Infectious Complications: Report of a Query to the Emerging Infections Network of the Infectious Diseases Society of America. <i>Open Forum Infectious Diseases</i> , 2015, 2, ofv002.	0.4	16
222	Colonization with <i>Escherichia coli</i> Strains among Female Sex Partners of Men with Febrile Urinary Tract Infection. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1947-1950.	1.8	16
223	PCR for Specific Detection of H7 Flagellar Variant of <i>fliC</i> among Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>Journal of Clinical Microbiology</i> , 2001, 39, 3712-3717.	1.8	15
224	OxyR contributes to the virulence of a Clonal Group A <i>Escherichia coli</i> strain (O17:K+:H18) in animal models of urinary tract infection, subcutaneous infection, and systemic sepsis. <i>Microbial Pathogenesis</i> , 2013, 64, 1-5.	1.3	15
225	Molecularly defined extraintestinal pathogenic <i>Escherichia coli</i> status predicts virulence in a murine sepsis model better than does virotype, individual virulence genes, or clonal subset among <i>E. coli</i> ST131 isolates. <i>Virulence</i> , 2020, 11, 327-336.	1.8	15
226	Antimicrobial resistance in <i>Escherichia coli</i> causing urinary tract infections in Costa Rica: a clinical dilemma. <i>International Journal of Antimicrobial Agents</i> , 2003, 21, 79-81.	1.1	14
227	In Vitro Activity of Available Antimicrobial Coated Foley Catheters Against <i>Escherichia coli</i> , Including Strains Resistant to Extended Spectrum Cephalosporins. <i>Journal of Urology</i> , 2010, 184, 2572-2577.	0.2	14
228	Urine Cultures among Hospitalized Veterans: Casting Too Broad a Net?. <i>Infection Control and Hospital Epidemiology</i> , 2014, 35, 574-576.	1.0	14
229	Global molecular epidemiology of carbapenem-resistant <i>Escherichia coli</i> (2002–2017). <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, , 1.	1.3	14
230	Bacteriophage Cocktail and Microcin-Producing Probiotic <i>Escherichia coli</i> Protect Mice Against Gut Colonization With Multidrug-Resistant <i>Escherichia coli</i> Sequence Type 131. <i>Frontiers in Microbiology</i> , 2022, 13, 887799.	1.5	14
231	Concurrent Fecal Colonization with Extraintestinal Pathogenic <i>Escherichia coli</i> in a Homosexual Man with Recurrent Urinary Tract Infection and in His Male Sex Partner. <i>Clinical Infectious Diseases</i> , 2002, 35, e65-e68.	2.9	13
232	Phylogenetic Backgrounds and Virulence-Associated Traits of <i>Escherichia coli</i> Isolates from Surface Waters and Diverse Animals in Minnesota and Wisconsin. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	1.4	13
233	Contribution of yersiniabactin to the virulence of an <i>Escherichia coli</i> sequence type 69 (clonal) Tj ETQq1 1.0.784314 rgBT /Overlock Pathogenesis, 2018, 120, 128-131.	1.3	13
234	Prevalence and Molecular Characteristics of <i>Clostridium difficile</i> in Retail Meats, Food-Producing and Companion Animals, and Humans in Minnesota. <i>Journal of Food Protection</i> , 2018, 81, 1635-1642.	0.8	13

#	ARTICLE	IF	CITATIONS
235	Activity of Imipenem-Relebactam against Carbapenem-Resistant Escherichia coli Isolates from the United States in Relation to Clonal Background, Resistance Genes, Coresistance, and Region. Antimicrobial Agents and Chemotherapy, 2020, 64, .	1.4	13
236	Intestinal population dynamics of UTI-causing Escherichia coli within heterosexual couples. Current Issues in Intestinal Microbiology, 2004, 5, 49-57.	2.5	13
237	Should All Catheterized Patients with Candiduria Be Treated?. Clinical Infectious Diseases, 1993, 17, 814-814.	2.9	12
238	Greater Ciprofloxacin Tolerance as a Possible Selectable Phenotype Underlying the Pandemic Spread of the H30 Subclone of Escherichia coli Sequence Type 131. Antimicrobial Agents and Chemotherapy, 2015, 59, 7132-7135.	1.4	12
239	Inactivation of extraintestinal pathogenic E. coli clinical and food isolates suspended in ground chicken meat by gamma radiation. Food Microbiology, 2019, 84, 103264.	2.1	12
240	Host and microbial factors in kidney transplant recipients with Escherichia coli acute pyelonephritis or asymptomatic bacteriuria: a prospective study using whole-genome sequencing. Nephrology Dialysis Transplantation, 2019, 34, 878-885.	0.4	12
241	Genomic analysis of fluoroquinolone-susceptible phylogenetic group B2 extraintestinal pathogenic Escherichia coli causing infections in cats. Veterinary Microbiology, 2020, 245, 108685.	0.8	12
242	Analysis of Urinary Escherichia coli Isolates for Ability To Produce Shiga Toxin. Journal of Clinical Microbiology, 2002, 40, 2247-2248.	1.8	11
243	Virulence Factors in Escherichia coli. Journal of Clinical Microbiology, 2005, 43, 6221-6222.	1.8	11
244	Important Complexities of the Antivirulence Target Paradigm: A Novel Ostensibly Resistance-Avoiding Approach for Treating Infections: Table 1.. Journal of Infectious Diseases, 2016, 213, 901-903.	1.9	11
245	Evaluation of CTX-M steady-state mRNA, mRNA half-life and protein production in various STs of Escherichia coli. Journal of Antimicrobial Chemotherapy, 2016, 71, 607-616.	1.3	11
246	Variation in Resistance Traits, Phylogenetic Backgrounds, and Virulence Genotypes among Escherichia coli Clinical Isolates from Adjacent Hospital Campuses Serving Distinct Patient Populations. Antimicrobial Agents and Chemotherapy, 2015, 59, 5331-5339.	1.4	10
247	Complete Genome Sequence of a CTX-M-15-Producing Escherichia coli Strain from the H30R Subclone of Sequence Type 131 from a Patient with Recurrent Urinary Tract Infections, Closely Related to a Lethal Urosepsis Isolate from the Patient's Sister. Genome Announcements, 2016, 4, .	0.8	10
248	Activity of Eravacycline against Escherichia coli Clinical Isolates Collected from U.S. Veterans in 2011 in Relation to Coresistance Phenotype and Sequence Type 131 Genotype. Antimicrobial Agents and Chemotherapy, 2016, 60, 1888-1891.	1.4	10
249	Inactivation of Transcriptional Regulators during Within-Household Evolution of Escherichia coli. Journal of Bacteriology, 2017, 199, .	1.0	10
250	Large Fecal Reservoir of Escherichia coli Sequence Type 131-H30 Subclone Strains That Are Shared Within Households and Resemble Clinical ST131-H30 Isolates. Journal of Infectious Diseases, 2020, 221, 1659-1668.	1.9	10
251	Activity of meropenem/vaborbactam against international carbapenem-resistant Escherichia coli isolates in relation to clonal background, resistance genes, resistance to comparators and region. Journal of Global Antimicrobial Resistance, 2021, 24, 190-197.	0.9	10
252	Host-pathogen interactions in Escherichia coli urinary tract infection. Current Opinion in Infectious Diseases, 1994, 7, 287-294.	1.3	9

#	ARTICLE	IF	CITATIONS
253	Escherichia coli ST131: Variations on a theme of clonal expansion. Enfermedades Infecciosas Y Microbiología Clínica, 2013, 31, 355-356.	0.3	9
254	The Niche for Escherichia coli Sequence Type 131 Among Veterans: Urinary Tract Abnormalities and Long-Term Care Facilities. Open Forum Infectious Diseases, 2016, 3, ofw138.	0.4	9
255	Escherichia coli Sequence Type 131 <i>H</i> 30 Is the Main Driver of Emerging Extended-Spectrum-β-Lactamase-Producing E. coli at a Tertiary Care Center. MSphere, 2016, 1, .	1.3	9
256	Bacteriuria/Pyuria of Clinically Undetermined Significance (BPCUS): Common, but Currently Nameless. American Journal of Medicine, 2017, 130, e201-e204.	0.6	9
257	Clinical and Molecular Correlates of Escherichia coli Bloodstream Infection from Two Geographically Diverse Centers in Rochester, Minnesota, and Singapore. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	9
258	Analysis of urine-specific antibiograms from veterans to guide empiric therapy for suspected urinary tract infection. Diagnostic Microbiology and Infectious Disease, 2019, 95, 114874.	0.8	9
259	Intestinal Persistence of Colonizing <i>Escherichia coli</i> Strains, Especially ST131- <i>H</i> 30, in Relation to Bacterial and Host Factors. Journal of Infectious Diseases, 2022, 225, 2197-2207.	1.9	9
260	Evolution of Pathogenic Escherichia coli. , 2002, , 55-77.		8
261	Clinical and Molecular Epidemiology of Escherichia coli Sequence Type 131 among Hospitalized Patients Colonized Intestinally with Fluoroquinolone-Resistant E. coli. Antimicrobial Agents and Chemotherapy, 2014, 58, 7003-7006.	1.4	8
262	Epidemiology and characteristics of Escherichia coli sequence type 131 (ST131) from long-term care facility residents colonized intestinally with fluoroquinolone-resistant Escherichia coli. Diagnostic Microbiology and Infectious Disease, 2017, 87, 275-280.	0.8	8
263	Inactivation of extraintestinal pathogenic E. coli suspended in ground chicken meat by high pressure processing and identification of virulence factors which may affect resistance to high pressure. Food Control, 2020, 111, 107070.	2.8	8
264	Successful Management of a Serious Group A Streptococcal Infection During the Third Trimester of Pregnancy. Clinical Infectious Diseases, 1995, 21, 1058-1059.	2.9	7
265	Fluoroquinolone-resistant extraintestinal Escherichia coli clinical isolates representing the O15:K52:H1 clonal group from humans and dogs in Australia. Comparative Immunology, Microbiology and Infectious Diseases, 2012, 35, 319-324.	0.7	7
266	Prevalence of Rectal Colonization with Multidrug-Resistant Enterobacteriaceae among International Patients Hospitalized at Mayo Clinic, Rochester, Minnesota. Infection Control and Hospital Epidemiology, 2014, 35, 182-186.	1.0	7
267	Extraintestinal Pathogenic and Antimicrobial-Resistant Escherichia coli Contamination of 56 Public Restrooms in the Greater Minneapolis-St. Paul Metropolitan Area. Applied and Environmental Microbiology, 2015, 81, 4498-4506.	1.4	7
268	Isolation and Characterization of Escherichia coli Sequence Type 131 and Other Antimicrobial-Resistant Gram-Negative Bacilli from Clinical Stool Samples from Veterans. Antimicrobial Agents and Chemotherapy, 2016, 60, 4638-4645.	1.4	7
269	Definitions of Complicated Urinary Tract Infection and Pyelonephritis. Clinical Infectious Diseases, 2017, 64, 390-390.	2.9	7
270	Thermal inactivation of extraintestinal pathogenic Escherichia coli suspended in ground chicken meat. Food Control, 2019, 104, 269-277.	2.8	7

#	ARTICLE	IF	CITATIONS
271	Virulence genes and subclone status as markers of experimental virulence in a murine sepsis model among <i>Escherichia coli</i> sequence type 131 clinical isolates from Spain. <i>PLoS ONE</i> , 2017, 12, e0188838.	1.1	7
272	Community-Associated <i>Escherichia coli</i> Harboring CTX-M β -Lactamases from Urine Cultures from Pediatric Patients. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 2209-2210.	1.4	6
273	Activity of plazomicin against carbapenem-intermediate or -resistant <i>Escherichia coli</i> isolates from the United States and international sites in relation to clonal background, resistance genes, co-resistance, and region. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2061-2070.	1.3	5
274	Delayed Treatment of Bacterial Meningitis. <i>Annals of Internal Medicine</i> , 1999, 131, 715.	2.0	4
275	Draft Genomic Sequencing of Six Potential Extraintestinal Pathogenic <i>Escherichia coli</i> Isolates from Retail Chicken Meat. <i>Genome Announcements</i> , 2018, 6, .	0.8	4
276	Bacterial "Virulence" Traits and Host Demographics Predict <i>Escherichia coli</i> Colonization Behaviors Within Households. <i>Open Forum Infectious Diseases</i> , 2020, 7, ofaa495.	0.4	4
277	Visual prototypes in the ventral stream are attuned to complexity and gaze behavior. <i>Nature Communications</i> , 2021, 12, 6723.	5.8	4
278	Evaluating new therapies in gastrointestinal stromal tumor using in vivo molecular optical imaging. <i>Cancer Biology and Therapy</i> , 2014, 15, 911-918.	1.5	3
279	Reservoirs of Extraintestinal Pathogenic <i>Escherichia coli</i> . , 0, , 159-177.		3
280	Acute Pyelonephritis in Adults. <i>New England Journal of Medicine</i> , 2018, 378, 1162-1162.	13.9	3
281	<i>Escherichia coli</i> Clonal Group A. <i>Clinical Infectious Diseases</i> , 2005, 41, 568-568.	2.9	2
282	BACTERIAL CHARACTERISTICS AS PREDICTORS OF POSTTHERAPY RECURRENT BACTERIURIA AMONG CHILDREN WITH ACUTE UNCOMPLICATED CYSTITIS CAUSED BY <i>ESCHERICHIA COLI</i> . <i>Pediatric Infectious Disease Journal</i> , 2007, 26, 1151-1153.	1.1	2
283	Extraintestinal Pathogenic <i>Escherichia coli</i> . , 2009, , 939-961.		2
284	Activity of ceftolozane-tazobactam against <i>Escherichia coli</i> isolates from U.S. veterans (2011) in relation to co-resistance and sequence type 131 (ST131) H30 and H30Rx status. <i>PLoS ONE</i> , 2018, 13, e0200442.	1.1	2
285	Activity of ceftazidime-avibactam against <i>Escherichia coli</i> isolates from U.S. veterans (2011) in relation to co-resistance and sequence type 131 (ST131) H30 and H30Rx status. <i>Diagnostic Microbiology and Infectious Disease</i> , 2020, 97, 115034.	0.8	2
286	Infectious Pneumothorax Due to <i>Clostridium septicum</i> . <i>Journal of General Internal Medicine</i> , 2020, 35, 2197-2198.	1.3	2
287	Comparative activity of plazomicin against extended-spectrum cephalosporin-resistant <i>Escherichia coli</i> clinical isolates (2012-2017) in relation to phylogenetic background, sequence type 131 subclones, blaCTX-M genotype, and resistance to comparator agents. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> . 2021, 40, 2069-2075.	1.3	2
288	Core Genome Multi-Locus Sequence Typing and Prediction of Antimicrobial Susceptibility Using Whole Genome Sequences of <i>Escherichia coli</i> Bloodstream Infection Isolates. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, e0113921.	1.4	2

#	ARTICLE	IF	CITATIONS
289	Occurrence and potential transmission of extended-spectrum beta-lactamase-producing extraintestinal pathogenic and enteropathogenic <i>Escherichia coli</i> in domestic dog faeces from Minnesota. <i>Zoonoses and Public Health</i> , 2022, 69, 888-895.	0.9	2
290	Investigation at a Veterans Affairs Medical Center of Spurious Legionella Environmental Testing Results and High Laboratory-to-Laboratory Variability Among Four Commercial Laboratories. <i>Open Forum Infectious Diseases</i> , 2016, 3, .	0.4	1
291	Treatment of ED patients with bacteriuria/pyuria of clinically undetermined significance. <i>American Journal of Emergency Medicine</i> , 2017, 35, 1561.	0.7	1
292	Predictive characteristics of methicillin-resistant <i>Staphylococcus aureus</i> nares screening tests for methicillin resistance among <i>S. aureus</i> clinical isolates from hospitalized veterans. <i>Infection Control and Hospital Epidemiology</i> , 2019, 40, 603-605.	1.0	1
293	Draft genome sequences of concurrent <i>Escherichia coli</i> blood and fecal isolates from a patient with bacteremia and diarrhea belie BioFire-based detection of fecal enteropathogenic <i>E. coli</i> . <i>Pathogens and Disease</i> , 2020, 78, .	0.8	1
294	Complete Genome Sequence of <i>Escherichia coli</i> Strain FEX669, a ColV Plasmid-Containing Isolate from Retail Chicken Meat. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.3	1
295	Molecular Characteristics, Ecology, and Zoonotic Potential of <i>Escherichia coli</i> Strains That Cause Hemorrhagic Pneumonia in Animals. <i>Applied and Environmental Microbiology</i> , 2021, 87, e0147121.	1.4	1
296	Molecular Epidemiology and Population Genetics of Extraintestinal Pathogenic <i>Escherichia coli</i> . <i>JAMA</i> , 2003, 290, 91-107.		1
297	Effect of 7 vs 14 Days of Antibiotics Among Afebrile Men With Urinary Tract Infection—Reply. <i>JAMA - Journal of the American Medical Association</i> , 2021, 326, 2080.	3.8	1
298	Novel Multiplex PCR Method and Genome Sequence-Based Analog for High-Resolution Subclonal Assignment and Characterization of <i>Escherichia coli</i> Sequence Type 131 Isolates. <i>Microbiology Spectrum</i> , 2022, 10, .	1.2	1
299	Prevalence and characteristics of multidrug-resistant <i>Escherichia coli</i> sequence type ST131 at two academic centers in Boston and Minneapolis, USA. <i>American Journal of Infection Control</i> , 2023, 51, 434-439.	1.1	1
300	Clonal identity of <i>Escherichia coli</i> as a predictive factor for recurrent cystitis. <i>Virulence</i> , 2011, 2, 495-497.	1.8	0
301	Molecular Characterization of Environmental <i>Escherichia coli</i> Isolates from Public Restrooms in the Minneapolis-St. Paul Area. <i>Open Forum Infectious Diseases</i> , 2014, 1, S365-S365.	0.4	0
302	Response to Giufre et al. <i>Journal of Infectious Diseases</i> , 2014, 209, 630-631.	1.9	0
303	Molecular Characterization of Fecal <i>Escherichia coli</i> Isolates From Households (HHs) of Veterans Within Which One or More Household (HH) Members Carries <i>E. coli</i> Sequence Type 131 (ST131) or Other Fluoroquinolone (FQ)-Resistant <i>E. coli</i> (FQREC). <i>Open Forum Infectious Diseases</i> , 2016, 3, .	0.4	0
304	Confusion and Bacteriuria in Long-Term Care Facility Residents. <i>Journal of the American Geriatrics Society</i> , 2018, 66, 1235-1235.	1.3	0
305	Asymptomatic Bacteriuria. <i>Clinical Infectious Diseases</i> , 2018, 66, 1816-1817.	2.9	0
306	2609. <i>Escherichia coli</i> Clonal Lineages and Virulence Factors Predict Fecal Colonization within Households. <i>Open Forum Infectious Diseases</i> , 2019, 6, S907-S907.	0.4	0

#	ARTICLE	IF	CITATIONS
307	1438. Escherichia coli (EC) ST131-H30 Clonal Group is Associated with Antimicrobial Resistance, Illness Severity, Host Compromise, and Non-Cure among Patients with Bacteriuria. Open Forum Infectious Diseases, 2019, 6, S525-S525.	0.4	0
308	2583. Short-term Impact of Antimicrobial Exposure on Fecal Carriage of Resistant Microorganisms. Open Forum Infectious Diseases, 2019, 6, S897-S897.	0.4	0
309	1429. Diagnosis and Management of Osteomyelitis Associated with Stage IV Pressure Ulcers: Report of a Query to the Emerging Infections Network of the Infectious Diseases Society of America. Open Forum Infectious Diseases, 2019, 6, S521-S522.	0.4	0
310	A novel ergonomic wheelchair reduces bacterial hand contamination. Disability and Rehabilitation: Assistive Technology, 2020, , 1-4.	1.3	0
311	Pyelonephritis and abscesses of the kidney. , 2010, , 605-614.		0
312	184. Seven vs. 14 Days Treatment Duration for Afebrile Men with Urinary Tract Infections; A Randomized Clinical Trial. Open Forum Infectious Diseases, 2020, 7, S220-S221.	0.4	0
313	Draft Genome Sequences of Sixteen Fluoroquinolone-Resistant Extraintestinal Escherichia coli Isolates from Human Patients. Microbiology Resource Announcements, 2022, , e0000322.	0.3	0