James R Johnson

List of Publications by Year in descending order

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313 papers 24,179 citations

83 h-index 9861 141 g-index

317 all docs

 $\begin{array}{c} 317 \\ \text{docs citations} \end{array}$

317 times ranked

12994 citing authors

#	Article	IF	CITATIONS
1	Extended Virulence Genotypes of (i) Escherichia coli (i) Strains from Patients with Urosepsis in Relation to Phylogeny and Host Compromise. Journal of Infectious Diseases, 2000, 181, 261-272.	4.0	1,091
2	Guidelines for Antimicrobial Treatment of Uncomplicated Acute Bacterial Cystitis and Acute Pyelonephritis in Women. Clinical Infectious Diseases, 1999, 29, 745-758.	5.8	1,005
3	Organised Genome Dynamics in the Escherichia coli Species Results in Highly Diverse Adaptive Paths. PLoS Genetics, 2009, 5, e1000344.	3. 5	1,005
4	Intercontinental emergence of Escherichia coli clone O25:H4-ST131 producing CTX-M-15. Journal of Antimicrobial Chemotherapy, 2007, 61, 273-281.	3.0	737
5	Medical and economic impact of extraintestinal infections due to Escherichia coli: focus on an increasingly important endemic problem. Microbes and Infection, 2003, 5, 449-456.	1.9	649
6	Widespread Distribution of Urinary Tract Infections Caused by a Multidrug-ResistantEscherichia coliClonal Group. New England Journal of Medicine, 2001, 345, 1007-1013.	27.0	470
7	<i>Escherichia coli</i> Sequence Type ST131 as the Major Cause of Serious Multidrugâ€Resistant <i>E. coli</i> Infections in the United States. Clinical Infectious Diseases, 2010, 51, 286-294.	5.8	457
8	The Epidemic of Extended-Spectrum- \hat{l}^2 -Lactamase-Producing Escherichia coli ST131 Is Driven by a Single Highly Pathogenic Subclone, <i>H</i> 30-Rx. MBio, 2013, 4, e00377-13.	4.1	380
9	Phylogenetic Distribution of Extraintestinal Virulenceâ€Associated Traits inEscherichia coli. Journal of Infectious Diseases, 2001, 183, 78-88.	4.0	356
10	The Genome Sequence of Avian Pathogenic Escherichia coli Strain O1:K1:H7 Shares Strong Similarities with Human Extraintestinal Pathogenic E. coli Genomes. Journal of Bacteriology, 2007, 189, 3228-3236.	2.2	342
11	Extraintestinal pathogenic Escherichia coli : "The other bad E coli ― Translational Research, 2002, 139, 155-162.	2.3	326
12	Isolation and Molecular Characterization of Nalidixic Acid-Resistant Extraintestinal Pathogenic Escherichia coli from Retail Chicken Products. Antimicrobial Agents and Chemotherapy, 2003, 47, 2161-2168.	3.2	301
13	Evolutionary History of the Global Emergence of the Escherichia coli Epidemic Clone ST131. MBio, 2016, 7, e02162.	4.1	289
14	Systematic Review: Antimicrobial Urinary Catheters To Prevent Catheter-Associated Urinary Tract Infection in Hospitalized Patients. Annals of Internal Medicine, 2006, 144, 116.	3.9	276
15	Comparison of Extraintestinal Pathogenic <i>Escherichia coli</i> Strains from Human and Avian Sources Reveals a Mixed Subset Representing Potential Zoonotic Pathogens. Applied and Environmental Microbiology, 2008, 74, 7043-7050.	3.1	256
16	Food-Borne Origins of Escherichia coli Causing Extraintestinal Infections. Clinical Infectious Diseases, 2012, 55, 712-719.	5.8	255
17	Epidemic Clonal Groups of <i>Escherichia coli</i> as a Cause of Antimicrobial-Resistant Urinary Tract Infections in Canada, 2002 to 2004. Antimicrobial Agents and Chemotherapy, 2009, 53, 2733-2739.	3.2	249
18	Abrupt Emergence of a Single Dominant Multidrug-Resistant Strain of Escherichia coli. Journal of Infectious Diseases, 2013, 207, 919-928.	4.0	247

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19	Extended-Spectrum β-Lactamase–Producing Escherichia coli From Retail Chicken Meat and Humans: Comparison of Strains, Plasmids, Resistance Genes, and Virulence Factors. Clinical Infectious Diseases, 2013, 56, 478-487.	5.8	233
20	Phylogenetic Origin and Virulence Genotype in Relation to Resistance to Fluoroquinolones and/or Extended‧pectrum Cephalosporins and Cephamycins amongEscherichia colilsolates from Animals and Humans. Journal of Infectious Diseases, 2003, 188, 759-768.	4.0	227
21	Escherichia coli Isolates That Carry <i>vat</i> , <i>fyuA</i> , <i>chuA</i> , and <i>yfcV</i> Efficiently Colonize the Urinary Tract. Infection and Immunity, 2012, 80, 4115-4122.	2.2	226
22	Loop-Mediated Isothermal Amplification Assay for Rapid Detection of Common Strains of <i>Escherichia coli</i> . Journal of Clinical Microbiology, 2008, 46, 2800-2804.	3.9	225
23	Antimicrobialâ€Resistant and Extraintestinal PathogenicEscherichia coliin Retail Foods. Journal of Infectious Diseases, 2005, 191, 1040-1049.	4.0	223
24	Molecular epidemiology of extraintestinal pathogenic (uropathogenic) Escherichia coli. International Journal of Medical Microbiology, 2005, 295, 383-404.	3.6	218
25	A New Clone Sweeps Clean: the Enigmatic Emergence of Escherichia coli Sequence Type 131. Antimicrobial Agents and Chemotherapy, 2014, 58, 4997-5004.	3.2	207
26	Antimicrobial Drug–Resistant <i>Escherichia coli</i> from Humans and Poultry Products, Minnesota and Wisconsin, 2002–2004. Emerging Infectious Diseases, 2007, 13, 838-846.	4.3	190
27	Escherichia coli ST131- <i>H</i> 22 as a Foodborne Uropathogen. MBio, 2018, 9, .	4.1	184
28	<i>In Silico</i> Genotyping of Escherichia coli Isolates for Extraintestinal Virulence Genes by Use of Whole-Genome Sequencing Data. Journal of Clinical Microbiology, 2020, 58, .	3.9	179
29	High-Resolution Two-Locus Clonal Typing of Extraintestinal Pathogenic Escherichia coli. Applied and Environmental Microbiology, 2012, 78, 1353-1360.	3.1	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. Journal of Clinical Microbiology, 2008, 46, 2529-2534.	3.9	169
31	Escherichia coli Sequence Type 131 (ST131) Subclone H30 as an Emergent Multidrug-Resistant Pathogen Among US Veterans. Clinical Infectious Diseases, 2013, 57, 1256-1265.	5.8	167
32	Molecular Epidemiology and Phylogenetic Distribution of the <i>Escherichia coli pks</i> Genomic Island. Journal of Clinical Microbiology, 2008, 46, 3906-3911.	3.9	157
33	Microbial virulence determinants and the pathogenesis of urinary tract infection. Infectious Disease Clinics of North America, 2003, 17, 261-278.	5.1	156
34	Zoonotic Potential of Escherichia coli Isolates from Retail Chicken Meat Products and Eggs. Applied and Environmental Microbiology, 2015, 81, 1177-1187.	3.1	156
35	Extended Virulence Genotypes and Phylogenetic Background of Escherichia colilsolates from Patients with Cystitis, Pyelonephritis, or Prostatitis. Journal of Infectious Diseases, 2005, 191, 46-50.	4.0	151
36	Molecular Epidemiological and Phylogenetic Associations of Two Novel Putative Virulence Genes, iha and iroN E. coli, among Escherichia coli Isolates from Patients with Urosepsis. Infection and Immunity, 2000, 68, 3040-3047.	2.2	150

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37	Phylogenetic Distribution of Virulenceâ€Associated Genes amongEscherichia colilsolates Associated with Neonatal Bacterial Meningitis in The Netherlands. Journal of Infectious Diseases, 2002, 185, 774-784.	4.0	150
38	Experimental Mouse Lethality of Escherichia colilsolates, in Relation to Accessory Traits, Phylogenetic Group, and Ecological Source. Journal of Infectious Diseases, 2006, 194, 1141-1150.	4.0	146
39	Acquisition of Avian Pathogenic Escherichia coli Plasmids by a Commensal E. coli Isolate Enhances Its Abilities To Kill Chicken Embryos, Grow in Human Urine, and Colonize the Murine Kidney. Infection and Immunity, 2006, 74, 6287-6292.	2.2	145
40	The genetic structure of Escherichia coli populations in primary and secondary habitats. Microbiology (United Kingdom), 2002, 148, 1513-1522.	1.8	142
41	Similarity between Human and ChickenEscherichia colilsolates in Relation to Ciprofloxacin Resistance Status. Journal of Infectious Diseases, 2006, 194, 71-78.	4.0	138
42	<i>Escherichia coli</i> Colonization Patterns among Human Household Members and Pets, with Attention to Acute Urinary Tract Infection. Journal of Infectious Diseases, 2008, 197, 218-224.	4.0	134
43	Rates of Mutation and Host Transmission for an Escherichia coli Clone over 3 Years. PLoS ONE, 2011, 6, e26907.	2.5	132
44	Quinolone, fluoroquinolone and trimethoprim/sulfamethoxazole resistance in relation to virulence determinants and phylogenetic background among uropathogenic Escherichia coli. Journal of Antimicrobial Chemotherapy, 2006, 57, 204-211.	3.0	131
45	Changes in Colonic Bile Acid Composition following Fecal Microbiota Transplantation Are Sufficient to Control Clostridium difficile Germination and Growth. PLoS ONE, 2016, 11, e0147210.	2.5	130
46	Multiple-Host Sharing, Long-Term Persistence, and Virulence of <i>Escherichia coli</i> Clones from Human and Animal Household Members. Journal of Clinical Microbiology, 2008, 46, 4078-4082.	3.9	127
47	Associations Between Multidrug Resistance, Plasmid Content, and Virulence Potential Among Extraintestinal Pathogenic and Commensal <i>Escherichia coli</i> from Humans and Poultry. Foodborne Pathogens and Disease, 2012, 9, 37-46.	1.8	126
48	Fimbrial Profiles Predict Virulence of Uropathogenic Escherichia coli Strains: Contribution of Ygi and Yad Fimbriae. Infection and Immunity, 2011, 79, 4753-4763.	2.2	121
49	<i>Escherichia coli</i> Sequence Type 131 Is a Dominant, Antimicrobial-Resistant Clonal Group Associated with Healthcare and Elderly Hosts. Infection Control and Hospital Epidemiology, 2013, 34, 361-369.	1.8	121
50	Identification of a New Iron-Regulated Virulence Gene, ireA , in an Extraintestinal Pathogenic Isolate of Escherichia coli. Infection and Immunity, 2001, 69, 6209-6216.	2.2	118
51	IroN Functions as a Siderophore Receptor and Is a Urovirulence Factor in an Extraintestinal Pathogenic Isolate of Escherichia coli. Infection and Immunity, 2002, 70, 7156-7160.	2.2	118
52	CTX-M-27- and CTX-M-14-producing, ciprofloxacin-resistant <i>Escherichia coli</i> of the <i>H</i> 30 subclonal group within ST131 drive a Japanese regional ESBL epidemic. Journal of Antimicrobial Chemotherapy, 2015, 70, 1639-1649.	3.0	118
53	Epidemiology of <i>Escherichia coli</i> Bacteremia: A Systematic Literature Review. Clinical Infectious Diseases, 2021, 72, 1211-1219.	5.8	116
54	Determination of Escherichia coli O types by allele-specific polymerase chain reaction: application to the O types involved in human septicemia. Diagnostic Microbiology and Infectious Disease, 2007, 57, 129-136.	1.8	115

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55	Evaluation of Escherichia coli isolates from healthy chickens to determine their potential risk to poultry and human health. PLoS ONE, 2017, 12, e0180599.	2.5	113
56	Commonality among Fluoroquinolone-Resistant Sequence Type ST131 Extraintestinal <i>Escherichia coli</i> Isolates from Humans and Companion Animals in Australia. Antimicrobial Agents and Chemotherapy, 2011, 55, 3782-3787.	3.2	112
57	Molecular Epidemiology of Escherichia coli Sequence Type 131 and Its H30 and H30-Rx Subclones among Extended-Spectrum-β-Lactamase-Positive and -Negative E. coli Clinical Isolates from the Chicago Region, 2007 to 2010. Antimicrobial Agents and Chemotherapy, 2013, 57, 6385-6388.	3.2	112
58	Extraintestinal Pathogenic Escherichia coli as a Cause of Invasive Nonurinary Infections. Journal of Clinical Microbiology, 2003, 41, 5798-5802.	3.9	111
59	Molecular Epidemiological Analysis of Escherichia coli Sequence Type ST131 (O25:H4) and <i>bla</i> _{CTX-M-15} among Extended-Spectrum-β-Lactamase-Producing E. coli from the United States, 2000 to 2009. Antimicrobial Agents and Chemotherapy, 2012, 56, 2364-2370.	3.2	107
60	Rapid and Specific Detection, Molecular Epidemiology, and Experimental Virulence of the O16 Subgroup within Escherichia coli Sequence Type 131. Journal of Clinical Microbiology, 2014, 52, 1358-1365.	3.9	107
61	Bacterial Characteristics in Relation to Clinical Source of Escherichia coli Isolates from Women with Acute Cystitis or Pyelonephritis and Uninfected Women. Journal of Clinical Microbiology, 2005, 43, 6064-6072.	3.9	106
62	Sharing of Virulent Escherichia coli Clones among Household Members of a Woman with Acute Cystitis. Clinical Infectious Diseases, 2006, 43, e101-e108.	5.8	106
63	Canine Feces as a Reservoir of Extraintestinal Pathogenic Escherichia coli. Infection and Immunity, 2001, 69, 1306-1314.	2.2	105
64	The IrgA Homologue Adhesin Iha Is an Escherichia coli Virulence Factor in Murine Urinary Tract Infection. Infection and Immunity, 2005, 73, 965-971.	2.2	105
65	A disseminated multidrug-resistant clonal group of uropathogenic Escherichia coli in pyelonephritis. Lancet, The, 2002, 359, 2249-2251.	13.7	104
66	Intermingled <i>Klebsiella pneumoniae </i> Populations Between Retail Meats and Human Urinary Tract Infections. Clinical Infectious Diseases, 2015, 61, 892-899.	5.8	104
67	Virulence Factor Profiles and Phylogenetic Background of Escherichia colilsolates from Veterans with Bacteremia and Uninfected Control Subjects. Journal of Infectious Diseases, 2004, 190, 2121-2128.	4.0	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. Journal of Clinical Microbiology, 2009, 47, 3721-3725.	3.9	102
69	Multidrug-resistant extraintestinal pathogenic Escherichia coli of sequence type ST131 in animals and foods. Veterinary Microbiology, 2011, 153, 99-108.	1.9	102
70	Virulence of Escherichia coli Clinical Isolates in a Murine Sepsis Model in Relation to Sequence Type ST131 Status, Fluoroquinolone Resistance, and Virulence Genotype. Infection and Immunity, 2012, 80, 1554-1562.	2.2	101
71	Host Characteristics and Bacterial Traits Predict Experimental Virulence for Escherichia coli Bloodstream Isolates From Patients With Urosepsis. Open Forum Infectious Diseases, 2015, 2, ofv083.	0.9	100
72	Ongoing Horizontal and Vertical Transmission of Virulence Genes and papA Alleles among Escherichia coliBlood Isolates from Patients with Diverse-Source Bacteremia. Infection and Immunity, 2001, 69, 5363-5374.	2,2	99

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73	Quinolone-Resistant Uropathogenic Escherichia coli Strains from Phylogenetic Group B2 Have Fewer Virulence Factors than Their Susceptible Counterparts. Journal of Clinical Microbiology, 2005, 43, 2962-2964.	3.9	99
74	Evidence of Commonality between Canine and Human Extraintestinal Pathogenic Escherichia coli Strains That Express papG Allele III. Infection and Immunity, 2000, 68, 3327-3336.	2.2	98
75	Separate F-Type Plasmids Have Shaped the Evolution of the $\langle i \rangle H \langle i \rangle$ 30 Subclone of Escherichia coli Sequence Type 131. MSphere, 2016, 1, .	2.9	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. Antimicrobial Agents and Chemotherapy, 1999, 43, 2990-2995.	3.2	96
77	Modulation of Host Innate Immune Response in the Bladder by Uropathogenic <i>Escherichia coli </i> Infection and Immunity, 2007, 75, 5353-5360.	2.2	96
78	Virulence Characteristics and Phylogenetic Background of Multidrugâ€Resistant and Antimicrobialâ€Susceptible Clinical Isolates of <1 > Escherichia coli < /1 > from across the United States, 2000–2001. Journal of Infectious Diseases, 2004, 190, 1739-1744.	4.0	95
79	Comparison of <i>Escherichia coli</i> ST131 Pulsotypes, by Epidemiologic Traits, 1967–2009. Emerging Infectious Diseases, 2012, 18, 598-607.	4.3	93
80	Selection Footprint in the FimH Adhesin Shows Pathoadaptive Niche Differentiation in Escherichia coli. Molecular Biology and Evolution, 2004, 21, 1373-1383.	8.9	91
81	Identification of Urovirulence Traits in Escherichia coli by Comparison of Urinary and Rectal E. coli Isolates from Dogs with Urinary Tract Infection. Journal of Clinical Microbiology, 2003, 41, 337-345.	3.9	89
82	Epidemiological Correlates of Virulence Genotype and Phylogenetic Background amongEscherichia coliBlood Isolates from Adults with Diverseâ€Source Bacteremia. Journal of Infectious Diseases, 2002, 185, 1439-1447.	4.0	88
83	Analysis of the F Antigen-Specific papA Alleles of Extraintestinal Pathogenic Escherichia coli Using a Novel Multiplex PCR-Based Assay. Infection and Immunity, 2000, 68, 1587-1599.	2.2	87
84	Identification of two previously unrecognized genes (guaAandargC) important for uropathogenesis. Molecular Microbiology, 1996, 22, 217-229.	2.5	86
85	Virulence Factors of Escherichia coli Isolates That Produce CTX-M-Type Extended-Spectrum β-Lactamases. Antimicrobial Agents and Chemotherapy, 2005, 49, 4667-4670.	3.2	85
86	Distribution and Characteristics of <i>Escherichia coli </i> Clonal Group A1. Emerging Infectious Diseases, 2005, 11, 141-145.	4.3	84
87	Contamination of Retail Foods, Particularly Turkey, from Community Markets (Minnesota, 1999–2000) with Antimicrobial-Resistant and Extraintestinal PathogenicEscherichia coli. Foodborne Pathogens and Disease, 2005, 2, 38-49.	1.8	84
88	Uropathogenic <i>Escherichia coli</i> Induces Chronic Pelvic Pain. Infection and Immunity, 2011, 79, 628-635.	2.2	83
89	Hepatitis Due to Herpes Simplex Virus in Marrow-Transplant Recipients. Clinical Infectious Diseases, 1992, 14, 38-45.	5.8	81
90	Rapid and Extensive Expansion in the United States of a New Multidrug-resistant <i>Escherichia coli</i> Clonal Group, Sequence Type 1193. Clinical Infectious Diseases, 2019, 68, 334-337.	5 . 8	81

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91	Improved Repetitive-Element PCR Fingerprinting for Resolving Pathogenic and Nonpathogenic Phylogenetic Groups within Escherichia coli. Vaccine Journal, 2000, 7, 265-273.	2.6	80
92	The Clonal Distribution and Diversity of Extraintestinal Escherichia coli Isolates Vary According to Patient Characteristics. Antimicrobial Agents and Chemotherapy, 2013, 57, 5912-5917.	3.2	80
93	The Pandemic H30 Subclone of Sequence Type 131 (ST131) as the Leading Cause of Multidrug-Resistant Escherichia coli Infections in the United States (2011–2012). Open Forum Infectious Diseases, 2017, 4, ofx089.	0.9	79
94	Phylogenetic relationships among clonal groups of extraintestinal pathogenic Escherichia coli as assessed by multi-locus sequence analysis. Microbes and Infection, 2006, 8, 1702-1713.	1.9	78
95	UropathogenicEscherichia colias Agents of Diverse Non–Urinary Tract Extraintestinal Infections. Journal of Infectious Diseases, 2002, 186, 859-864.	4.0	77
96	Enteroaggregative Escherichia coli O78:H10, the Cause of an Outbreak of Urinary Tract Infection. Journal of Clinical Microbiology, 2012, 50, 3703-3711.	3.9	77
97	Prevalence and Characteristics of the Epidemic Multiresistant Escherichia coli ST131 Clonal Group among Extended-Spectrum Beta-Lactamase-Producing E. coli Isolates in Copenhagen, Denmark. Journal of Clinical Microbiology, 2013, 51, 1779-1785.	3.9	77
98	Clonal analysis reveals high rate of structural mutations in fimbrial adhesins of extraintestinal pathogenic Escherichia coli. Molecular Microbiology, 2006, 59, 975-988.	2.5	76
99	Four Main Virotypes among Extended-Spectrum-β-Lactamase-Producing Isolates of Escherichia coli O25b:H4-B2-ST131: Bacterial, Epidemiological, and Clinical Characteristics. Journal of Clinical Microbiology, 2013, 51, 3358-3367.	3.9	76
100	Virulence Genotype and Phylogenetic Origin in Relation to Antibiotic Resistance Profile among Escherichia coli Urine Sample Isolates from Israeli Women with Acute Uncomplicated Cystitis. Antimicrobial Agents and Chemotherapy, 2005, 49, 26-31.	3.2	72
101	Reservoirs of Extraintestinal Pathogenic <i>Escherichia coli</i> li>. Microbiology Spectrum, 2015, 3, .	3.0	71
102	<i>Escherichia coli</i> Pyomyositis: An Emerging Infectious Disease among Patients with Hematologic Malignancies. Clinical Infectious Diseases, 2010, 50, 374-380.	5.8	70
103	Carbapenemase-producing bacteria in companion animals: a public health concern on the horizon. Journal of Antimicrobial Chemotherapy, 2014, 69, 1155-1157.	3.0	68
104	Intensity and Mechanisms of Fluoroquinolone Resistance within the $\langle i \rangle H \langle i \rangle$ 30 and $\langle i \rangle H \langle i \rangle$ 30Rx Subclones of Escherichia coli Sequence Type 131 Compared with Other Fluoroquinolone-Resistant E. coli. Antimicrobial Agents and Chemotherapy, 2015, 59, 4471-4480.	3.2	68
105	Virulence Genotypes and Phylogenetic Background of <i>Escherichia coli</i> Serogroup O6 Isolates from Humans, Dogs, and Cats. Journal of Clinical Microbiology, 2008, 46, 417-422.	3.9	67
106	Structure and urovirulence characteristics of the fecal Escherichia coli population among healthy women. Microbes and Infection, 2009, 11 , $274-280$.	1.9	67
107	Prominence of an O75 Clonal Group (Clonal Complex 14) among Non-ST131 Fluoroquinolone-Resistant Escherichia coli Causing Extraintestinal Infections in Humans and Dogs in Australia. Antimicrobial Agents and Chemotherapy, 2012, 56, 3898-3904.	3.2	66
108	Ciprofloxacin-Resistant Gram-Negative Bacilli in the Fecal Microflora of Children. Antimicrobial Agents and Chemotherapy, 2006, 50, 3325-3329.	3. 2	64

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109	Phylogenomic Analysis of Extraintestinal Pathogenic <i>Escherichia coli</i> Sequence Type 1193, an Emerging Multidrug-Resistant Clonal Group. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	64
110	Transmission of an Extended-Spectrum-Beta-Lactamase-Producing <i>Escherichia coli</i> (Sequence) Tj ETQq0 0 Pyelonephritis. Journal of Clinical Microbiology, 2009, 47, 3780-3782.	0 rgBT /O 3.9	verlock 10 Tf 63
111	Genetic Diversity and Virulence Profiles of <i>Escherichia coli</i> Isolates Causing Spontaneous Bacterial Peritonitis and Bacteremia in Patients with Cirrhosis. Journal of Clinical Microbiology, 2010, 48, 2709-2714.	3.9	63
112	<i>Escherichia coli</i> /i> Serotype O15:K52:H1 as a Uropathogenic Clone. Journal of Clinical Microbiology, 2000, 38, 201-209.	3.9	63
113	Phylogenetic Background and Virulence Profiles of Fluoroquinoloneâ€Resistant ClinicalEscherichia colilsolates from The Netherlands. Journal of Infectious Diseases, 2002, 186, 1852-1856.	4.0	62
114	Predictive Diagnostics for Escherichia coli Infections Based on the Clonal Association of Antimicrobial Resistance and Clinical Outcome. Journal of Clinical Microbiology, 2013, 51, 2991-2999.	3.9	62
115	Household Clustering of Escherichia coli Sequence Type 131 Clinical and Fecal Isolates According to Whole Genome Sequence Analysis. Open Forum Infectious Diseases, 2016, 3, ofw129.	0.9	62
116	Clonal Origin, Virulence Factors, and Virulence. Infection and Immunity, 2000, 68, 424-425.	2.2	61
117	Escherichia coli Sequence Type 131 as a Prominent Cause of Antibiotic Resistance among Urinary Escherichia coli Isolates from Reproductive-Age Women. Journal of Clinical Microbiology, 2013, 51, 3270-3276.	3.9	61
118	Rapid and Specific Detection of Escherichia coli Clonal Group A by Gene-Specific PCR. Journal of Clinical Microbiology, 2004, 42, 2618-2622.	3.9	60
119	<i>Escherichia coli</i> Sequence Type ST131 as an Emerging Fluoroquinolone-Resistant Uropathogen among Renal Transplant Recipients. Antimicrobial Agents and Chemotherapy, 2010, 54, 546-550.	3.2	60
120	Gut Colonization of Healthy Children and Their Mothers With Pathogenic Ciprofloxacin-Resistant <i>Escherichia coli</i> Iournal of Infectious Diseases, 2015, 212, 1862-1868.	4.0	60
121	Urinary Tract Infections. Primary Care - Clinics in Office Practice, 2008, 35, 345-367.	1.6	59
122	Phylogenetic diversity, antimicrobial susceptibility and virulence characteristics of phylogroup F Escherichia coli in Australia. Microbiology (United Kingdom), 2016, 162, 1904-1912.	1.8	59
123	Molecular Epidemiology of Extraintestinal Pathogenic <i>Escherichia coli</i> EcoSal Plus, 2018, 8, .	5.4	57
124	Iha from an Escherichia coli Urinary Tract Infection Outbreak Clonal Group A Strain Is Expressed In Vivo in the Mouse Urinary Tract and Functions as a Catecholate Siderophore Receptor. Infection and Immunity, 2006, 74, 3427-3436.	2,2	56
125	Clonal group distribution of fluoroquinolone-resistant Escherichia coli among humans and companion animals in Australia. Journal of Antimicrobial Chemotherapy, 2010, 65, 1936-1938.	3.0	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. Journal of Clinical Microbiology, 2019, 57, .	3.9	56

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127	Human-associated fluoroquinolone-resistant Escherichia coli clonal lineages, including ST354, isolated from canine feces and extraintestinal infections in Australia. Microbes and Infection, 2015, 17, 266-274.	1.9	55
128	<i>papG</i> Alleles among <i>Escherichia coli</i> Strains Causing Urosepsis: Associations with Other Bacterial Characteristics and Host Compromise. Infection and Immunity, 1998, 66, 4568-4571.	2.2	55
129	Predictors and Molecular Epidemiology of Community-Onset Extended-Spectrum β-Lactamase–Producing ⟨i⟩Escherichia coli⟨/i⟩ Infection in a Midwestern Community. Infection Control and Hospital Epidemiology, 2013, 34, 947-953.	1.8	54
130	Colonisation with ESBL-producing and carbapenemase-producing Enterobacteriaceae, vancomycin-resistant enterococci, and meticillin-resistant Staphylococcus aureus in a long-term care facility over one year. BMC Infectious Diseases, 2015, 15, 168.	2.9	54
131	Global Molecular Epidemiology of the O15:K52:H1 Extraintestinal Pathogenic Escherichia coli Clonal Group: Evidence of Distribution beyond Europe. Journal of Clinical Microbiology, 2002, 40, 1913-1923.	3.9	53
132	Phylogenetic and Pathotypic Comparison of Concurrent Urine and Rectal Escherichia coli Isolates from Men with Febrile Urinary Tract Infection. Journal of Clinical Microbiology, 2005, 43, 3895-3900.	3.9	51
133	Three-Decade Epidemiological Analysis of <i>Escherichia coli</i> O15:K52:H1. Journal of Clinical Microbiology, 2009, 47, 1857-1862.	3.9	51
134	Presence of Putative Repeat-in-Toxin Gene <i>tosA</i> in Escherichia coli Predicts Successful Colonization of the Urinary Tract. MBio, 2011, 2, e00066-11.	4.1	51
135	Occurrence of Antibiotic-Resistant Uropathogenic Escherichia coli Clonal Group A in Wastewater Effluents. Applied and Environmental Microbiology, 2007, 73, 4180-4184.	3.1	50
136	Role of Homologous Recombination in Adaptive Diversification of Extraintestinal Escherichia coli. Journal of Bacteriology, 2013, 195, 231-242.	2.2	50
137	Risk Factors for Trimethoprim-Sulfamethoxazole Resistance in Patients with Acute Uncomplicated Cystitis. Antimicrobial Agents and Chemotherapy, 2008, 52, 846-851.	3.2	49
138	Clinical and Microbiological Determinants of Infection After Transrectal Prostate Biopsy. Clinical Infectious Diseases, 2015, 60, 979-987.	5.8	49
139	Genetic Structure and Antimicrobial Resistance of Escherichia coli and Cryptic Clades in Birds with Diverse Human Associations. Applied and Environmental Microbiology, 2015, 81, 5123-5133.	3.1	49
140	Molecular Analysis of <i>Escherichia coli</i> from Retail Meats (2002–2004) from the United States National Antimicrobial Resistance Monitoring System. Clinical Infectious Diseases, 2009, 49, 195-201.	5.8	48
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