Richard Goering

List of Publications by Year
in descending order

64 papers	4,145 citations	159585 30 h-index	123424 61 g-index
65 all docs	65 docs citations	65 times ranked	4898 citing authors

#	Article	IF	Citations
1	Methicillin-resistant Staphylococcus aureus strain USA300: origin and epidemiology. Journal of Antimicrobial Chemotherapy, 2009, 64, 441-446.	3.0	380
2	Whole genome sequencing options for bacterial strain typing and epidemiologic analysis based on single nucleotide polymorphism versus gene-by-geneâ€“based approaches. Clinical Microbiology and Infection, 2018, 24, 350-354.	6.0	373
3	Molecular Typing of Methicillin-ResistantStaphylococcus aureusby Pulsed-Field Gel Electrophoresis: Comparison of Results Obtained in a Multilaboratory Effort Using Identical Protocols and MRSA Strains. Microbial Drug Resistance, 2000, 6, 189-198.	2.0	267
4	Bacterial Whole-Genome Sequencing Revisited: Portable, Scalable, and Standardized Analysis for Typing and Detection of Virulence and Antibiotic Resistance Genes. Journal of Clinical Microbiology, 2014, 52, 2365-2370.	3.9	250
5	Frequent emergence and limited geographic dispersal of methicillin-resistant Staphylococcus aureus. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 14130-14135.	7.1	239
6	Pulsed field gel electrophoresis: A review of application and interpretation in the molecular epidemiology of infectious disease. Infection, Genetics and Evolution, 2010, 10, 866-875.	2.3	200
7	Assessment of Resolution and Intercenter Reproducibility of Results of Genotyping Staphylococcus aureus by Pulsed-Field Gel Electrophoresis of Sma I Macrorestriction Fragments: a Multicenter Study. Journal of Clinical Microbiology, 1998, 36, 1653-1659.	3.9	186
8	Impact of Strain Type on Detection of Toxigenic Clostridium difficile: Comparison of Molecular Diagnostic and Enzyme Immunoassay Approaches. Journal of Clinical Microbiology, 2010, 48, 3719-3724.	3.9	177
9	Guidelines for Reporting Novel mecA Gene Homologues. Antimicrobial Agents and Chemotherapy, 2012, 56, 4997-4999.	3.2	144
10	Phenotypic and Enzymatic Comparative Analysis of the Novel KPC Variant KPC-5 and Its Evolutionary Variants, KPC-2 and KPC-4. Antimicrobial Agents and Chemotherapy, 2009, 53, 557-562.	3.2	119
11	Molecular Epidemiology of Methicillin-Resistant and Methicillin-Susceptible Staphylococcus aureus Isolates from Global Clinical Trials. Journal of Clinical Microbiology, 2008, 46, 2842-2847.	3.9	113
12	Characterization of Nasal and Blood Culture Isolates of Methicillin-Resistant Staphylococcus aureus from Patients in United States Hospitals. Antimicrobial Agents and Chemotherapy, 2012, 56, 1324-1330.	3.2	105
13	Strain Types and Antimicrobial Resistance Patterns of Clostridium difficile Isolates from the United States, 2011 to 2013. Antimicrobial Agents and Chemotherapy, 2014, 58, 4214-4218.	3.2	103
14	Usefulness of mec-associated direct repeat unit (dru) typing in the epidemiological analysis of highly clonal methicillin-resistant Staphylococcus aureus in Scotland. Clinical Microbiology and Infection, 2008, 14, 964-969.	6.0	92
15	Microbial Typing by Matrix-Assisted Laser Desorption Ionizationâ€“Time of Flight Mass Spectrometry: Do We Need Guidance for Data Interpretation?. Journal of Clinical Microbiology, 2015, 53, 760-765.	3.9	92
16	Outbreak of Carbapenem-Resistant Klebsiella pneumoniae in Puerto Rico Associated with a Novel Carbapenemase Variant. Infection Control and Hospital Epidemiology, 2010, 31, 476-484.	1.8	89
17	Surveillance of Carbapenem-Resistant Pseudomonas aeruginosa Isolates from Puerto Rican Medical Center Hospitals: Dissemination of KPC and IMP-18 Î²-Lactamases. Antimicrobial Agents and Chemotherapy, 2009, 53, 1660-1664.	3.2	88
18	Comparison of Strain Typing Results for Clostridium difficile Isolates from North America. Journal of Clinical Microbiology, 2011, 49, 1831-1837.	3.9	86

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19	Characterization of a Novel Arginine Catabolic Mobile Element (ACME) and Staphylococcal Chromosomal Cassette mec Composite Island with Significant Homology to Staphylococcus epidermidis ACME Type II in Methicillin-Resistant Staphylococcus aureus Genotype ST22-MRSA-IV. Antimicrobial Agents and Chemotherapy, 2011, 55, 1896-1905.	3.2	83
20	Transduction of Staphylococcal Cassette Chromosome mec Elements between Strains of Staphylococcus aureus. Antimicrobial Agents and Chemotherapy, 2013, 57, 5233-5238.	3.2	64
21	Wide geographical dissemination of the multiresistant Staphylococcus capitis NRCS-A clone in neonatal intensive-care units. Clinical Microbiology and Infection, 2016, 22, 46-52.	6.0	58
22	Enhanced Discrimination of Highly Clonal ST22-Methicillin-Resistant Staphylococcus aureus IV Isolates Achieved by Combining spa , dru , and Pulsed-Field Gel Electrophoresis Typing Data. Journal of Clinical Microbiology, 2010, 48, 1839-1852.	3.9	55
23	Evolution and Global Transmission of a Multidrug-Resistant, Community-Associated Methicillin-Resistant Staphylococcus aureus Lineage from the Indian Subcontinent. MBio, 2019, 10, .	4.1	50
24	Two Distinct Clones of Methicillin-Resistant Staphylococcus aureus (MRSA) with the Same USA300 Pulsed-Field Gel Electrophoresis Profile: a Potential Pitfall for Identification of USA300 Community-Associated MRSA. Journal of Clinical Microbiology, 2009, 47, 3765-3768.	3.9	46
25	Activity of Tedizolid (TR-700) against Well-Characterized Methicillin-Resistant Staphylococcus aureus Strains of Diverse Epidemiological Origins. Antimicrobial Agents and Chemotherapy, 2013, 57, 2892-2895.	3.2	44
26	Prevalence of Toxic Shock Syndrome Toxin 1 (TSST-1)-Producing Strains of Staphylococcus aureus and Antibody to TSST-1 among Healthy Japanese Women. Journal of Clinical Microbiology, 2008, 46, 2731-2738.	3.9	43
27	Subpopulations of Staphylococcus aureus Clonal Complex 121 Are Associated with Distinct Clinical Entities. PLoS ONE, 2013, 8, e58155.	2.5	43
28	Molecular characterization of methicillin-resistant Staphylococcus aureus isolated over a 2-year period in a Qatari hospital from multinational patients. Clinical Microbiology and Infection, 2014, 20, 169-173.	6.0	40
29	Changes in molecular epidemiology and antimicrobial resistance profiles of Clostridioides (Clostridium) difficile strains in the United States between 2011 and 2017. Anaerobe, 2019, 60, 102050.	2.1	35
30	Linezolid-Resistant Staphylococcus aureus Strain 1128105, the First Known Clinical Isolate Possessing thecfrMultidrug Resistance Gene. Antimicrobial Agents and Chemotherapy, 2014, 58, 6592-6598.	3.2	34
31	Emergence of Oxacillin Resistance in Stealth Methicillin-Resistant Staphylococcus aureus Due to mecA Sequence Instability. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	34
32	Emergence of Carbapenem Resistance inPseudomonas aeruginosaIsolates from a Patient with Cystic Fibrosis in the Absence of Carbapenem Therapy. Clinical Infectious Diseases, 2008, 46, e137-e141.	5.8	31
33	Multicenter Evaluation of the Verigene Clostridium difficile Nucleic Acid Assay. Journal of Clinical Microbiology, 2013, 51, 4120-4125.	3.9	31
34	Stenotrophomonas maltophilia in Malaysia: molecular epidemiology and trimethoprimâ€“sulfamethoxazole resistance. International Journal of Infectious Diseases, 2012, 16, e603-e607.	3.3	30
35	Continued expansion of USA300-like methicillin-resistant Staphylococcus aureus (MRSA) among hospitalized patients in the United States. Diagnostic Microbiology and Infectious Disease, 2017, 88, 342-347.	1.8	28
36	Characterization of a Novel Composite Staphylococcal Cassette Chromosomemec(SCCmec-SCCcad/ars/cop) in the Neonatal Sepsis-Associated Staphylococcus capitis Pulsotype NRCS-A. Antimicrobial Agents and Chemotherapy, 2013, 57, 6354-6357.	3.2	26

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37	ST2249-MRSA-III: a second major recombinant methicillin-resistant Staphylococcus aureus clone causing healthcare infection in the 1970s. Clinical Microbiology and Infection, 2015, 21, 444-450.	6.0	22
38	Outbreak of Skin Infections in College Football Team Members Due to an Unusual Strain of Community-Acquired Methicillin-Susceptible Staphylococcus aureus. Journal of Clinical Microbiology, 2010, 48, 609-611.	3.9	20
39	Identification and Characterization of Linezolid-Resistantcfr-Positive Staphylococcus aureus USA300 Isolates from a New York City Medical Center. Antimicrobial Agents and Chemotherapy, 2014, 58, 6949-6952.	3.2	20
40	Comparative genomic analysis of European and Middle Eastern community-associated methicillin-resistant Staphylococcus aureus (CC80:ST80-IV) isolates by high-density microarray. Clinical Microbiology and Infection, 2009, 15, 748-755.	6.0	18
41	Two Novel Class I Integron Arrays Containing IMP-18 Metallo-Î²-Lactamase Gene in Pseudomonas aeruginosa Clinical Isolates from Puerto Rico. Antimicrobial Agents and Chemotherapy, 2012, 56, 2119-2121.	3.2	18
42	Direct Repeat Unit (dru) Typing of Methicillin-Resistant Staphylococcus pseudintermedius from Dogs and Cats. Journal of Clinical Microbiology, 2015, 53, 3760-3765.	3.9	18
43	Associations between dru Types and SCCmec Cassettes. PLoS ONE, 2013, 8, e61860.	2.5	17
44	Temporal changes in the genotypes of methicillin-resistant Staphylococcus aureus strains isolated from a tertiary Malaysian hospital based on MLST, spa, and mec-associated dru typing. Diagnostic Microbiology and Infectious Disease, 2012, 74, 106-112.	1.8	16
45	From theory to practice: molecular strain typing for the clinical and public health setting. Eurosurveillance, 2013, 18, 20383.	7.0	16
46	Isolation and Characterization of an Epidemic Methicillin-Resistant Staphylococcus aureus 15 Variant in the Central United States. Journal of Clinical Microbiology, 2008, 46, 3548-3549.	3.9	15
47	Toxic Shock Syndrome: Characterization of Human Immune Responses to TSST-1 and Evidence for Sensitivity Thresholds. Toxicological Sciences, 2013, 134, 49-63.	3.1	12
48	Wound infections caused by inducible meticillin-resistant Staphylococcus aureus strains. Journal of Global Antimicrobial Resistance, 2013, 1, 79-83.	2.2	11
49	Complex Clonal Diversity ofStaphylococcus aureusNasal Colonization among Community Personnel, Healthcare Workers, and Clinical Students in the Eastern Province, Saudi Arabia. BioMed Research International, 2018, 2018, 1-9.	1.9	11
50	Daptomycin non-susceptible Staphylococcus aureus at a US medical centre. Clinical Microbiology and Infection, 2013, 19, 1169-1172.	6.0	9
51	A critical review of the in-vitro activity of teicoplanin. International Journal of Antimicrobial Agents, 1995, 5, 169-177.	2.5	7
52	Lineage II (Serovar 1/2a and 1/2c) Human Listeria monocytogenes Pulsed-Field Gel Electrophoresis Types Divided into PFGE Groups Using the Band Patterns Below 145.5â€‰kb. Foodborne Pathogens and Disease, 2017, 14, 8-16.	1.8	5
53	Comparison of the microbiological efficacy of tedizolid and linezolid in acute bacterial skin and skin structure infections: pooled data from phase 3 clinical trials. Diagnostic Microbiology and Infectious Disease, 2019, 94, 277-286.	1.8	5
54	Mobile genetic elements responsible for discordant Staphylococcus aureus phenotypes and genotypes in the same blood culture bottle. Diagnostic Microbiology and Infectious Disease, 2020, 98, 115175.	1.8	5

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55	Presence of Clostridioides difficile and multidrug-resistant healthcare-associated pathogens in stool specimens from hospitalized patients in the USA. Journal of Hospital Infection, 2020, 106, 179-185.	2.9	4
56	Characterization of Clostridioides difficile isolates recovered from two Phase 3 surotomycin treatment trials by restriction endonuclease analysis, PCR ribotyping and antimicrobial susceptibilities. Journal of Antimicrobial Chemotherapy, 2020, 75, 3120-3125.	3.0	4
57	Editorial: New Insights and Updates on the Molecular Epidemiology and Antimicrobial Resistance of MRSA in Humans in the Whole-Genome Sequencing Era. Frontiers in Microbiology, 2019, 10, 637.	3.5	3
58	Characterization of methicillin-resistant Staphylococcus aureus isolated at Tripoli Medical Center, Libya, between 2008 and 2014. Journal of Medical Microbiology, 2016, 65, 1472-1475.	1.8	3
59	Division of HumanListeria monocytogenesPulsed-Field Gel Electrophoresis (PFGE) Types Belonging to Lineage I (Serovar 4b, 1/2b, and 3b) into PFGE Groups. Foodborne Pathogens and Disease, 2015, 12, 447-453.	1.8	2
60	Retrospective Definition of Clostridioides difficile PCR Ribotypes on the Basis of Whole Genome Polymorphisms: A Proof of Principle Study. Diagnostics, 2020, 10, 1078.	2.6	2
61	Characterization of SCC mec Instability in Methicillin-Resistant Staphylococcus aureus Affecting Adjacent Chromosomal Regions, Including the Gene for Staphylococcal Protein A () Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf	3.2	2
62	Whole genome mapping of the first reported case of KPC-2â€“positive Klebsiella pneumoniae ST258 in Nebraska. Diagnostic Microbiology and Infectious Disease, 2014, 79, 384-386.	1.8	1
63	2399. Ribotype Diversity of Clostridioides difficile strains obtained during screening tests. Open Forum Infectious Diseases, 2019, 6, S828-S829.	0.9	0
64	Molecular Epidemiology of Pneumococcal Infections in Adults and Children From the Kansas City Area Between 2010 and 2013. Open Forum Infectious Diseases, 2015, 2, .	0.9	0