

Richard Goering

List of Publications by Year in descending order

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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	Characterization of SCC <i>mecA</i> Instability in Methicillin-Resistant <i>Staphylococcus aureus</i> Affecting Adjacent Chromosomal Regions, Including the Gene for Staphylococcal Protein A (<i>spa</i>). <i>Journal of Hospital Infection</i> , 2020, 106, 179-185.	1.0	10
2	Presence of <i>Clostridioides difficile</i> and multidrug-resistant healthcare-associated pathogens in stool specimens from hospitalized patients in the USA. <i>Journal of Hospital Infection</i> , 2020, 106, 179-185.	2.9	4
3	Mobile genetic elements responsible for discordant <i>Staphylococcus aureus</i> phenotypes and genotypes in the same blood culture bottle. <i>Diagnostic Microbiology and Infectious Disease</i> , 2020, 98, 115-117.	1.8	5
4	Characterization of <i>Clostridioides difficile</i> isolates recovered from two Phase 3 surotomycin treatment trials by restriction endonuclease analysis, PCR ribotyping and antimicrobial susceptibilities. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 3120-3125.	3.0	4
5	Retrospective Definition of <i>Clostridioides difficile</i> PCR Ribotypes on the Basis of Whole Genome Polymorphisms: A Proof of Principle Study. <i>Diagnostics</i> , 2020, 10, 1078.	2.6	2
6	Comparison of the microbiological efficacy of tedizolid and linezolid in acute bacterial skin and skin structure infections: pooled data from phase 3 clinical trials. <i>Diagnostic Microbiology and Infectious Disease</i> , 2019, 94, 277-286.	1.8	5
7	Emergence of Oxacillin Resistance in Stealth Methicillin-Resistant <i>Staphylococcus aureus</i> Due to <i>mecA</i> Sequence Instability. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	34
8	Changes in molecular epidemiology and antimicrobial resistance profiles of <i>Clostridioides (Clostridium) difficile</i> strains in the United States between 2011 and 2017. <i>Anaerobe</i> , 2019, 60, 102050.	2.1	35
9	Editorial: New Insights and Updates on the Molecular Epidemiology and Antimicrobial Resistance of MRSA in Humans in the Whole-Genome Sequencing Era. <i>Frontiers in Microbiology</i> , 2019, 10, 637.	3.5	3
10	2399. Ribotype Diversity of <i>Clostridioides difficile</i> strains obtained during screening tests. <i>Open Forum Infectious Diseases</i> , 2019, 6, S828-S829.	0.9	0
11	Evolution and Global Transmission of a Multidrug-Resistant, Community-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> Lineage from the Indian Subcontinent. <i>MBio</i> , 2019, 10, .	4.1	50
12	Whole genome sequencing options for bacterial strain typing and epidemiologic analysis based on single nucleotide polymorphism versus gene-by-gene based approaches. <i>Clinical Microbiology and Infection</i> , 2018, 24, 350-354.	6.0	373
13	Complex Clonal Diversity of <i>Staphylococcus aureus</i> Nasal Colonization among Community Personnel, Healthcare Workers, and Clinical Students in the Eastern Province, Saudi Arabia. <i>BioMed Research International</i> , 2018, 2018, 1-9.	1.9	11
14	Continued expansion of USA300-like methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) among hospitalized patients in the United States. <i>Diagnostic Microbiology and Infectious Disease</i> , 2017, 88, 342-347.	1.8	28
15	Lineage II (Serovar 1/2a and 1/2c) Human <i>Listeria monocytogenes</i> Pulsed-Field Gel Electrophoresis Types Divided into PFGE Groups Using the Band Patterns Below 145.5 kb. <i>Foodborne Pathogens and Disease</i> , 2017, 14, 8-16.	1.8	5
16	Wide geographical dissemination of the multiresistant <i>Staphylococcus capitis</i> NRCS-A clone in neonatal intensive-care units. <i>Clinical Microbiology and Infection</i> , 2016, 22, 46-52.	6.0	58
17	Characterization of methicillin-resistant <i>Staphylococcus aureus</i> isolated at Tripoli Medical Center, Libya, between 2008 and 2014. <i>Journal of Medical Microbiology</i> , 2016, 65, 1472-1475.	1.8	3
18	Division of Human <i>Listeria monocytogenes</i> Pulsed-Field Gel Electrophoresis (PFGE) Types Belonging to Lineage I (Serovar 4b, 1/2b, and 3b) into PFGE Groups. <i>Foodborne Pathogens and Disease</i> , 2015, 12, 447-453.	1.8	2

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19	ST2249-MRSA-III: a second major recombinant methicillin-resistant <i>Staphylococcus aureus</i> clone causing healthcare infection in the 1970s. <i>Clinical Microbiology and Infection</i> , 2015, 21, 444-450.	6.0	22
20	Direct Repeat Unit (<i>dru</i>) Typing of Methicillin-Resistant <i>Staphylococcus pseudintermedius</i> from Dogs and Cats. <i>Journal of Clinical Microbiology</i> , 2015, 53, 3760-3765.	3.9	18
21	Microbial Typing by Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry: Do We Need Guidance for Data Interpretation?. <i>Journal of Clinical Microbiology</i> , 2015, 53, 760-765.	3.9	92
22	Molecular Epidemiology of Pneumococcal Infections in Adults and Children From the Kansas City Area Between 2010 and 2013. <i>Open Forum Infectious Diseases</i> , 2015, 2, .	0.9	0
23	Whole genome mapping of the first reported case of KPC-2–positive <i>Klebsiella pneumoniae</i> ST258 in Nebraska. <i>Diagnostic Microbiology and Infectious Disease</i> , 2014, 79, 384-386.	1.8	1
24	Strain Types and Antimicrobial Resistance Patterns of <i>Clostridium difficile</i> Isolates from the United States, 2011 to 2013. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 4214-4218.	3.2	103
25	Bacterial Whole-Genome Sequencing Revisited: Portable, Scalable, and Standardized Analysis for Typing and Detection of Virulence and Antibiotic Resistance Genes. <i>Journal of Clinical Microbiology</i> , 2014, 52, 2365-2370.	3.9	250
26	Linezolid-Resistant <i>Staphylococcus aureus</i> Strain 1128105, the First Known Clinical Isolate Possessing the <i>cr</i> Multidrug Resistance Gene. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 6592-6598.	3.2	34
27	Identification and Characterization of Linezolid-Resistant <i>cr</i> -Positive <i>Staphylococcus aureus</i> USA300 Isolates from a New York City Medical Center. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 6949-6952.	3.2	20
28	Molecular characterization of methicillin-resistant <i>Staphylococcus aureus</i> isolated over a 2-year period in a Qatari hospital from multinational patients. <i>Clinical Microbiology and Infection</i> , 2014, 20, 169-173.	6.0	40
29	Wound infections caused by inducible methicillin-resistant <i>Staphylococcus aureus</i> strains. <i>Journal of Global Antimicrobial Resistance</i> , 2013, 1, 79-83.	2.2	11
30	Activity of Tedizolid (TR-700) against Well-Characterized Methicillin-Resistant <i>Staphylococcus aureus</i> Strains of Diverse Epidemiological Origins. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 2892-2895.	3.2	44
31	Toxic Shock Syndrome: Characterization of Human Immune Responses to TSST-1 and Evidence for Sensitivity Thresholds. <i>Toxicological Sciences</i> , 2013, 134, 49-63.	3.1	12
32	Multicenter Evaluation of the Verigene <i>Clostridium difficile</i> Nucleic Acid Assay. <i>Journal of Clinical Microbiology</i> , 2013, 51, 4120-4125.	3.9	31
33	Associations between <i>dru</i> Types and SCCmec Cassettes. <i>PLoS ONE</i> , 2013, 8, e61860.	2.5	17
34	Daptomycin non-susceptible <i>Staphylococcus aureus</i> at a US medical centre. <i>Clinical Microbiology and Infection</i> , 2013, 19, 1169-1172.	6.0	9
35	Transduction of Staphylococcal Cassette Chromosome <i>mec</i> Elements between Strains of <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 5233-5238.	3.2	64
36	Characterization of a Novel Composite Staphylococcal Cassette Chromosome <i>mec</i> (SCCmec-SCCcad/ars/cop) in the Neonatal Sepsis-Associated <i>Staphylococcus capitis</i> Pulsotype NRCS-A. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 6354-6357.	3.2	26

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37	Subpopulations of <i>Staphylococcus aureus</i> Clonal Complex 121 Are Associated with Distinct Clinical Entities. <i>PLoS ONE</i> , 2013, 8, e58155.	2.5	43
38	From theory to practice: molecular strain typing for the clinical and public health setting. <i>Eurosurveillance</i> , 2013, 18, 20383.	7.0	16
39	Characterization of Nasal and Blood Culture Isolates of Methicillin-Resistant <i>Staphylococcus aureus</i> from Patients in United States Hospitals. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 1324-1330.	3.2	105
40	Two Novel Class I Integron Arrays Containing IMP-18 Metallo- β -Lactamase Gene in <i>Pseudomonas aeruginosa</i> Clinical Isolates from Puerto Rico. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 2119-2121.	3.2	18
41	Guidelines for Reporting Novel <i>mecA</i> Gene Homologues. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 4997-4999.	3.2	144
42	<i>Stenotrophomonas maltophilia</i> in Malaysia: molecular epidemiology and trimethoprim-sulfamethoxazole resistance. <i>International Journal of Infectious Diseases</i> , 2012, 16, e603-e607.	3.3	30
43	Temporal changes in the genotypes of methicillin-resistant <i>Staphylococcus aureus</i> strains isolated from a tertiary Malaysian hospital based on MLST, <i>spa</i> , and <i>mec</i> -associated <i>dru</i> typing. <i>Diagnostic Microbiology and Infectious Disease</i> , 2012, 74, 106-112.	1.8	16
44	Characterization of a Novel Arginine Catabolic Mobile Element (ACME) and Staphylococcal Chromosomal Cassette <i>mec</i> Composite Island with Significant Homology to <i>Staphylococcus epidermidis</i> ACME Type II in Methicillin-Resistant <i>Staphylococcus aureus</i> Genotype ST22-MRSA-IV. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 1896-1905.	3.2	83
45	Comparison of Strain Typing Results for <i>Clostridium difficile</i> Isolates from North America. <i>Journal of Clinical Microbiology</i> , 2011, 49, 1831-1837.	3.9	86
46	Pulsed field gel electrophoresis: A review of application and interpretation in the molecular epidemiology of infectious disease. <i>Infection, Genetics and Evolution</i> , 2010, 10, 866-875.	2.3	200
47	Enhanced Discrimination of Highly Clonal ST22-Methicillin-Resistant <i>Staphylococcus aureus</i> IV Isolates Achieved by Combining <i>spa</i> , <i>dru</i> , and Pulsed-Field Gel Electrophoresis Typing Data. <i>Journal of Clinical Microbiology</i> , 2010, 48, 1839-1852.	3.9	55
48	Outbreak of Skin Infections in College Football Team Members Due to an Unusual Strain of Community-Acquired Methicillin-Susceptible <i>Staphylococcus aureus</i> . <i>Journal of Clinical Microbiology</i> , 2010, 48, 609-611.	3.9	20
49	Impact of Strain Type on Detection of Toxigenic <i>Clostridium difficile</i> : Comparison of Molecular Diagnostic and Enzyme Immunoassay Approaches. <i>Journal of Clinical Microbiology</i> , 2010, 48, 3719-3724.	3.9	177
50	Outbreak of Carbapenem-Resistant <i>Klebsiella pneumoniae</i> in Puerto Rico Associated with a Novel Carbapenemase Variant. <i>Infection Control and Hospital Epidemiology</i> , 2010, 31, 476-484.	1.8	89
51	Surveillance of Carbapenem-Resistant <i>Pseudomonas aeruginosa</i> Isolates from Puerto Rican Medical Center Hospitals: Dissemination of KPC and IMP-18 β -Lactamases. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 1660-1664.	3.2	88
52	Two Distinct Clones of Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) with the Same USA300 Pulsed-Field Gel Electrophoresis Profile: a Potential Pitfall for Identification of USA300 Community-Associated MRSA. <i>Journal of Clinical Microbiology</i> , 2009, 47, 3765-3768.	3.9	46
53	Phenotypic and Enzymatic Comparative Analysis of the Novel KPC Variant KPC-5 and Its Evolutionary Variants, KPC-2 and KPC-4. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 557-562.	3.2	119
54	Comparative genomic analysis of European and Middle Eastern community-associated methicillin-resistant <i>Staphylococcus aureus</i> (CC80:ST80-IV) isolates by high-density microarray. <i>Clinical Microbiology and Infection</i> , 2009, 15, 748-755.	6.0	18

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55	Methicillin-resistant <i>Staphylococcus aureus</i> strain USA300: origin and epidemiology. <i>Journal of Antimicrobial Chemotherapy</i> , 2009, 64, 441-446.	3.0	380
56	Usefulness of mec-associated direct repeat unit (dru) typing in the epidemiological analysis of highly clonal methicillin-resistant <i>Staphylococcus aureus</i> in Scotland. <i>Clinical Microbiology and Infection</i> , 2008, 14, 964-969.	6.0	92
57	Frequent emergence and limited geographic dispersal of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 14130-14135.	7.1	239
58	Emergence of Carbapenem Resistance in <i>Pseudomonas aeruginosa</i> Isolates from a Patient with Cystic Fibrosis in the Absence of Carbapenem Therapy. <i>Clinical Infectious Diseases</i> , 2008, 46, e137-e141.	5.8	31
59	Molecular Epidemiology of Methicillin-Resistant and Methicillin-Susceptible <i>Staphylococcus aureus</i> Isolates from Global Clinical Trials. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2842-2847.	3.9	113
60	Prevalence of Toxic Shock Syndrome Toxin 1 (TSST-1)-Producing Strains of <i>Staphylococcus aureus</i> and Antibody to TSST-1 among Healthy Japanese Women. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2731-2738.	3.9	43
61	Isolation and Characterization of an Epidemic Methicillin-Resistant <i>Staphylococcus aureus</i> 15 Variant in the Central United States. <i>Journal of Clinical Microbiology</i> , 2008, 46, 3548-3549.	3.9	15
62	Molecular Typing of Methicillin-Resistant <i>Staphylococcus aureus</i> by Pulsed-Field Gel Electrophoresis: Comparison of Results Obtained in a Multilaboratory Effort Using Identical Protocols and MRSA Strains. <i>Microbial Drug Resistance</i> , 2000, 6, 189-198.	2.0	267
63	Assessment of Resolution and Intercenter Reproducibility of Results of Genotyping <i>Staphylococcus aureus</i> by Pulsed-Field Gel Electrophoresis of <i>Sma</i> I Macrorestriction Fragments: a Multicenter Study. <i>Journal of Clinical Microbiology</i> , 1998, 36, 1653-1659.	3.9	186
64	A critical review of the in-vitro activity of teicoplanin. <i>International Journal of Antimicrobial Agents</i> , 1995, 5, 169-177.	2.5	7