

Anna C Shore

List of Publications by Year in descending order

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41
papers

2,931
citations

218677

26
h-index

276875

41
g-index

42
all docs

42
docs citations

42
times ranked

3034
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic analysis of 600 vancomycin-resistant <i>Enterococcus faecium</i> reveals a high prevalence of ST80 and spread of similar <i>vanA</i> regions via IS1216E and plasmid transfer in diverse genetic lineages in Ireland. <i>Journal of Antimicrobial Chemotherapy</i> , 2022, 77, 320-330.	3.0	13
2	Screening the nose, throat and the naso-pharynx for methicillin-resistant <i>Staphylococcus aureus</i> : a pilot study. <i>Journal of Infection Prevention</i> , 2020, 21, 155-158.	0.9	2
3	Linezolid resistance in <i>Enterococcus faecium</i> and <i>Enterococcus faecalis</i> from hospitalized patients in Ireland: high prevalence of the MDR genes <i>optrA</i> and <i>poxA</i> in isolates with diverse genetic backgrounds. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 1704-1711.	3.0	48
4	A novel multidrug-resistant PVL-negative CC1-MRSA-IV clone emerging in Ireland and Germany likely originated in South-Eastern Europe. <i>Infection, Genetics and Evolution</i> , 2019, 69, 117-126.	2.3	20
5	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
6	A molecular epidemiological investigation of methicillin-susceptible <i>Staphylococcus aureus</i> causing bloodstream infections in Ireland, 2006–2017. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2019, 38, 927-936.	2.9	8
7	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
8	Range Expansion and the Origin of USA300 North American Epidemic Methicillin-Resistant <i>Staphylococcus aureus</i> . <i>MBio</i> , 2018, 9, .	4.1	42
9	Intra-Hospital, Inter-Hospital and Intercontinental Spread of ST78 MRSA From Two Neonatal Intensive Care Unit Outbreaks Established Using Whole-Genome Sequencing. <i>Frontiers in Microbiology</i> , 2018, 9, 1485.	3.5	26
10	Molecular Typing of ST239-MRSA-III From Diverse Geographic Locations and the Evolution of the SCCmec III Element During Its Intercontinental Spread. <i>Frontiers in Microbiology</i> , 2018, 9, 1436.	3.5	45
11	Observational cross-sectional study of nasal staphylococcal species of medical students of diverse geographical origin, prior to healthcare exposure: prevalence of SCC <i>mec</i> , <i>fusC</i> , <i>fusB</i> and the arginine catabolite mobile element (ACME) in the absence of selective antibiotic pressure. <i>BMI Open</i> , 2018, 8, e020391.	1.9	13
12	Molecular Characterization of Nasal Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates Showing Increasing Prevalence of Mupirocin Resistance and Associated Multidrug Resistance following Attempted Decolonization. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	6
13	Novel multiresistance <i>cfr</i> plasmids in linezolid-resistant methicillin-resistant <i>Staphylococcus epidermidis</i> and vancomycin-resistant <i>Enterococcus faecium</i> (VRE) from a hospital outbreak: co-location of <i>cfr</i> and <i>optrA</i> in VRE. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 3252-3257.	3.0	80
14	The recent emergence in hospitals of multidrug-resistant community-associated sequence type 1 and spa type t127 methicillin-resistant <i>Staphylococcus aureus</i> investigated by whole-genome sequencing: Implications for screening. <i>PLoS ONE</i> , 2017, 12, e0175542.	2.5	45
15	Dissemination of high-level mupirocin-resistant CC22-MRSA-IV in Saxony. <i>GMS Hygiene and Infection Control</i> , 2017, 12, Doc19.	0.3	2
16	Diversity of <i>Staphylococcus aureus</i> Isolates in European Wildlife. <i>PLoS ONE</i> , 2016, 11, e0168433.	2.5	94
17	Enhanced Tracking of Nosocomial Transmission of Endemic Sequence Type 22 Methicillin-Resistant <i>Staphylococcus aureus</i> Type IV Isolates among Patients and Environmental Sites by Use of Whole-Genome Sequencing. <i>Journal of Clinical Microbiology</i> , 2016, 54, 445-448.	3.9	19
18	First Report of <i>cfr</i> -Carrying Plasmids in the Pandemic Sequence Type 22 Methicillin-Resistant <i>Staphylococcus aureus</i> Staphylococcal Cassette Chromosome <i>mec</i> Type IV Clone. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 3007-3015.	3.2	37

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19	In vitro activity of ceftaroline against mecC-positive MRSA isolates. <i>Journal of Global Antimicrobial Resistance</i> , 2016, 5, 3-6.	2.2	1
20	The Emergence and Spread of Multiple Livestock-Associated Clonal Complex 398 Methicillin-Resistant and Methicillin-Susceptible <i>Staphylococcus aureus</i> Strains among Animals and Humans in the Republic of Ireland, 2010–2014. <i>PLoS ONE</i> , 2016, 11, e0149396.	2.5	21
21	Comparative Genotypes, Staphylococcal Cassette Chromosome mec (SCCmec) Genes and Antimicrobial Resistance amongst <i>Staphylococcus epidermidis</i> and <i>Staphylococcus haemolyticus</i> Isolates from Infections in Humans and Companion Animals. <i>PLoS ONE</i> , 2015, 10, e0138079.	2.5	66
22	Extensive Genetic Diversity Identified among Sporadic Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates Recovered in Irish Hospitals between 2000 and 2012. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 1907-1917.	3.2	37
23	Panton-Valentine Leukocidin-Positive <i>Staphylococcus aureus</i> in Ireland from 2002 to 2011: 21 Clones, Frequent Importation of Clones, Temporal Shifts of Predominant Methicillin-Resistant <i>S. aureus</i> Clones, and Increasing Multiresistance. <i>Journal of Clinical Microbiology</i> , 2014, 52, 859-870.	3.9	68
24	Staphylococcal cassette chromosome mec: Recent advances and new insights. <i>International Journal of Medical Microbiology</i> , 2013, 303, 350-359.	3.6	135
25	Emergence of Sequence Type 779 Methicillin-Resistant <i>Staphylococcus aureus</i> Harboring a Novel Pseudo Staphylococcal Cassette Chromosome <i>mec</i> (SCC <i>mec</i>)-SCC-SCC _{CRISPR} Composite Element in Irish Hospitals. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 524-531.	3.2	72
26	Detection of mecC-Positive <i>Staphylococcus aureus</i> (CC130-MRSA-XI) in Diseased European Hedgehogs (<i>Erinaceus europaeus</i>) in Sweden. <i>PLoS ONE</i> , 2013, 8, e66166.	2.5	74
27	Emergence of Hospital- and Community-Associated Panton-Valentine Leukocidin-Positive Methicillin-Resistant <i>Staphylococcus aureus</i> Genotype ST772-MRSA-V in Ireland and Detailed Investigation of an ST772-MRSA-V Cluster in a Neonatal Intensive Care Unit. <i>Journal of Clinical Microbiology</i> , 2012, 50, 841-847.	3.9	67
28	DNA Microarray Profiling of a Diverse Collection of Nosocomial Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates Assigns the Majority to the Correct Sequence Type and Staphylococcal Cassette Chromosome <i>mec</i> (SCC <i>mec</i>) Type and Results in the Subsequent Identification and Characterization of Novel SCC <i>mec</i> -SCC _{M1} Composite Islands. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 5340-5355.	3.2	29
29	Evaluation of screening risk and nonrisk patients for methicillin-resistant <i>Staphylococcus aureus</i> on admission in an acute care hospital. <i>American Journal of Infection Control</i> , 2012, 40, 411-415.	2.3	19
30	Distribution of SCCmec-associated phenol-soluble modulins in staphylococci. <i>Molecular and Cellular Probes</i> , 2012, 26, 99-103.	2.1	23
31	A Field Guide to Pandemic, Epidemic and Sporadic Clones of Methicillin-Resistant <i>Staphylococcus aureus</i> . <i>PLoS ONE</i> , 2011, 6, e17936.	2.5	734
32	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
33	DNA Microarray Genotyping and Virulence and Antimicrobial Resistance Gene Profiling of Methicillin-Resistant <i>Staphylococcus aureus</i> Bloodstream Isolates from Renal Patients. <i>Journal of Clinical Microbiology</i> , 2011, 49, 4349-4351.	3.9	13
34	Detection of Staphylococcal Cassette Chromosome <i>mec</i> Type XI Carrying Highly Divergent <i>mecA</i> , <i>mecI</i> , <i>mecR1</i> , <i>blaZ</i> , and <i>ccr</i> Genes in Human Clinical Isolates of Clonal Complex 130 Methicillin-Resistant <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 3765-3773.	3.2	336
35	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
36	Identification and Characterization of the Multidrug Resistance Gene <i>cfr</i> in a Panton-Valentine Leukocidin-Positive Sequence Type 8 Methicillin-Resistant <i>Staphylococcus aureus</i> IVa (USA300) Isolate. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 4978-4984.	3.2	91

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37	The Effect of Rapid Screening for Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) on the Identification and Earlier Isolation of MRSA-Positive Patients. <i>Infection Control and Hospital Epidemiology</i> , 2010, 31, 374-381.	1.8	34
38	Molecular typing of nasal carriage isolates of <i>Staphylococcus aureus</i> from an Irish university student population based on toxin gene PCR, <i>agr</i> locus types and multiple locus, variable number tandem repeat analysis. <i>Journal of Medical Microbiology</i> , 2008, 57, 348-358.	1.8	43
39	Detection of Staphylococcal Cassette Chromosome <i>mec</i> -Associated DNA Segments in Multiresistant Methicillin-Susceptible <i>Staphylococcus aureus</i> (MSSA) and Identification of <i>Staphylococcus epidermidis ccrAB4</i> in both Methicillin-Resistant <i>S. aureus</i> and MSSA. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 4407-4419.	3.2	65
40	The Emergence and Importation of Diverse Genotypes of Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) Harboring the Panton-Valentine Leukocidin Gene (<i>pvl</i>) Reveal that <i>pvl</i> Is a Poor Marker for Community-Acquired MRSA Strains in Ireland. <i>Journal of Clinical Microbiology</i> , 2007, 45, 2554-2563.	3.9	154
41	Seven Novel Variants of the Staphylococcal Chromosomal Cassette <i>mec</i> in Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates from Ireland. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 2070-2083.	3.2	157