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	A Field Guide to Pandemic, Epidemic and Sporadic Clones of Methicillin-Resistant Staphylococcus aureus. PLoS ONE, 2011, 6, e17936.	2.5	734
2	Detection of Staphylococcal Cassette Chromosome mec Type XI Carrying Highly Divergent mecA , mecI , mecR1 , blaZ , and ccr Genes in Human Clinical Isolates of Clonal Complex 130 Methicillin-Resistant Staphylococcus aureus. Antimicrobial Agents and Chemotherapy, 2011, 55, 3765-3773.	3.2	336
3	Seven Novel Variants of the Staphylococcal Chromosomal Cassette mec in Methicillin-Resistant Staphylococcus aureus Isolates from Ireland. Antimicrobial Agents and Chemotherapy, 2005, 49, 2070-2083.	3.2	157
4	The Emergence and Importation of Diverse Genotypes of Methicillin-Resistant Staphylococcus aureus (MRSA) Harboring the Panton-Valentine Leukocidin Gene (pvl) Reveal that pvl Is a Poor Marker for Community-Acquired MRSA Strains in Ireland. Journal of Clinical Microbiology, 2007, 45, 2554-2563.	3.9	154
5	Staphylococcal cassette chromosome mec: Recent advances and new insights. International Journal of Medical Microbiology, 2013, 303, 350-359.	3.6	135
6	Diversity of Staphylococcus aureus Isolates in European Wildlife. PLoS ONE, 2016, 11, e0168433.	2.5	94
7	Identification and Characterization of the Multidrug Resistance Gene cfr in a Panton-Valentine Leukocidin-Positive Sequence Type 8 Methicillin-Resistant Staphylococcus aureus IVa (USA300) Isolate. Antimicrobial Agents and Chemotherapy, 2010, 54, 4978-4984.	3.2	91
8	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
9	Novel multiresistance cfr plasmids in linezolid-resistant methicillin-resistant Staphylococcus epidermidis and vancomycin-resistant Enterococcus faecium (VRE) from a hospital outbreak: co-location of cfr and optrA in VRE. Journal of Antimicrobial Chemotherapy, 2017, 72, 3252-3257.	3.0	80
10	Detection of mecC-Positive Staphylococcus aureus (CC130-MRSA-XI) in Diseased European Hedgehogs (Erinaceus europaeus) in Sweden. PLoS ONE, 2013, 8, e66166.	2.5	74
11	Emergence of Sequence Type 779 Methicillin-Resistant Staphylococcus aureus Harboring a Novel Pseudo Staphylococcal Cassette Chromosome mec (SCC mec)-SCC-SCC _{CRISPR} Composite Element in Irish Hospitals. Antimicrobial Agents and Chemotherapy, 2013, 57, 524-531.	3.2	72
12	Panton-Valentine Leukocidin-Positive Staphylococcus aureus in Ireland from 2002 to 2011: 21 Clones, Frequent Importation of Clones, Temporal Shifts of Predominant Methicillin-Resistant S. aureus Clones, and Increasing Multiresistance. Journal of Clinical Microbiology, 2014, 52, 859-870.	3.9	68
13	Emergence of Hospital- and Community-Associated Panton-Valentine Leukocidin-Positive Methicillin-Resistant Staphylococcus aureus Genotype ST772-MRSA-V in Ireland and Detailed Investigation of an ST772-MRSA-V Cluster in a Neonatal Intensive Care Unit. Journal of Clinical Microbiology, 2012, 50, 841-847.	3.9	67
14	Comparative Genotypes, Staphylococcal Cassette Chromosome mec (SCCmec) Genes and Antimicrobial Resistance amongst Staphylococcus epidermidis and Staphylococcus haemolyticus Isolates from Infections in Humans and Companion Animals. PLoS ONE, 2015, 10, e0138079.	2.5	66
15	Detection of Staphylococcal Cassette Chromosome mec -Associated DNA Segments in Multiresistant Methicillin-Susceptible Staphylococcus aureus (MSSA) and Identification of Staphylococcus epidermidis ccrAB4 in both Methicillin-Resistant S. aureus and MSSA. Antimicrobial Agents and Chemotherapy, 2008, 52, 4407-4419.	3.2	65
16	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
17	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
18	Linezolid resistance in Enterococcus faecium and Enterococcus faecalis from hospitalized patients in Ireland: high prevalence of the MDR genes optrA and poxtA in isolates with diverse genetic backgrounds. Journal of Antimicrobial Chemotherapy, 2020, 75, 1704-1711.	3.0	48

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19	The recent emergence in hospitals of multidrug-resistant community-associated sequence type 1 and spa type t127 methicillin-resistant Staphylococcus aureus investigated by whole-genome sequencing: Implications for screening. PLoS ONE, 2017, 12, e0175542.	2.5	45
20	Molecular Typing of ST239-MRSA-III From Diverse Geographic Locations and the Evolution of the SCCmec III Element During Its Intercontinental Spread. Frontiers in Microbiology, 2018, 9, 1436.	3.5	45
21	Molecular typing of nasal carriage isolates of Staphylococcus aureus from an Irish university student population based on toxin gene PCR, agr locus types and multiple locus, variable number tandem repeat analysis. Journal of Medical Microbiology, 2008, 57, 348-358.	1.8	43
22	Range Expansion and the Origin of USA300 North American Epidemic Methicillin-Resistant Staphylococcus aureus. MBio, 2018, 9, .	4.1	42
23	Extensive Genetic Diversity Identified among Sporadic Methicillin-Resistant Staphylococcus aureus Isolates Recovered in Irish Hospitals between 2000 and 2012. Antimicrobial Agents and Chemotherapy, 2014, 58, 1907-1917.	3.2	37
24	First Report of cfr -Carrying Plasmids in the Pandemic Sequence Type 22 Methicillin-Resistant Staphylococcus aureus Staphylococcal Cassette Chromosome mec Type IV Clone. Antimicrobial Agents and Chemotherapy, 2016, 60, 3007-3015.	3.2	37
25	The Effect of Rapid Screening for Methicillin-ResistantStaphylococcus aureus(MRSA) on the Identification and Earlier Isolation of MRSA-Positive Patients. Infection Control and Hospital Epidemiology, 2010, 31, 374-381.	1.8	34
26	DNA Microarray Profiling of a Diverse Collection of Nosocomial Methicillin-Resistant Staphylococcus aureus Isolates Assigns the Majority to the Correct Sequence Type and Staphylococcal Cassette Chromosome mec (SCC mec) Type and Results in the Subsequent Identification and Characterization of Novel SCC mec -SCC _{M1} Composite Islands. Antimicrobial Agents and Chemotherapy, 2012, 56, 5340-5355.	3.2	29
27	Intra-Hospital, Inter-Hospital and Intercontinental Spread of ST78 MRSA From Two Neonatal Intensive Care Unit Outbreaks Established Using Whole-Genome Sequencing. Frontiers in Microbiology, 2018, 9, 1485.	3.5	26
28	Distribution of SCCmec-associated phenol-soluble modulin in staphylococci. Molecular and Cellular Probes, 2012, 26, 99-103.	2.1	23
29	The Emergence and Spread of Multiple Livestock-Associated Clonal Complex 398 Methicillin-Resistant and Methicillin-Susceptible Staphylococcus aureus Strains among Animals and Humans in the Republic of Ireland, 2010â€“2014. PLoS ONE, 2016, 11, e0149396.	2.5	21
30	A novel multidrug-resistant PVL-negative CC1-MRSA-IV clone emerging in Ireland and Germany likely originated in South-Eastern Europe. Infection, Genetics and Evolution, 2019, 69, 117-126.	2.3	20
31	Evaluation of screening risk and nonrisk patients for methicillin-resistant Staphylococcus aureus on admission in an acute care hospital. American Journal of Infection Control, 2012, 40, 411-415.	2.3	19
32	Enhanced Tracking of Nosocomial Transmission of Endemic Sequence Type 22 Methicillin-Resistant Staphylococcus aureus Type IV Isolates among Patients and Environmental Sites by Use of Whole-Genome Sequencing. Journal of Clinical Microbiology, 2016, 54, 445-448.	3.9	19
33	DNA Microarray Genotyping and Virulence and Antimicrobial Resistance Gene Profiling of Methicillin-Resistant Staphylococcus aureus Bloodstream Isolates from Renal Patients. Journal of Clinical Microbiology, 2011, 49, 4349-4351.	3.9	13
34	Observational cross-sectional study of nasal staphylococcal species of medical students of diverse geographical origin, prior to healthcare exposure: prevalence of SCCmec,fusC,fusBand the arginine catabolite mobile element (ACME) in the absence of selective antibiotic pressure. BMJ Open, 2018, 8, e020391.	1.9	13
35	Genomic analysis of 600 vancomycin-resistant Enterococcus faecium reveals a high prevalence of ST80 and spread of similar vanA regions via IS1216E and plasmid transfer in diverse genetic lineages in Ireland. Journal of Antimicrobial Chemotherapy, 2022, 77, 320-330.	3.0	13
36	A molecular epidemiological investigation of methicillin-susceptible Staphylococcus aureus causing bloodstream infections in Ireland, 2006â€“2017. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 927-936.	2.9	8

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37	Molecular Characterization of Nasal Methicillin-Resistant Staphylococcus aureus Isolates Showing Increasing Prevalence of Mupirocin Resistance and Associated Multidrug Resistance following Attempted Decolonization. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	6
38	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
39	Screening the nose, throat and the naso-pharynx for methicillin-resistant Staphylococcus aureus: a pilot study. Journal of Infection Prevention, 2020, 21, 155-158.	0.9	2
40	Dissemination of high-level mupirocin-resistant CC22-MRSA-IV in Saxony. GMS Hygiene and Infection Control, 2017, 12, Doc19.	0.3	2
41	In vitro activity of ceftaroline against mecC-positive MRSA isolates. Journal of Global Antimicrobial Resistance, 2016, 5, 3-6.	2.2	1