

Geoffrey W Coombs

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175
papers

6,289
citations

37
h-index

75
g-index

186
ext. papers

7,506
ext. citations

5.8
avg, IF

5.28
L-index

#	Paper	IF	Citations
175	Dissemination of new methicillin-resistant <i>Staphylococcus aureus</i> clones in the community. <i>Journal of Clinical Microbiology</i> , 2002 , 40, 4289-94	9.7	733
174	A field guide to pandemic, epidemic and sporadic clones of methicillin-resistant <i>Staphylococcus aureus</i> . <i>PLoS ONE</i> , 2011 , 6, e17936	3.7	563
173	A genomic portrait of the emergence, evolution, and global spread of a methicillin-resistant <i>Staphylococcus aureus</i> pandemic. <i>Genome Research</i> , 2013 , 23, 653-64	9.7	325
172	Evolution of multidrug resistance during <i>Staphylococcus aureus</i> infection involves mutation of the essential two component regulator WalKR. <i>PLoS Pathogens</i> , 2011 , 7, e1002359	7.6	213
171	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, 14130-5	11.5	205
170	Antibiotic choice may not explain poorer outcomes in patients with <i>Staphylococcus aureus</i> bacteremia and high vancomycin minimum inhibitory concentrations. <i>Journal of Infectious Diseases</i> , 2011 , 204, 340-7	7	189
169	Antimicrobial resistance: Not community-associated methicillin-resistant <i>Staphylococcus aureus</i> (CA-MRSA)! A clinician's guide to community MRSA - its evolving antimicrobial resistance and implications for therapy. <i>Clinical Infectious Diseases</i> , 2011 , 52, 99-114	11.6	146
168	<i>Staphylococcus aureus</i> bacteraemia: a major cause of mortality in Australia and New Zealand. <i>Medical Journal of Australia</i> , 2009 , 191, 368-73	4	141
167	Genetic diversity among community methicillin-resistant <i>Staphylococcus aureus</i> strains causing outpatient infections in Australia. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 4735-43	9.7	139
166	Diversity among community isolates of methicillin-resistant <i>Staphylococcus aureus</i> in Australia. <i>Journal of Clinical Microbiology</i> , 2004 , 42, 3185-90	9.7	120
165	Combination of Vancomycin and β -Lactam Therapy for Methicillin-Resistant <i>Staphylococcus aureus</i> Bacteremia: A Pilot Multicenter Randomized Controlled Trial. <i>Clinical Infectious Diseases</i> , 2016 , 62, 173-180	11.6	117
164	Genomic insights to control the emergence of vancomycin-resistant enterococci. <i>MBio</i> , 2013 , 4,	7.8	112
163	Community-associated methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) in Australia. <i>International Journal of Antimicrobial Agents</i> , 2008 , 31, 401-10	14.3	110
162	Increasing tolerance of hospital to handwash alcohols. <i>Science Translational Medicine</i> , 2018 , 10,	17.5	102
161	Methicillin-resistant <i>Staphylococcus aureus</i> in the Australian community: an evolving epidemic. <i>Medical Journal of Australia</i> , 2006 , 184, 384-8	4	95
160	Comparative genomics and DNA array-based genotyping of pandemic <i>Staphylococcus aureus</i> strains encoding Panton-Valentine leukocidin. <i>Clinical Microbiology and Infection</i> , 2007 , 13, 236-49	9.5	84
159	Epidemiology and antimicrobial susceptibility profiles of pathogens causing urinary tract infections in the Asia-Pacific region: Results from the Study for Monitoring Antimicrobial Resistance Trends (SMART), 2010-2013. <i>International Journal of Antimicrobial Agents</i> , 2016 , 47, 328-34	14.3	81

158	Origin, evolution, and global transmission of community-acquired ST8. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E10596-E10604	11.5	80
157	Genomic insights into the emergence and spread of international clones of healthcare-, community- and livestock-associated methicillin-resistant <i>Staphylococcus aureus</i> : Blurring of the traditional definitions. <i>Journal of Global Antimicrobial Resistance</i> , 2016 , 6, 95-101	3.4	80
156	Methicillin-resistant <i>Staphylococcus aureus</i> clones, Western Australia. <i>Emerging Infectious Diseases</i> , 2006 , 12, 241-7	10.2	71
155	Emergence of community-acquired methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) infection in Queensland, Australia. <i>International Journal of Infectious Diseases</i> , 2003 , 7, 259-64	10.5	67
154	Community-acquired pneumonia due to pandemic A(H1N1)2009 influenza virus and methicillin resistant <i>Staphylococcus aureus</i> co-infection. <i>PLoS ONE</i> , 2010 , 5, e8705	3.7	63
153	Differentiation of clonal complex 59 community-associated methicillin-resistant <i>Staphylococcus aureus</i> in Western Australia. <i>Antimicrobial Agents and Chemotherapy</i> , 2010 , 54, 1914-21	5.9	60
152	Evolution and diversity of community-associated methicillin-resistant <i>Staphylococcus aureus</i> in a geographical region. <i>BMC Microbiology</i> , 2011 , 11, 215	4.5	59
151	Eradication of a large outbreak of a single strain of vanB vancomycin-resistant <i>Enterococcus faecium</i> at a major Australian teaching hospital. <i>Infection Control and Hospital Epidemiology</i> , 2004 , 25, 384-90	2	59
150	An updated view of plasmid conjugation and mobilization in <i>Staphylococcus</i> . <i>Mobile Genetic Elements</i> , 2016 , 6, e1208317		58
149	The dominant Australian community-acquired methicillin-resistant <i>Staphylococcus aureus</i> clone ST93-IV [2B] is highly virulent and genetically distinct. <i>PLoS ONE</i> , 2011 , 6, e25887	3.7	56
148	Antimicrobial-resistant CC17 <i>Enterococcus faecium</i> : The past, the present and the future. <i>Journal of Global Antimicrobial Resistance</i> , 2019 , 16, 36-47	3.4	55
147	Molecular epidemiology of enterococcal bacteremia in Australia. <i>Journal of Clinical Microbiology</i> , 2014 , 52, 897-905	9.7	53
146	Origin-of-transfer sequences facilitate mobilisation of non-conjugative antimicrobial-resistance plasmids in <i>Staphylococcus aureus</i> . <i>Nucleic Acids Research</i> , 2015 , 43, 7971-83	20.1	52
145	Community-associated versus healthcare-associated methicillin-resistant <i>Staphylococcus aureus</i> bacteraemia: a 10-year retrospective review. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2009 , 28, 353-61	5.3	52
144	Molecular typing of methicillin-resistant staphylococci isolated from cats and dogs. <i>Journal of Antimicrobial Chemotherapy</i> , 2006 , 58, 428-31	5.1	47
143	Convergent adaptation in the dominant global hospital clone ST239 of methicillin-resistant <i>Staphylococcus aureus</i> . <i>MBio</i> , 2015 , 6, e00080	7.8	44
142	Incidence, risk factors, and outcomes of Pantone-Valentine leukocidin-positive methicillin-susceptible <i>Staphylococcus aureus</i> infections in Auckland, New Zealand. <i>Journal of Clinical Microbiology</i> , 2010 , 48, 3470-4	9.7	43
141	Clinical and molecular epidemiology of methicillin-resistant <i>Staphylococcus aureus</i> in New Zealand: rapid emergence of sequence type 5 (ST5)-SCCmec-IV as the dominant community-associated MRSA clone. <i>PLoS ONE</i> , 2013 , 8, e62020	3.7	41

140	Comparison of a multiplexed MassARRAY system with real-time allele-specific PCR technology for genotyping of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Clinical Microbiology and Infection</i> , 2011 , 17, 1804-10	9.5	38
139	Hyperexpression of Hemolysin explains enhanced virulence of sequence type 93 community-associated methicillin-resistant <i>Staphylococcus aureus</i> . <i>BMC Microbiology</i> , 2014 , 14, 31	4.5	37
138	A British epidemic strain of methicillin-resistant <i>Staphylococcus aureus</i> (UK EMRSA-15) in Western Australia. <i>Medical Journal of Australia</i> , 2001 , 174, 662	4	37
137	Population dynamics of methicillin-susceptible and -resistant <i>Staphylococcus aureus</i> in remote communities. <i>Journal of Antimicrobial Chemotherapy</i> , 2009 , 64, 684-93	5.1	35
136	A multidrug-resistant <i>Staphylococcus epidermidis</i> clone (ST2) is an ongoing cause of hospital-acquired infection in a Western Australian hospital. <i>Journal of Clinical Microbiology</i> , 2012 , 50, 2147-51	9.7	35
135	Methicillin-resistant <i>Staphylococcus aureus</i> in a population of horses in Australia. <i>Australian Veterinary Journal</i> , 2011 , 89, 221-5	1.2	34
134	Clonal diversity and geographic distribution of methicillin-resistant <i>Staphylococcus pseudintermedius</i> from Australian animals: Discovery of novel sequence types. <i>Veterinary Microbiology</i> , 2018 , 213, 58-65	3.3	34
133	Methicillin-resistant <i>Staphylococcus aureus</i> , Western Australia. <i>Emerging Infectious Diseases</i> , 2005 , 11, 1584-90	10.2	33
132	Epidemiology and trends in the antibiotic susceptibilities of Gram-negative bacilli isolated from patients with intra-abdominal infections in the Asia-Pacific region, 2010-2013. <i>International Journal of Antimicrobial Agents</i> , 2017 , 49, 734-739	14.3	32
131	<i>Candida auris</i> Sternal Osteomyelitis in a Man from Kenya Visiting Australia, 2015. <i>Emerging Infectious Diseases</i> , 2019 , 25, 192-194	10.2	32
130	<i>Staphylococcus aureus</i> Down Under contemporary epidemiology of <i>S. aureus</i> in Australia, New Zealand, and the South West Pacific. <i>Clinical Microbiology and Infection</i> , 2014 , 20, 597-604	9.5	32
129	The molecular epidemiology of the highly virulent ST93 Australian community <i>Staphylococcus aureus</i> strain. <i>PLoS ONE</i> , 2012 , 7, e43037	3.7	32
128	Molecular Typing of ST239-MRSA-III From Diverse Geographic Locations and the Evolution of the SCC III Element During Its Intercontinental Spread. <i>Frontiers in Microbiology</i> , 2018 , 9, 1436	5.7	31
127	Rapid detection of Pantone-Valentine leukocidin in <i>Staphylococcus aureus</i> cultures by use of a lateral flow assay based on monoclonal antibodies. <i>Journal of Clinical Microbiology</i> , 2013 , 51, 487-95	9.7	31
126	The molecular epidemiology and evolution of the Pantone-Valentine leukocidin-positive, methicillin-resistant <i>Staphylococcus aureus</i> strain USA300 in Western Australia. <i>Clinical Microbiology and Infection</i> , 2009 , 15, 770-6	9.5	31
125	Evolutionary origins of the emergent ST796 clone of vancomycin resistant. <i>PeerJ</i> , 2017 , 5, e2916	3.1	31
124	Genetic and molecular predictors of high vancomycin MIC in <i>Staphylococcus aureus</i> bacteremia isolates. <i>Journal of Clinical Microbiology</i> , 2014 , 52, 3384-93	9.7	30
123	Genome sequencing and molecular characterisation of <i>Staphylococcus aureus</i> ST772-MRSA-V, "Bengal Bay Clone". <i>BMC Research Notes</i> , 2013 , 6, 548	2.3	29

122	Controlling a multicenter outbreak involving the New York/Japan methicillin-resistant <i>Staphylococcus aureus</i> clone. <i>Infection Control and Hospital Epidemiology</i> , 2007 , 28, 845-52	2	29
121	Prevalence of MRSA strains among <i>Staphylococcus aureus</i> isolated from outpatients, 2006. <i>Communicable Diseases Intelligence Quarterly Report</i> , 2009 , 33, 10-20		28
120	Evaluation of robenidine analog NCL195 as a novel broad-spectrum antibacterial agent. <i>PLoS ONE</i> , 2017 , 12, e0183457	3.7	26
119	Range Expansion and the Origin of USA300 North American Epidemic Methicillin-Resistant. <i>MBio</i> , 2018 , 9,	7.8	26
118	Type V staphylococcal cassette chromosome mec in community staphylococci from Australia. <i>Antimicrobial Agents and Chemotherapy</i> , 2005 , 49, 5129-32	5.9	26
117	Molecular Epidemiology of Methicillin-Resistant <i>Staphylococcus aureus</i> Isolated from Australian Veterinarians. <i>PLoS ONE</i> , 2016 , 11, e0146034	3.7	26
116	Adaptive change inferred from genomic population analysis of the ST93 epidemic clone of community-associated methicillin-resistant <i>Staphylococcus aureus</i> . <i>Genome Biology and Evolution</i> , 2014 , 6, 366-78	3.9	25
115	Outbreak of invasive methicillin-resistant <i>Staphylococcus aureus</i> infection associated with acupuncture and joint injection. <i>Infection Control and Hospital Epidemiology</i> , 2008 , 29, 859-65	2	25
114	Proteome analysis of <i>Helicobacter pylori</i> : major proteins of type strain NCTC 11637. <i>Pathology</i> , 2001 , 33, 365-374	1.6	25
113	<i>Staphylococcus aureus</i> plasmids without mobilization genes are mobilized by a novel conjugative plasmid from community isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2015 , 70, 649-52	5.1	23
112	Clonal Expansion of New Penicillin-Resistant Clade of <i>Neisseria meningitidis</i> Serogroup W Clonal Complex 11, Australia. <i>Emerging Infectious Diseases</i> , 2017 , 23, 1364-1367	10.2	23
111	Identification of source and sink populations for the emergence and global spread of the East-Asia clone of community-associated MRSA. <i>Genome Biology</i> , 2016 , 17, 160	18.3	23
110	Illness severity in community-onset invasive <i>Staphylococcus aureus</i> infection and the presence of virulence genes. <i>Journal of Infectious Diseases</i> , 2012 , 205, 1840-8	7	23
109	Proteome analysis of highly immunoreactive proteins of <i>Helicobacter pylori</i> . <i>Helicobacter</i> , 2002 , 7, 175-82	4.9	23
108	Molecular Characterization of Methicillin-Resistant <i>Staphylococcus aureus</i> Isolated from Australian Animals and Veterinarians. <i>Microbial Drug Resistance</i> , 2018 , 24, 203-212	2.9	22
107	<i>Staphylococcus aureus</i> ST398 detected in pigs in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2014 , 69, 1426-8	5.1	22
106	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,	7.8	22
105	A rapid (20 h) solid screening medium for detecting methicillin-resistant <i>Staphylococcus aureus</i> . <i>Journal of Hospital Infection</i> , 1998 , 40, 67-72	6.9	21

104	A prospective, randomized, controlled trial comparing transparent polyurethane and hydrocolloid dressings for central venous catheters. <i>American Journal of Infection Control</i> , 1999 , 27, 488-96	3.8	21
103	Epidemiological, clinical, outcome and antibiotic susceptibility differences between PVL positive and PVL negative <i>Staphylococcus aureus</i> infections in Western Australia: a case control study. <i>BMC Infectious Diseases</i> , 2015 , 15, 10	4	20
102	Isolation of mecC MRSA in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2016 , 71, 2348-9	5.1	20
101	Low vancomycin MICs and fecal densities reduce the sensitivity of screening methods for vancomycin resistance in Enterococci. <i>Journal of Clinical Microbiology</i> , 2014 , 52, 2829-33	9.7	20
100	Distribution of SCCmec-associated phenol-soluble modulin in staphylococci. <i>Molecular and Cellular Probes</i> , 2012 , 26, 99-103	3.3	20
99	Molecular characterization of endocarditis-associated <i>Staphylococcus aureus</i> . <i>Journal of Clinical Microbiology</i> , 2013 , 51, 2131-8	9.7	20
98	Polyclonal emergence of vanA vancomycin-resistant <i>Enterococcus faecium</i> in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2017 , 72, 998-1001	5.1	19
97	Cousins, siblings, or copies: the genomics of recurrent <i>Staphylococcus aureus</i> infections in chronic rhinosinusitis. <i>International Forum of Allergy and Rhinology</i> , 2014 , 4, 953-60	6.3	18
96	Antimicrobial susceptibility of <i>Staphylococcus aureus</i> and molecular epidemiology of methicillin-resistant <i>S. aureus</i> isolated from Australian hospital inpatients: Report from the Australian Group on Antimicrobial Resistance 2011 <i>Staphylococcus aureus</i> Surveillance Programme. <i>Journal of Global Antimicrobial Resistance</i> , 2013 , 1, 149-156	3.4	18
95	Community-acquired methicillin-resistant <i>Staphylococcus aureus</i> pneumonia: a clinical audit. <i>Respirology</i> , 2011 , 16, 926-31	3.6	18
94	Characterization of methicillin-resistant and methicillin-susceptible isolates of <i>Staphylococcus pseudintermedius</i> from cases of canine pyoderma in Australia. <i>Journal of Medical Microbiology</i> , 2014 , 63, 1228-1233	3.2	17
93	Emergence and molecular characterization of clonal complex 398 (CC398) methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) in New Zealand. <i>Journal of Antimicrobial Chemotherapy</i> , 2014 , 69, 1428-30	5.1	17
92	Genomic epidemiology and population structure of <i>Neisseria gonorrhoeae</i> from remote highly endemic Western Australian populations. <i>BMC Genomics</i> , 2018 , 19, 165	4.5	16
91	Global Scale Dissemination of ST93: A Divergent Epidemic Lineage That Has Recently Emerged From Remote Northern Australia. <i>Frontiers in Microbiology</i> , 2018 , 9, 1453	5.7	16
90	Misidentification of <i>Staphylococcus aureus</i> by the Cepheid Xpert MRSA/SA BC Assay Due to Deletions in the Gene. <i>Journal of Clinical Microbiology</i> , 2018 , 56,	9.7	16
89	Non-multiresistant methicillin-resistant <i>Staphylococcus aureus</i> bacteraemia in Sydney, Australia: emergence of EMRSA-15, Oceania, Queensland and Western Australian MRSA strains. <i>Pathology</i> , 2006 , 38, 239-44	1.6	15
88	Survey of methicillin-resistant <i>Staphylococcus aureus</i> strains from two hospitals in El Paso, Texas. <i>Journal of Clinical Microbiology</i> , 2005 , 43, 2969-72	9.7	15
87	Genomic, Antimicrobial Resistance, and Public Health Insights into spp. from Australian Chickens. <i>Journal of Clinical Microbiology</i> , 2019 , 57,	9.7	14

86	Genomic characterization of coagulase-negative staphylococci including methicillin-resistant <i>Staphylococcus sciuri</i> causing bovine mastitis. <i>Veterinary Microbiology</i> , 2018 , 219, 17-22	3.3	14
85	Genetic organization of <i>mecA</i> and <i>mecA</i> -regulatory genes in epidemic methicillin-resistant <i>Staphylococcus aureus</i> from Australia and England. <i>Journal of Antimicrobial Chemotherapy</i> , 2002 , 50, 819-24	5.1	14
84	Characterisation of Australian MRSA strains ST75- and ST883-MRSA-IV and analysis of their accessory gene regulator locus. <i>PLoS ONE</i> , 2010 , 5, e14025	3.7	14
83	Vancomycin-resistant sequence type 796 - rapid international dissemination of a new epidemic clone. <i>Antimicrobial Resistance and Infection Control</i> , 2018 , 7, 44	6.2	13
82	Development and usage of protein microarrays for the quantitative measurement of Panton-Valentine leukocidin. <i>Molecular and Cellular Probes</i> , 2014 , 28, 123-32	3.3	13
81	Linezolid-resistant ST872 <i>Enterococcus faecium</i> harbouring <i>optrA</i> and <i>cfr</i> (D) oxazolidinone resistance genes. <i>International Journal of Antimicrobial Agents</i> , 2020 , 55, 105831	14.3	13
80	Rapid detection of H and R Panton-Valentine leukocidin isoforms in <i>Staphylococcus aureus</i> by high-resolution melting analysis. <i>Diagnostic Microbiology and Infectious Disease</i> , 2010 , 67, 399-401	2.9	12
79	Characterization of Staphylococcal Cassette Chromosome Elements from Methicillin-Resistant Infections in Australian Animals. <i>MSphere</i> , 2018 , 3,	5	12
78	Preventing the introduction of methicillin-resistant <i>Staphylococcus aureus</i> into hospitals. <i>Journal of Global Antimicrobial Resistance</i> , 2014 , 2, 260-268	3.4	11
77	Community-onset <i>Staphylococcus aureus</i> Surveillance Programme annual report, 2012. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E59-69		11
76	Reverse zoonotic transmission of community-associated MRSA ST1-IV to a dairy cow. <i>International Journal of Antimicrobial Agents</i> , 2017 , 50, 125-126	14.3	10
75	Carriage of critically important antimicrobial resistant bacteria and zoonotic parasites amongst camp dogs in remote Western Australian indigenous communities. <i>Scientific Reports</i> , 2018 , 8, 8725	4.9	10
74	Australian Group on Antimicrobial Resistance Hospital-onset <i>Staphylococcus aureus</i> Surveillance Programme annual report, 2011. <i>Communicable Diseases Intelligence</i> , 2013 , 37, E210-8		10
73	Australian <i>Staphylococcus aureus</i> Sepsis Outcome Programme annual report, 2013. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E309-19		10
72	Australian Group on Antimicrobial Resistance Australian <i>Staphylococcus aureus</i> Sepsis Outcome Programme annual report, 2014. <i>Communicable Diseases Intelligence</i> , 2016 , 40, E244-54		9
71	Increased EMRSA-15 health-care worker colonization demonstrated in retrospective review of EMRSA hospital outbreaks. <i>Antimicrobial Resistance and Infection Control</i> , 2014 , 3, 7	6.2	8
70	Multiple introductions of methicillin-resistant <i>Staphylococcus aureus</i> ST612 into Western Australia associated both with human and equine reservoirs. <i>International Journal of Antimicrobial Agents</i> , 2019 , 54, 681-685	14.3	7
69	Evolution of a 72-Kilobase Cointegrant, Conjugative Multiresistance Plasmid in Community-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates from the Early 1990s. <i>Antimicrobial Agents and Chemotherapy</i> , 2019 , 63,	5.9	7

68	Characterisation of <i>Staphylococcus felis</i> isolated from cats using whole genome sequencing. <i>Veterinary Microbiology</i> , 2018 , 222, 98-104	3.3	7
67	A change in the molecular epidemiology of vancomycin resistant enterococci in Western Australia. <i>Pathology</i> , 2014 , 46, 73-5	1.6	7
66	Australian Group on Antimicrobial Resistance (AGAR) Australian <i>Staphylococcus aureus</i> Sepsis Outcome Programme (ASSOP) Annual Report 2017. <i>Communicable Diseases Intelligence (2018)</i> , 2019 , 43,	1.9	7
65	Peptidyl-Prolyl Isomerase Is Essential for Proteome Homeostasis and Virulence in <i>Burkholderia pseudomallei</i> . <i>Infection and Immunity</i> , 2019 , 87,	3.7	6
64	Genomic epidemiology of methicillin-susceptible <i>Staphylococcus aureus</i> across colonisation and skin and soft tissue infection. <i>Journal of Infection</i> , 2017 , 75, 326-335	18.9	6
63	Australian Group on Antimicrobial Resistance Australian Enterococcal Sepsis Outcome Programme annual report, 2014. <i>Communicable Diseases Intelligence</i> , 2016 , 40, E236-43		6
62	A clonal complex 12 methicillin-resistant <i>Staphylococcus aureus</i> strain, West Australian MRSA-59, harbors a novel pseudo-SCCmec element. <i>Antimicrobial Agents and Chemotherapy</i> , 2015 , 59, 7142-4	5.9	5
61	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related <i>Staphylococcus aureus</i> . <i>PLoS ONE</i> , 2020 , 15, e0228676	3.7	5
60	Intrafamilial transmission of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Emerging Infectious Diseases</i> , 2009 , 15, 1687-9	10.2	5
59	Australian Group on Antimicrobial Resistance (AGAR) Australian Gram-negative Sepsis Outcome Programme (GNSOP) Annual Report 2017. <i>Communicable Diseases Intelligence (2018)</i> , 2019 , 43,	1.9	5
58	Australian Group on Antimicrobial Resistance (AGAR) Australian Enterococcal Sepsis Outcome Programme (AESOP) Annual Report 2017. <i>Communicable Diseases Intelligence (2018)</i> , 2019 , 43,	1.9	5
57	Australian Group on Antimicrobial Resistance (AGAR) Australian Enterococcal Sepsis Outcome Programme (AESOP) Annual Report 2018. <i>Communicable Diseases Intelligence (2018)</i> , 2020 , 44,	1.9	5
56	PROTEOME ANALYSIS OF <i>HELICOBACTER PYLORI</i> : MAJOR PROTEINS OF TYPE STRAIN NCTC 11637. <i>Pathology</i> , 2001 , 33, 365-374	1.6	5
55	Severe Disease Caused by Community-Associated MRSA ST398 Type V, Australia, 2017. <i>Emerging Infectious Diseases</i> , 2019 , 25, 190-192	10.2	5
54	Australian Group on Antimicrobial Resistance Community-onset Gram-negative Surveillance Program annual report, 2010. <i>Communicable Diseases Intelligence</i> , 2013 , 37, E219-23		5
53	Australian Enterococcal Sepsis Outcome Programme annual report, 2013. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E320-6		5
52	Australian Group on Antimicrobial Resistance Australian Enterobacteriaceae Sepsis Outcome Programme annual report, 2014. <i>Communicable Diseases Intelligence</i> , 2016 , 40, E229-35		5
51	Sulfamethoxazole/trimethoprim resistance overcall by VITEK [®] 2 and BD Phoenix [™] community-associated MRSA and MSSA. <i>Journal of Antimicrobial Chemotherapy</i> , 2019 , 74, 3639-3641	5.1	4

50	Molecular characterization and evolution of the first outbreak of vancomycin-resistant <i>Enterococcus faecium</i> in Western Australia. <i>International Journal of Antimicrobial Agents</i> , 2019 , 53, 814-819	14.3	4
49	Meningococcal Disease-Associated Prophage-Like Elements Are Present in <i>Neisseria gonorrhoeae</i> and Some Commensal <i>Neisseria</i> Species. <i>Genome Biology and Evolution</i> , 2020 , 12, 3938-3950	3.9	4
48	A three-year whole genome sequencing perspective of <i>Enterococcus faecium</i> sepsis in Australia. <i>PLoS ONE</i> , 2020 , 15, e0228781	3.7	4
47	Macrolide, lincosamide and streptogramin B resistance in a dominant clone of Australian community methicillin-resistant <i>Staphylococcus aureus</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2005 , 56, 985-6	5.1	4
46	Alterations in phage-typing patterns in vancomycin-intermediate <i>Staphylococcus aureus</i> . <i>Journal of Medical Microbiology</i> , 2003 , 52, 711-714	3.2	4
45	Antimicrobial resistance and genomic insights into bovine mastitis-associated <i>Staphylococcus aureus</i> in Australia. <i>Veterinary Microbiology</i> , 2020 , 250, 108850	3.3	4
44	Genomic and Epidemiological Evidence of a Dominant Panton-Valentine Leucocidin-Positive Methicillin Resistant Lineage in Sri Lanka and Presence Among Isolates From the United Kingdom and Australia. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019 , 9, 123	5.9	3
43	Antibiotic susceptibility and molecular epidemiology of Panton-Valentine leukocidin-positive methicillin-resistant <i>Staphylococcus aureus</i> : An international survey. <i>Journal of Global Antimicrobial Resistance</i> , 2014 , 2, 43-47	3.4	3
42	Evaluation of the BD GeneOhm MRSA ACP Assay and the Cepheid GeneXpert MRSA Assay to detect genetically diverse CA-MRSA. <i>Pathology</i> , 2013 , 45, 713-5	1.6	3
41	Detection of <i>Helicobacter pylori</i> antigen in faeces by enzyme immunoassay. <i>Pathology</i> , 2001 , 33, 496-7	1.6	3
40	Australian Group on Antimicrobial Resistance (AGAR) Australian <i>Staphylococcus aureus</i> Sepsis Outcome Programme (ASSOP) Annual Report 2018. <i>Communicable Diseases Intelligence (2018)</i> , 2020 , 44,	1.9	3
39	Increasing tolerance of hospital <i>Enterococcus faecium</i> to hand-wash alcohols		3
38	Identification and characterisation of fosfomycin resistance in <i>Escherichia coli</i> urinary tract infection isolates from Australia. <i>International Journal of Antimicrobial Agents</i> , 2020 , 56, 106121	14.3	3
37	Complete Genome Sequence of a <i>Staphylococcus aureus</i> Sequence Type 612 Isolate from an Australian Horse. <i>Microbiology Resource Announcements</i> , 2018 , 7,	1.3	3
36	Australian Enterococcal Sepsis Outcome Programme, 2011. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E247-52		3
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33	Pediatric <i>Staphylococcus aureus</i> bacteremia: clinical spectrum and predictors of poor outcome. <i>Clinical Infectious Diseases</i> , 2021 ,	11.6	2

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31	Genomic characterisation of CC398 MRSA causing severe disease in Australia.. <i>International Journal of Antimicrobial Agents</i> , 2022 , 106577	14.3	2
30	Admissions for antibiotic-resistant infections in cancer patients during first year of cancer diagnosis: a cross-sectional study. <i>Internal Medicine Journal</i> , 2017 , 47, 1306-1310	1.6	1
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28	Activity of ceftaroline against community associated and healthcare associated methicillin resistant Staphylococcus aureus. <i>Pathology</i> , 2014 , 46, 71-3	1.6	1
27	Prevalence of methicillin-resistant Staphylococcus aureus colonisation in Tasmanian rural hospitals. <i>Healthcare Infection</i> , 2009 , 14, 159-163		1
26	Reply to De Angelis et al. <i>Clinical Infectious Diseases</i> , 2011 , 52, 1472-1472	11.6	1
25	Australian Group on Antimicrobial Resistance (AGAR) Australian Gram-negative Sepsis Outcome Programme (GNSOP) Annual Report 2019. <i>Communicable Diseases Intelligence (2018)</i> , 2020 , 44,	1.9	1
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23	Evolution and global transmission of a multidrug-resistant, community-associated MRSA lineage from the Indian subcontinent		1
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19	Enterobacteriaceae Sepsis Outcome Programme annual report, 2013. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E327-33		1
18	Australian Group on Antimicrobial Resistance (AGAR) Australian Staphylococcus aureus Sepsis Outcome Programme (ASSOP) Annual Report 2016. <i>Communicable Diseases Intelligence (2018)</i> , 2018 , 42,	1.9	1
17	Australian Group on Antimicrobial Resistance (AGAR) Australian Enterococcal Sepsis Outcome Programme(AESOP) Annual Report 2015. <i>Communicable Diseases Intelligence (2018)</i> , 2018 , 42,	1.9	1
16	Australian Group on Antimicrobial Resistance (AGAR) Australian Enterococcal Sepsis Outcome Programme (AESOP) Annual Report 2016. <i>Communicable Diseases Intelligence (2018)</i> , 2018 , 42,	1.9	1
15	Molecular confirmation of Escherichia coli classified as fosfomycin-resistant by the revised EUCAST MIC breakpoint.. <i>Pathology</i> , 2022 ,	1.6	1

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6	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related <i>Staphylococcus aureus</i> 2020 , 15, e0228676		
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4	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related <i>Staphylococcus aureus</i> 2020 , 15, e0228676		
3	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related <i>Staphylococcus aureus</i> 2020 , 15, e0228676		
2	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related <i>Staphylococcus aureus</i> 2020 , 15, e0228676		
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