

Geoffrey W Coombs

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/7964011/geoffrey-w-coombs-publications-by-year.pdf>

Version: 2024-04-29

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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	Apophysomyces Variabilis Infection in Transplant Recipients due to Unrecognized Infection in an Intravenous Drug-Using Donor.. <i>Transplantation</i> , 2022 , 106, e169-e171	1.8	0
174	Genomic characterisation of CC398 MRSA causing severe disease in Australia.. <i>International Journal of Antimicrobial Agents</i> , 2022 , 106577	14.3	2
173	Molecular confirmation of Escherichia coli classified as fosfomycin-resistant by the revised EUCAST MIC breakpoint.. <i>Pathology</i> , 2022 ,	1.6	1
172	The changing molecular epidemiology of Enterococcus faecium harbouring the van operon at a teaching hospital in Western Australia: A fifteen-year retrospective study.. <i>International Journal of Medical Microbiology</i> , 2021 , 312, 151546	3.7	
171	Antimicrobial Resistance in Porcine Enterococci in Australia and the Ramifications for Human Health. <i>Applied and Environmental Microbiology</i> , 2021 , 87,	4.8	1
170	Pediatric Staphylococcus aureus bacteremia: clinical spectrum and predictors of poor outcome. <i>Clinical Infectious Diseases</i> , 2021 ,	11.6	2
169	Genome-wide association studies reveal candidate genes associated to bacteraemia caused by ST93-IV CA-MRSA. <i>BMC Genomics</i> , 2021 , 22, 418	4.5	
168	Complete Genome Sequence of Community-Associated Methicillin-Resistant Staphylococcus aureus Sequence Type 1, SCC IV[2B], Isolated in the 1990s from Northern Western Australia. <i>Microbiology Resource Announcements</i> , 2021 , 10, e0079621	1.3	
167	Molecular characterization of fosfomycin-resistant Escherichia coli urinary tract infection isolates from Australia. <i>Clinical Microbiology and Infection</i> , 2021 , 27, 1360-1361	9.5	1
166	Progress towards a coordinated, national paediatric antimicrobial resistance surveillance programme: Staphylococcus aureus, enterococcal and Gram-negative bacteraemia in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2020 , 75, 1639-1644	5.1	2
165	Meningococcal Disease-Associated Prophage-Like Elements Are Present in Neisseria gonorrhoeae and Some Commensal Neisseria Species. <i>Genome Biology and Evolution</i> , 2020 , 12, 3938-3950	3.9	4
164	A three-year whole genome sequencing perspective of Enterococcus faecium sepsis in Australia. <i>PLoS ONE</i> , 2020 , 15, e0228781	3.7	4
163	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus. <i>PLoS ONE</i> , 2020 , 15, e0228676	3.7	5
162	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
161	Australian Group on Antimicrobial Resistance (AGAR) Australian Staphylococcus aureus Sepsis Outcome Programme (ASSOP) Annual Report 2018. <i>Communicable Diseases Intelligence (2018)</i> , 2020 , 44,	1.9	3
160	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
159	Australian Group on Antimicrobial Resistance (AGAR) Australian Gram-negative Sepsis Outcome Programme (GNSOP) Annual Report 2018. <i>Communicable Diseases Intelligence (2018)</i> , 2020 , 44,	1.9	1

158	Linezolid-resistant ST872 Enterococcus faecium harbouring optrA and cfr (D) oxazolidinone resistance genes. <i>International Journal of Antimicrobial Agents</i> , 2020 , 55, 105831	14.3	13
157	Antimicrobial resistance and genomic insights into bovine mastitis-associated Staphylococcus aureus in Australia. <i>Veterinary Microbiology</i> , 2020 , 250, 108850	3.3	4
156	Identification and characterisation of fosfomycin resistance in Escherichia coli urinary tract infection isolates from Australia. <i>International Journal of Antimicrobial Agents</i> , 2020 , 56, 106121	14.3	3
155	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus 2020 , 15, e0228676		
154	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus 2020 , 15, e0228676		
153	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus 2020 , 15, e0228676		
152	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus 2020 , 15, e0228676		
151	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus 2020 , 15, e0228676		
150	Diversity of bacteriophages encoding Panton-Valentine leukocidin in temporally and geographically related Staphylococcus aureus 2020 , 15, e0228676		
149	Sulfamethoxazole/trimethoprim resistance overcall by VITEK ² and BD Phoenix ^{IN} community-associated MRSA and MSSA. <i>Journal of Antimicrobial Chemotherapy</i> , 2019 , 74, 3639-3641	5.1	4
148	Multiple introductions of methicillin-resistant Staphylococcus aureus 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
147	Evolution of a 72-Kilobase Cointegrant, Conjugative Multiresistance Plasmid in Community-Associated Methicillin-Resistant Staphylococcus aureus Isolates from the Early 1990s. <i>Antimicrobial Agents and Chemotherapy</i> , 2019 , 63,	5.9	7
146	Genomic, Antimicrobial Resistance, and Public Health Insights into spp. from Australian Chickens. <i>Journal of Clinical Microbiology</i> , 2019 , 57,	9.7	14
145	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
144	The Detroit 562 Pharyngeal Immortalized Cell Line Model for the Assessment of Infectivity of Pathogenic Neisseria sp. <i>Methods in Molecular Biology</i> , 2019 , 1969, 123-133	1.4	0
143	Molecular characterization and evolution of the first outbreak of vancomycin-resistant Enterococcus faecium in Western Australia. <i>International Journal of Antimicrobial Agents</i> , 2019 , 53, 814-819	14.3	4
142	Candida auris Sternal Osteomyelitis in a Man from Kenya Visiting Australia, 2015. <i>Emerging Infectious Diseases</i> , 2019 , 25, 192-194	10.2	32
141	Peptidyl-Prolyl Isomerase Is Essential for Proteome Homeostasis and Virulence in Burkholderia pseudomallei. <i>Infection and Immunity</i> , 2019 , 87,	3.7	6

140	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
139	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
138	Australian Group on Antimicrobial Resistance (AGAR) Australian Staphylococcus aureus Sepsis Outcome Programme (ASSOP) Annual Report 2017. <i>Communicable Diseases Intelligence (2018)</i> , 2019 , 43,	1.9	7
137	Evolution and Global Transmission of a Multidrug-Resistant, Community-Associated Methicillin-Resistant Staphylococcus aureus Lineage from the Indian Subcontinent. <i>MBio</i> , 2019 , 10,	7.8	22
136	Antimicrobial-resistant CC17 Enterococcus faecium: The past, the present and the future. <i>Journal of Global Antimicrobial Resistance</i> , 2019 , 16, 36-47	3.4	55
135	Severe Disease Caused by Community-Associated MRSA ST398 Type V, Australia, 2017. <i>Emerging Infectious Diseases</i> , 2019 , 25, 190-192	10.2	5
134	Genomic characterization of coagulase-negative staphylococci including methicillin-resistant Staphylococcus sciuri causing bovine mastitis. <i>Veterinary Microbiology</i> , 2018 , 219, 17-22	3.3	14
133	Range Expansion and the Origin of USA300 North American Epidemic Methicillin-Resistant. <i>MBio</i> , 2018 , 9,	7.8	26
132	Genomic epidemiology and population structure of Neisseria gonorrhoeae from remote highly endemic Western Australian populations. <i>BMC Genomics</i> , 2018 , 19, 165	4.5	16
131	Molecular Characterization of Methicillin-Resistant Staphylococcus aureus Isolated from Australian Animals and Veterinarians. <i>Microbial Drug Resistance</i> , 2018 , 24, 203-212	2.9	22
130	Increasing tolerance of hospital to handwash alcohols. <i>Science Translational Medicine</i> , 2018 , 10,	17.5	102
129	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
128	Characterisation of Staphylococcus felis isolated from cats using whole genome sequencing. <i>Veterinary Microbiology</i> , 2018 , 222, 98-104	3.3	7
127	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
126	Misidentification of Staphylococcus aureus 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
125	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
124	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
123	Clonal diversity and geographic distribution of methicillin-resistant Staphylococcus pseudintermedius from Australian animals: Discovery of novel sequence types. <i>Veterinary Microbiology</i> , 2018 , 213, 58-65	3.3	34

122	Complete Genome Sequence of a Staphylococcus aureus Sequence Type 612 Isolate from an Australian Horse. <i>Microbiology Resource Announcements</i> , 2018 , 7,	1.3	3
121	Characterization of Staphylococcal Cassette Chromosome Elements from Methicillin-Resistant Infections in Australian Animals. <i>MSphere</i> , 2018 , 3,	5	12
120	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
119	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
118	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
117	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
116	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
115	Evaluation of robenidine analog NCL195 as a novel broad-spectrum antibacterial agent. <i>PLoS ONE</i> , 2017 , 12, e0183457	3.7	26
114	Clonal Expansion of New Penicillin-Resistant Clade of Neisseria meningitidis Serogroup W Clonal Complex 11, Australia. <i>Emerging Infectious Diseases</i> , 2017 , 23, 1364-1367	10.2	23
113	Genomic epidemiology of methicillin-susceptible Staphylococcus aureus across colonisation and skin and soft tissue infection. <i>Journal of Infection</i> , 2017 , 75, 326-335	18.9	6
112	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
111	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
110	Methicillin-resistant Staphylococcus aureus in Papua New Guinea: a community nasal colonization prevalence study. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2017 , 111, 360-362	2	1
109	Evolutionary origins of the emergent ST796 clone of vancomycin resistant. <i>PeerJ</i> , 2017 , 5, e2916	3.1	31
108	Polyclonal emergence of vanA vancomycin-resistant Enterococcus faecium in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2017 , 72, 998-1001	5.1	19
107	Combination of Vancomycin and β -Lactam Therapy for Methicillin-Resistant Staphylococcus aureus Bacteremia: A Pilot Multicenter Randomized Controlled Trial. <i>Clinical Infectious Diseases</i> , 2016 , 62, 173-180	11.6	117
106	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
105	Isolation of mecC MRSA in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2016 , 71, 2348-9	5.1	20

104	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
103	A Prescription for Resistance: Management of Staphylococcal Skin Abscesses by General Practitioners in Australia. <i>Frontiers in Microbiology</i> , 2016 , 7, 802	5.7	
102	Molecular Epidemiology of Methicillin-Resistant Staphylococcus aureus Isolated from Australian Veterinarians. <i>PLoS ONE</i> , 2016 , 11, e0146034	3.7	26
101	An updated view of plasmid conjugation and mobilization in Staphylococcus. <i>Mobile Genetic Elements</i> , 2016 , 6, e1208317		58
100	Genomic insights into the emergence and spread of international clones of healthcare-, community- and livestock-associated methicillin-resistant Staphylococcus aureus: Blurring of the traditional definitions. <i>Journal of Global Antimicrobial Resistance</i> , 2016 , 6, 95-101	3.4	80
99	Australian Group on Antimicrobial Resistance Australian Enterobacteriaceae Sepsis Outcome Programme annual report, 2014. <i>Communicable Diseases Intelligence</i> , 2016 , 40, E229-35		5
98	Australian Group on Antimicrobial Resistance Australian Enterococcal Sepsis Outcome Programme annual report, 2014. <i>Communicable Diseases Intelligence</i> , 2016 , 40, E236-43		6
97	Australian Group on Antimicrobial Resistance Australian Staphylococcus aureus Sepsis Outcome Programme annual report, 2014. <i>Communicable Diseases Intelligence</i> , 2016 , 40, E244-54		9
96	Epidemiological, clinical, outcome and antibiotic susceptibility differences between PVL positive and PVL negative Staphylococcus aureus infections in Western Australia: a case control study. <i>BMC Infectious Diseases</i> , 2015 , 15, 10	4	20
95	Convergent adaptation in the dominant global hospital clone ST239 of methicillin-resistant Staphylococcus aureus. <i>MBio</i> , 2015 , 6, e00080	7.8	44
94	Origin-of-transfer sequences facilitate mobilisation of non-conjugative antimicrobial-resistance plasmids in Staphylococcus aureus. <i>Nucleic Acids Research</i> , 2015 , 43, 7971-83	20.1	52
93	A clonal complex 12 methicillin-resistant Staphylococcus aureus strain, West Australian MRSA-59, harbors a novel pseudo-SCCmec element. <i>Antimicrobial Agents and Chemotherapy</i> , 2015 , 59, 7142-4	5.9	5
92	Staphylococcus aureus 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
91	Hyperexpression of Ehemolysin explains enhanced virulence of sequence type 93 community-associated methicillin-resistant Staphylococcus aureus. <i>BMC Microbiology</i> , 2014 , 14, 31	4.5	37
90	Staphylococcus aureus ST398 detected in pigs in Australia. <i>Journal of Antimicrobial Chemotherapy</i> , 2014 , 69, 1426-8	5.1	22
89	Characterization of methicillin-resistant and methicillin-susceptible isolates of Staphylococcus pseudintermedius from cases of canine pyoderma in Australia. <i>Journal of Medical Microbiology</i> , 2014 , 63, 1228-1233	3.2	17
88	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
87	Can interchangeability of lincosamides be assumed in clinical practice? Comparative MICs of clindamycin and lincomycin for Streptococcus pyogenes, Streptococcus agalactiae and Staphylococcus aureus. <i>Journal of Antimicrobial Chemotherapy</i> , 2014 , 69, 856-7	5.1	2

86	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
85	Molecular epidemiology of enterococcal bacteremia in Australia. <i>Journal of Clinical Microbiology</i> , 2014 , 52, 897-905	9.7	53
84	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
83	<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
82	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
81	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
80	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
79	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
78	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
77	A change in the molecular epidemiology of vancomycin resistant enterococci in Western Australia. <i>Pathology</i> , 2014 , 46, 73-5	1.6	7
76	Activity of ceftaroline against community associated and healthcare associated methicillin resistant <i>Staphylococcus aureus</i> . <i>Pathology</i> , 2014 , 46, 71-3	1.6	1
75	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
74	Australian Enterococcal Sepsis Outcome Programme, 2011. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E247-52		3
73	Hospital-onset Gram-negative Surveillance Program annual report, 2011. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E49-53		
72	Community-onset Gram-negative Surveillance Program annual report, 2012. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E54-8		2
71	Community-onset <i>Staphylococcus aureus</i> Surveillance Programme annual report, 2012. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E59-69		11
70	Australian <i>Staphylococcus aureus</i> Sepsis Outcome Programme annual report, 2013. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E309-19		10
69	Australian Enterococcal Sepsis Outcome Programme annual report, 2013. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E320-6		5

68	Enterobacteriaceae Sepsis Outcome Programme annual report, 2013. <i>Communicable Diseases Intelligence</i> , 2014 , 38, E327-33		1
67	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
66	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
65	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
64	Rapid detection of Panton-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
63	Genomic insights to control the emergence of vancomycin-resistant enterococci. <i>MBio</i> , 2013 , 4,	7.8	112
62	Molecular characterization of endocarditis-associated <i>Staphylococcus aureus</i> . <i>Journal of Clinical Microbiology</i> , 2013 , 51, 2131-8	9.7	20
61	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
60	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
59	Australian Group on Antimicrobial Resistance Enterococcus Surveillance Programme annual report, 2010. <i>Communicable Diseases Intelligence</i> , 2013 , 37, E199-209		1
58	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
57	Australian Group on Antimicrobial Resistance Community-onset Gram-negative Surveillance Program annual report, 2010. <i>Communicable Diseases Intelligence</i> , 2013 , 37, E219-23		5
56	Distribution of SCCmec-associated phenol-soluble modulins in staphylococci. <i>Molecular and Cellular Probes</i> , 2012 , 26, 99-103	3.3	20
55	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
54	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
53	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
52	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
51	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

50	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
49	Community-acquired methicillin-resistant <i>Staphylococcus aureus</i> pneumonia: a clinical audit. <i>Respirology</i> , 2011 , 16, 926-31	3.6	18
48	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
47	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
46	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
45	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
44	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
43	Reply to De Angelis et al. <i>Clinical Infectious Diseases</i> , 2011 , 52, 1472-1472	11.6	1
42	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
41	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
40	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
39	Rapid detection of H and R Pantone-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
38	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
37	<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
36	Prevalence of methicillin-resistant <i>Staphylococcus aureus</i> colonisation in Tasmanian rural hospitals. <i>Healthcare Infection</i> , 2009 , 14, 159-163		1
35	Intrafamilial transmission of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Emerging Infectious Diseases</i> , 2009 , 15, 1687-9	10.2	5
34	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
33	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

32	The molecular epidemiology and evolution of the Panton-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
31	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
30	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
29	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
28	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
27	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
26	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
25	Molecular typing of methicillin-resistant staphylococci isolated from cats and dogs. <i>Journal of Antimicrobial Chemotherapy</i> , 2006 , 58, 428-31	5.1	47
24	Methicillin-resistant <i>Staphylococcus aureus</i> clones, Western Australia. <i>Emerging Infectious Diseases</i> , 2006 , 12, 241-7	10.2	71
23	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
22	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
21	Methicillin-resistant <i>Staphylococcus aureus</i> , Western Australia. <i>Emerging Infectious Diseases</i> , 2005 , 11, 1584-90	10.2	33
20	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
19	Type V staphylococcal cassette chromosome mec in community staphylococci from Australia. <i>Antimicrobial Agents and Chemotherapy</i> , 2005 , 49, 5129-32	5.9	26
18	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
17	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
16	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
15	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

14	All methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) are not equal. <i>Healthcare Infection</i> , 2004 , 9, 17-28		
13	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
12	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
11	Proteome analysis of highly immunoreactive proteins of <i>Helicobacter pylori</i> . <i>Helicobacter</i> , 2002 , 7, 175-82	4.9	23
10	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
9	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
8	Detection of <i>Helicobacter pylori</i> antigen in faeces by enzyme immunoassay. <i>Pathology</i> , 2001 , 33, 496-7	1.6	3
7	Proteome analysis of <i>Helicobacter pylori</i> : major proteins of type strain NCTC 11637. <i>Pathology</i> , 2001 , 33, 365-374	1.6	25
6	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
5	PROTEOME ANALYSIS OF <i>HELICOBACTER PYLORI</i> : MAJOR PROTEINS OF TYPE STRAIN NCTC 11637. <i>Pathology</i> , 2001 , 33, 365-374	1.6	5
4	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
3	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
2	Increasing tolerance of hospital <i>Enterococcus faecium</i> to hand-wash alcohols		3
1	Evolution and global transmission of a multidrug-resistant, community-associated MRSA lineage from the Indian subcontinent		1