Anders Rhod Larsen

List of Publications by Year
in descending order

130	7,018	50244	64755
papers	citations	46 h-index	79 g-index

132	132	132	6406
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Heterogeneity of Staphylococcus epidermidis in prosthetic joint infections: time to reevaluate microbiological criteria?. European Journal of Clinical Microbiology and Infectious Diseases, 2022, 41, 87-97.	1.3	9
2	MRSA surveillance programmes worldwide: moving towards a harmonised international approach. International Journal of Antimicrobial Agents, 2022, 59, 106538.	1.1	8
3	Emergence of methicillin resistance predates the clinical use of antibiotics. Nature, 2022, 602, 135-141.	13.7	138
4	Dabigatran and the Risk of Staphylococcus aureus Bacteremia: A Nationwide Cohort Study. Clinical Infectious Diseases, 2021, 73, 480-486.	2.9	9
5	Evaluation of methods for detection of Î²-lactamase production in MSSA. Journal of Antimicrobial Chemotherapy, 2021, 76, 1487-1494.	1.3	3
6	Whole Genome Sequencing and Antimicrobial Resistance of Staphylococcus aureus from Surgical Site Infections in Ghana. Pathogens, 2021, 10, 196.	1.2	4
7	Increasing Incidences and Clonal Diversity of Methicillin-Resistant Staphylococcus aureus in the Nordic Countries - Results From the Nordic MRSA Surveillance. Frontiers in Microbiology, 2021, 12, 668900.	1.5	18
8	Airborne Spread of Methicillin Resistant Staphylococcus aureus From a Swine Farm. Frontiers in Veterinary Science, 2021, 8, 644729.	0.9	8
9	Dynamics of the Human Nasal Microbiota and Staphylococcus aureus CC398 Carriage in Pig Truck Drivers across One Workweek. Applied and Environmental Microbiology, 2021, 87, e0122521.	1.4	8
10	Possible misinterpretation of penicillin susceptibility in Staphylococcus aureus blood isolate due to in vitro loss of the blaZ gene. European Journal of Clinical Microbiology and Infectious Diseases, 2021, , 1.	1.3	0
11	Complete Genome Sequences of Methicillin-Resistant Staphylococcus aureus Strains 110900 and 128254, Two Representatives of the CRISPR-Cas-Carrying Sequence Type 630/ spa Type t4549 Lineage. Microbiology Resource Announcements, 2020, 9, .	0.3	5
12	Arthritis Caused by MRSA CC398 in Patient without Animal Contact, Japan. Emerging Infectious Diseases, 2020, 26, 3104-3105.	2.0	2
13	Phage-Mediated Immune Evasion and Transmission of Livestock-Associated Methicillin-ResistantStaphylococcus aureusin Humans. Emerging Infectious Diseases, 2020, 26, .	2.0	21
14	Clinical Manifestations in Children with Staphylococcal Bacteremia Positive for Panton-Valentine Leucocidin. Pediatric Infectious Disease Journal, 2020, 39, e274-e276.	1.1	3
15	Risk of hospitalization and death within 2Âyears after methicillinâ€resistant Staphylococcus aureus (MRSA) diagnosis in persons colonized or infected with livestock and nonâ€“livestockâ€associated MRSAâ€”A nationwide registerâ€based cohort study. Zoonoses and Public Health, 2020, 67, 814-822.	0.9	0
16	Whole-genome sequence profiling of antibiotic-resistant Staphylococcus aureus isolates from livestock and farm attendants in Ghana. Journal of Global Antimicrobial Resistance, 2020, 22, 527-532.	0.9	16
17	Spread of LA-MRSA CC398 in Danish mink (Neovison vison) and mink farm workers. Veterinary Microbiology, 2020, 245, 108705.	0.8	12
18	Evolution and Population Dynamics of Clonal Complex 152 Community-Associated Methicillin-Resistant Staphylococcus aureus. MSphere, 2020, 5, .	1.3	16

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19	Staphylococcus aureus induces cell-surface expression of immune stimulatory NKG2D ligands on human monocytes. Journal of Biological Chemistry, 2020, 295, 11803-11821.	1.6	10
20	Investigation of the human nasal microbiome in persons with long- and short-term exposure to methicillin-resistant Staphylococcus aureus and other bacteria from the pig farm environment. PLoS ONE, 2020, 15, e0232456.	1.1	13
21	European hedgehogs (Erinaceus europaeus) as a natural reservoir of methicillin-resistant Staphylococcus aureus carrying mecC in Denmark. PLoS ONE, 2019, 14, e0222031.	1.1	30
22	Genomic identification of cryptic susceptibility to penicillins and Î²-lactamase inhibitors in methicillin-resistant Staphylococcus aureus. Nature Microbiology, 2019, 4, 1680-1691.	5.9	47
23	Increased risk of diabetes mellitus five years after an episode of Staphylococcus aureus bacteraemia. Infectious Diseases, 2019, 51, 512-518.	1.4	2
24	Age-Dependent Increase in Incidence of Staphylococcus aureus Bacteremia, Denmark, 2008â€“2015. Emerging Infectious Diseases, 2019, 25, .	2.0	25
25	Increased risk of Staphylococcus aureus bacteremia in hemodialysisâ€”A nationwide study. Hemodialysis International, 2019, 23, 230-238.	0.4	12
26	LA-MRSA CC398 in Dairy Cattle and Veal Calf Farms Indicates Spillover From Pig Production. Frontiers in Microbiology, 2019, 10, 2733.	1.5	30
27	Genome investigations show host adaptation and transmission of LA-MRSA CC398 from pigs into Danish healthcare institutions. Scientific Reports, 2019, 9, 18655.	1.6	51
28	International travel as source of a hospital outbreak with an unusual meticillin-resistant Staphylococcus aureus clonal complex 398, Denmark, 2016. Eurosurveillance, 2019, 24, .	3.9	22
29	Age-Dependent Increase in Incidence of Staphylococcus aureus Bacteremia, Denmark, 2008â€“2015. Emerging Infectious Diseases, 2019, 25, .	2.0	2
30	SCC mec Finder, a Web-Based Tool for Typing of Staphylococcal Cassette Chromosome mec in Staphylococcus aureus Using Whole-Genome Sequence Data. MSphere, 2018, 3, .	1.3	197
31	Antibiotic resistance and molecular characteristics of Staphylococcus aureus isolated from backyard-raised pigs and pig workers. Tropical Animal Health and Production, 2018, 50, 1565-1571.	0.5	16
32	Survival of LA-MRSA in Dust from Swine Farms. Annals of Work Exposures and Health, 2018, 62, 147-156.	0.6	51
33	Range Expansion and the Origin of USA300 North American Epidemic Methicillin-Resistant Staphylococcus aureus. MBio, 2018, 9, .	1.8	42
34	Novel SCC mec type XIII (9A) identified in an ST152 methicillin-resistant Staphylococcus aureus. Infection, Genetics and Evolution, 2018, 61, 74-76.	1.0	97
35	Identification of a PVL-negative SCC mec -IVa sublineage of the methicillin-resistant Staphylococcus aureus CC80 lineage: understanding the clonal origin of CA-MRSA. Clinical Microbiology and Infection, 2018, 24, 273-278.	2.8	15
36	Drivers and Dynamics of Methicillin-Resistant Livestock-Associated Staphylococcus aureus CC398 in Pigs and Humans in Denmark. MBio, 2018, 9, .	1.8	74

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37	Human genetic variation in GLS2 is associated with development of complicated Staphylococcus aureus bacteremia. PLoS Genetics, 2018, 14, e1007667.	1.5	16
38	Staphylococcus aureus Bacteremia in Children Aged 5-18 Yearsâ€"Risk Factors in the New Millennium. Journal of Pediatrics, 2018, 203, 108-115.e3.	0.9	12
39	Global spread of three multidrug-resistant lineages of Staphylococcus epidermidis. Nature Microbiology, 2018, 3, 1175-1185.	5.9	206
40	European external quality assessments for identification, molecular typing and characterization of Staphylococcus aureus. Journal of Antimicrobial Chemotherapy, 2018, 73, 2662-2666.	1.3	6
41	Controlling Transmission of MRSA to Humans During Short-Term Visits to Swine Farms Using Dust Masks. Frontiers in Microbiology, 2018, 9, 3361.	1.5	15
42	Variable performance of four commercial chromogenic media for detection of methicillin-resistant Staphylococcus aureus isolates harbouring mecC. International Journal of Antimicrobial Agents, 2017, 50, 263-265.	1.1	2
43	Detection of mecC -Positive Staphylococcus aureus: What To Expect from Immunological Tests Targeting PBP2a?. Journal of Clinical Microbiology, 2017, 55, 1961-1963.	1.8	12
44	Transmission of Methicillin-Resistant Staphylococcus aureus to Human Volunteers Visiting a Swine Farm. Applied and Environmental Microbiology, 2017, 83, .	1.4	50
45	Comparison of Automated Antimicrobial Susceptibility Testing Systems To Detect mecC -Positive Methicillin-Resistant Staphylococcus aureus. Journal of Clinical Microbiology, 2017, 55, 3554-3556.	1.8	4
46	Origin, evolution, and global transmission of community-acquired Staphylococcus aureus ST8. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E10596-E10604.	3.3	136
47	Staphylococcal aureus Enterotoxin C and Enterotoxin-Like L Associated with Post-partum Mastitis. Frontiers in Microbiology, 2017, 8, 173.	1.5	16
48	Nasal and pharyngeal carriage of methicillin-resistant Staphylococcus sciuri among hospitalised patients and healthcare workers in a Serbian university hospital. PLoS ONE, 2017, 12, e0185181.	1.1	11
49	Livestock-associated methicillin-resistant Staphylococcus aureus is widespread in farmed mink () Tj ETQq1 1 0.784314 rgBT /Overlock	0.8	19
50	Signatures of cytoplasmic proteins in the exoproteome distinguish community- and hospital-associated methicillin-resistantStaphylococcus aureusUSA300 lineages. Virulence, 2017, 8, 891-907.	1.8	19
51	Diabetes increases the risk of disease and death due to Staphylococcus aureus bacteremia. A matched case-control and cohort study. Infectious Diseases, 2017, 49, 689-697.	1.4	15
52	Emergence of Livestock-Associated Methicillin-Resistant Staphylococcus aureus Bloodstream Infections in Denmark. Clinical Infectious Diseases, 2017, 65, 1072-1076.	2.9	78
53	Whole-genome sequencing of bloodstream Staphylococcus aureus isolates does not distinguish bacteraemia from endocarditis. Microbial Genomics, 2017, 3, .	1.0	21
54	Evaluation of a widely used culture-based method for detection of livestock-associated meticillin-resistant Staphylococcus aureus (MRSA), Denmark and Norway, 2014 to 2016. Eurosurveillance, 2017, 22, .	3.9	13

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55	Increased Age-Dependent Risk of Death Associated With lukF-PV-Positive Staphylococcus aureus Bacteremia. Open Forum Infectious Diseases, 2016, 3, ofw220.	0.4	5
56	Whole-Genome Sequencing for Routine Pathogen Surveillance in Public Health: a Population Snapshot of Invasive Staphylococcus aureus in Europe. MBio, 2016, 7, .	1.8	192
57	Evidence for Human Adaptation and Foodborne Transmission of Livestock-Associated Methicillin-ResistantStaphylococcus aureus: Table 1.. Clinical Infectious Diseases, 2016, 63, 1349-1352.	2.9	89
58	Vancomycin gene selection in the microbiome of urbanRattus norvegicusfrom hospital environment. Evolution, Medicine and Public Health, 2016, 2016, 219-226.	1.1	9
59	Long-term mortality and causes of death associated with Staphylococcus aureus bacteremia. A matched cohort study. Journal of Infection, 2016, 73, 346-357.	1.7	26
60	Copresence oftet(K) andtet(M) in Livestock-Associated Methicillin-Resistant Staphylococcus aureus Clonal Complex 398 Is Associated with Increased Fitness during Exposure to Sublethal Concentrations of Tetracycline. Antimicrobial Agents and Chemotherapy, 2016, 60, 4401-4403.	1.4	44
61	Characterization of Staphylococcus aureus from Human Immunodeficiency Virus (HIV) patients in Accra, Ghana. Journal of Infection in Developing Countries, 2016, 10, 453-456.	0.5	14
62	Ability of the GENSPEED^{Â®}MRSA test kit to detect the novelmecAhomologuemecCinStaphylococcus aureus. Apmis, 2015, 123, 478-481.	0.9	2
63	Carriage and Genetic Diversity of Methicillin-Resistant Staphylococcus aureus among Patients and Healthcare Workers in a Serbian University Hospital. PLoS ONE, 2015, 10, e0127347.	1.1	32
64	Meticillin-resistant Staphylococcus aureus CC398 is an increasing cause of disease in people with no livestock contact in Denmark, 1999 to 2011. Eurosurveillance, 2015, 20, .	3.9	130
65	Increased risk of arterial thromboembolic events after Staphylococcus aureus bacteremia: A matched cohort study. Journal of Infection, 2015, 71, 167-178.	1.7	10
66	Risk and prognosis of Staphylococcus aureus bacteremia among individuals with and without end-stage renal disease: a Danish, population-based cohort study. BMC Infectious Diseases, 2015, 15, 6.	1.3	48
67	Methicillin-resistant Staphylococcus aureus strains from Ghana include USA300. Journal of Global Antimicrobial Resistance, 2015, 3, 26-30.	0.9	26
68	Long-term persistence of a multi-resistant methicillin-susceptible Staphylococcus aureus (MR-MSSA) clone at a university hospital in southeast Sweden, without further transmission within the region. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 1415-1422.	1.3	5
69	Methicillin-resistant Staphylococcus aureus biofilm formation on dacryocystorhinostomy silicone tubes depends on the genetic lineage. Graefe's Archive for Clinical and Experimental Ophthalmology, 2015, 253, 77-82.	1.0	7
70	Inhibitory Effect of Newly-Synthesized Chalcones on Hemolytic Activity of Methicillin-Resistant Staphylococcus aureus. Polish Journal of Microbiology, 2015, 64, 379-382.	0.6	3
71	Inhibitory Effect of Newly-Synthesized Chalcones on Hemolytic Activity of Methicillin-Resistant Staphylococcus aureus. Polish Journal of Microbiology, 2015, 64, 379-82.	0.6	0
72	Utility of a newly developed Mueller-Hinton E agar for the detection of MRSA carrying the novel mecA homologue mecC. Journal of Antimicrobial Chemotherapy, 2014, 70, 1256-7.	1.3	2

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73	Origin and Evolution of European Community-Acquired Methicillin-Resistant Staphylococcus aureus. MBio, 2014, 5, e01044-14.	1.8	112
74	Increased risk of venous thromboembolism within the first year after <scp>S</scp>taphylococcus aureus bacteraemia: a nationwide observational matched cohort study. Journal of Internal Medicine, 2014, 275, 387-397.	2.7	20
75	Association between susceptibility to photodynamic oxidation and the genetic background of Staphylococcus aureus. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 577-586.	1.3	15
76	Phenotypic detection of mecC-MRSA: cefoxitin is more reliable than oxacillin. Journal of Antimicrobial Chemotherapy, 2014, 69, 133-135.	1.3	50
77	Comparing Whole-Genome Sequencing with Sanger Sequencing for spa Typing of Methicillin-Resistant Staphylococcus aureus. Journal of Clinical Microbiology, 2014, 52, 4305-4308.	1.8	179
78	Novel mutations in penicillin-binding protein genes in clinical Staphylococcus aureus isolates that are methicillin resistant on susceptibility testing, but lack the mec gene. Journal of Antimicrobial Chemotherapy, 2014, 69, 594-597.	1.3	80
79	Importance of a Diverse Isolate Collection When Defining Genotype-Specific Mass Spectra in Staphylococcus aureus. Journal of Clinical Microbiology, 2014, 52, 2738-2739.	1.8	0
80	Antibiotic susceptibility and molecular epidemiology of Pantonâ€“Valentine leukocidin-positive meticillin-resistant Staphylococcus aureus: An international survey. Journal of Global Antimicrobial Resistance, 2014, 2, 43-47.	0.9	6
81	Long-term mortality after Staphylococcus aureus spondylodiscitis: A Danish nationwide population-based cohort study. Journal of Infection, 2014, 69, 252-258.	1.7	25
82	Methicillin-Resistant Staphylococcus aureus Colonization: A Three-Year Prospective Study in a Neonatal Intensive Care Unit in Italy. PLoS ONE, 2014, 9, e87760.	1.1	28
83	Molecular Epidemiology and Antimicrobial Susceptibility of Clinical Staphylococcus aureus from Healthcare Institutions in Ghana. PLoS ONE, 2014, 9, e89716.	1.1	82
84	Insights into Nasal Carriage of Staphylococcus aureus in an Urban and a Rural Community in Ghana. PLoS ONE, 2014, 9, e96119.	1.1	52
85	Carriage frequency, diversity and methicillin resistance of Staphylococcus aureus in Danish small ruminants. Veterinary Microbiology, 2013, 163, 110-115.	0.8	69
86	Prevalence of nasal carriage and diversity of Staphylococcus aureus among inpatients and hospital staff at Korle Bu Teaching Hospital, Ghana. Journal of Global Antimicrobial Resistance, 2013, 1, 189-193.	0.9	45
87	A nationwide study of comorbidity and risk of reinfection after Staphylococcus aureus bacteraemia. Journal of Infection, 2013, 67, 199-205.	1.7	39
88	Rapid and high-resolution distinction of community-acquired and nosocomial Staphylococcus aureus isolates with identical pulsed-field gel electrophoresis patterns and spa types. International Journal of Medical Microbiology, 2013, 303, 70-75.	1.5	12
89	Epidemiology of methicillin-resistant Staphylococcus aureus carrying the novel mecC gene in Denmark corroborates a zoonotic reservoir with transmission to humans. Clinical Microbiology and Infection, 2013, 19, E16-E22.	2.8	153
90	Whole genome sequencing identifies zoonotic transmission of MRSA isolates with the novel mecA homologue mecC. EMBO Molecular Medicine, 2013, 5, 509-515.	3.3	192

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91	Livestock Origin for a Human Pandemic Clone of Community-Associated Methicillin-Resistant Staphylococcus aureus. MBio, 2013, 4, .	1.8	177
92	Use of Vitek 2 Antimicrobial Susceptibility Profile To Identify mecC in Methicillin-Resistant Staphylococcus aureus. Journal of Clinical Microbiology, 2013, 51, 2732-2734.	1.8	53
93	A Staphylococcus xylosus Isolate with a New mecC Allotype. Antimicrobial Agents and Chemotherapy, 2013, 57, 1524-1528.	1.4	67
94	Evaluation of a Modular Multiplex-PCR Methicillin-Resistant Staphylococcus aureus Detection Assay Adapted formecCDetection. Journal of Clinical Microbiology, 2013, 51, 1917-1919.	1.8	26
95	Rapid Differentiation between Livestock-Associated and Livestock-Independent Staphylococcus aureus CC398 Clades. PLoS ONE, 2013, 8, e79645.	1.1	78
96	National surveillance reveals findings of Pantonâ€“Valentine leukocidin positive meticillin-resistant Staphylococcus aureus in Serbia. Journal of Medical Microbiology, 2013, 62, 342-344.	0.7	13
97	Incidence and Characterisation of Methicillin-Resistant Staphylococcus aureus (MRSA) from Nasal Colonisation in Participants Attending a Cattle Veterinary Conference in the UK. PLoS ONE, 2013, 8, e68463.	1.1	28
98	Multilocus Sequence Typing Scheme for Staphylococcus aureus: Revision of the gmk Locus. Journal of Clinical Microbiology, 2012, 50, 2538-2539.	1.8	24
99	The newly described mecA homologue, mecALGA251, is present in methicillin-resistant Staphylococcus aureus isolates from a diverse range of host species. Journal of Antimicrobial Chemotherapy, 2012, 67, 2809-2813.	1.3	153
100	Genome Sequence of Staphylococcus aureus Strain 11819-97, an ST80-IV European Community-Acquired Methicillin-Resistant Isolate. Journal of Bacteriology, 2012, 194, 1625-1626.	1.0	31
101	Development of a real-time quadruplex PCR assay for simultaneous detection of nuc, Panton-Valentine leucocidin (PVL), mecA and homologue mecALGA251. Journal of Antimicrobial Chemotherapy, 2012, 67, 2338-2341.	1.3	93
102	Rapid detection, differentiation and typing of methicillin-resistant Staphylococcus aureus harbouring either mecA or the new mecA homologue mecALGA251. Clinical Microbiology and Infection, 2012, 18, 395-400.	2.8	322
103	Stable incidence and continued improvement in short term mortality of Staphylococcus aureus bacteraemia between 1995 and 2008. BMC Infectious Diseases, 2012, 12, 260.	1.3	51
104	Patients transferred from Libya to Denmark carried OXA-48-producing Klebsiella pneumoniae, NDM-1-producing Acinetobacter baumannii and meticillin-resistant Staphylococcus aureus. International Journal of Antimicrobial Agents, 2012, 40, 191-192.	1.1	41
105	Genetic Variability in Beta-Defensins Is Not Associated with Susceptibility to Staphylococcus aureus Bacteremia. PLoS ONE, 2012, 7, e32315.	1.1	8
106	Methicillin-Resistant Staphylococcus aureus ST9 in Pigs in Thailand. PLoS ONE, 2012, 7, e31245.	1.1	62
107	Meticillin-resistant Staphylococcus aureus with a novel mecA homologue in human and bovine populations in the UK and Denmark: a descriptive study. Lancet Infectious Diseases, The, 2011, 11, 595-603.	4.6	751
108	Molecular Epidemiology of Panton-Valentine Leukocidin-Positive Staphylococcus aureus in Spain: Emergence of the USA300 Clone in an Autochthonous Population. Journal of Clinical Microbiology, 2011, 49, 433-436.	1.8	52

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109	Distribution of Fusidic Acid Resistance Determinants in Methicillin-Resistant Staphylococcus aureus. Antimicrobial Agents and Chemotherapy, 2011, 55, 1173-1176.	1.4	48
110	Novel Types of Staphylococcal Cassette Chromosome mec Elements Identified in Clonal Complex 398 Methicillin-Resistant Staphylococcus aureus Strains. Antimicrobial Agents and Chemotherapy, 2011, 55, 3046-3050.	1.4	136
111	A Common Variant of Staphylococcal Cassette Chromosome mec Type IVa in Isolates from Copenhagen, Denmark, Is Not Detected by the BD GeneOhm Methicillin-Resistant Staphylococcus aureus Assay. Journal of Clinical Microbiology, 2009, 47, 1524-1527.	1.8	64
112	Two Distinct Clones of Methicillin-Resistant Staphylococcus aureus (MRSA) with the Same USA300 Pulsed-Field Gel Electrophoresis Profile: a Potential Pitfall for Identification of USA300 Community-Associated MRSA. Journal of Clinical Microbiology, 2009, 47, 3765-3768.	1.8	46
113	Emergence and Characterization of Community-Associated Methicillin-Resistant Staphyloccocus aureus Infections in Denmark, 1999 to 2006. Journal of Clinical Microbiology, 2009, 47, 73-78.	1.8	89
114	Comparative genomic analysis of European and Middle Eastern community-associated methicillin-resistant Staphylococcus aureus (CC80:ST80-IV) isolates by high-density microarray. Clinical Microbiology and Infection, 2009, 15, 748-755.	2.8	18
115	spa typing directly from a mecA, spa and pvl multiplex PCR assayâ€"a cost-effective improvement for methicillin-resistant Staphylococcus aureus surveillance. Clinical Microbiology and Infection, 2008, 14, 611-614.	2.8	88
116	Epidemiological differences between the UK and Ireland versus France in Staphylococcus aureus isolates resistant to fusidic acid from community-acquired skin and soft tissue infections. Journal of Antimicrobial Chemotherapy, 2008, 61, 589-594.	1.3	29
117	Presence of the epidemic European fusidic acid-resistant impetigo clone (EEFIC) of Staphylococcus aureus in France--joint authors' response. Journal of Antimicrobial Chemotherapy, 2008, 63, 421-421.	1.3	0
118	Epidemiology of European Community-Associated Methicillin-Resistant Staphylococcus aureus Clonal Complex 80 Type IV Strains Isolated in Denmark from 1993 to 2004. Journal of Clinical Microbiology, 2008, 46, 62-68.	1.8	74
119	Characterization of the Epidemic European Fusidic Acid-Resistant Impetigo Clone of Staphylococcus aureus. Journal of Clinical Microbiology, 2007, 45, 1505-1510.	1.8	90
120	Changing Epidemiology of Pediatric Staphylococcus aureus Bacteremia in Denmark From 1971 Through 2000. Pediatric Infectious Disease Journal, 2007, 26, 398-405.	1.1	59
121	Rapid Increase of Genetically Diverse Methicillin-ResistantStaphylococcus aureus,Copenhagen, Denmark. Emerging Infectious Diseases, 2007, 13, 1533-1540.	2.0	76
122	Increasing incidence but decreasing in-hospital mortality of adult Staphylococcus aureus bacteraemia between 1981 and 2000. Clinical Microbiology and Infection, 2007, 13, 257-263.	2.8	129
123	Control of a methicillin-resistant Staphylococcus aureus (MRSA) outbreak in a day-care institution. Journal of Hospital Infection, 2006, 63, 84-92.	1.4	24
124	Phenotypic Detection of Methicillin Resistance in Staphylococcus aureus by Disk Diffusion Testing and Etest on Mueller-Hinton Agar. Journal of Clinical Microbiology, 2006, 44, 4395-4399.	1.8	62
125	Epidemiology of Emerging Methicillin-Resistant Staphylococcus aureus (MRSA) in Denmark: a Nationwide Study in a Country with Low Prevalence of MRSA Infection. Journal of Clinical Microbiology, 2005, 43, 1836-1842.	1.8	152
126	Evaluation of cefoxitin 5 and 10â€‰Î¼g discs for the detection of methicillin resistance in staphylococci. Journal of Antimicrobial Chemotherapy, 2005, 55, 157-161.	1.3	40

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127	A Fusidic Acid-Resistant Epidemic Strain of Staphylococcus aureus Carries the fusB Determinant, whereas fusA Mutations Are Prevalent in Other Resistant Isolates. Antimicrobial Agents and Chemotherapy, 2004, 48, 3594-3597.	1.4	72
128	Evaluation of different disk diffusion/media combinations for detection of methicillin resistance in Staphylococcus aureus and coagulase-negative staphylococci. Apmis, 2003, 111, 905-914.	0.9	11
129	Evaluation of a cefoxitin 30 Âg disc on Iso-Sensitest agar for detection of methicillin-resistant Staphylococcus aureus. Journal of Antimicrobial Chemotherapy, 2003, 52, 204-207.	1.3	100
130	Distinct Determinants of Human Immunodeficiency Virus Type 1 RNA and DNA Loads in Vaginal and Cervical Secretions. Journal of Infectious Diseases, 1998, 177, 1214-1220.	1.9	111