

# Dominique A Caugant

## List of Publications by Year in descending order

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239  
papers

17,544  
citations

20817

60  
h-index

16650

123  
g-index

244  
all docs

244  
docs citations

244  
times ranked

10987  
citing authors

#	ARTICLE	IF	CITATIONS
1	Surveillance and control of meningococcal disease in the COVID-19 era: A Global Meningococcal Initiative review. <i>Journal of Infection</i> , 2022, 84, 289-296.	3.3	26
2	Antibiotic Treatment Regimes as a Driver of the Global Population Dynamics of a Major Gonorrhoea Lineage. <i>Molecular Biology and Evolution</i> , 2021, 38, 1249-1261.	8.9	10
3	Cost-effectiveness of meningococcal vaccination of Norwegian teenagers with a quadrivalent ACWY conjugate vaccine. <i>Human Vaccines and Immunotherapeutics</i> , 2021, 17, 2777-2787.	3.3	5
4	Molecular diagnostic assays for the detection of common bacterial meningitis pathogens: A narrative review. <i>EBioMedicine</i> , 2021, 65, 103274.	6.1	15
5	Detection of <i>Streptococcus pneumoniae</i> , <i>Neisseria meningitidis</i> and <i>Haemophilus influenzae</i> in Culture Negative Cerebrospinal Fluid Samples from Meningitis Patients Using a Multiplex Polymerase Chain Reaction in Nepal. <i>Infectious Disease Reports</i> , 2021, 13, 173-180.	3.1	2
6	Meningococcal Carriage among Household Contacts of Patients with Invasive Meningococcal Disease in Kathmandu, Nepal: A Longitudinal Study. <i>Pathogens</i> , 2021, 10, 781.	2.8	0
7	The effect of recombination on the evolution of a population of <i>Neisseria meningitidis</i> . <i>Genome Research</i> , 2021, 31, 1258-1268.	5.5	4
8	The impact of global lineage dynamics, border restrictions, and emergence of the B.1.1.7 lineage on the SARS-CoV-2 epidemic in Norway. <i>Virus Evolution</i> , 2021, 7, veab086.	4.9	6
9	<i>Neisseria meningitidis</i> : using genomics to understand diversity, evolution and pathogenesis. <i>Nature Reviews Microbiology</i> , 2020, 18, 84-96.	28.6	68
10	Next generation rapid diagnostic tests for meningitis diagnosis. <i>Journal of Infection</i> , 2020, 81, 712-718.	3.3	16
11	Antimicrobial susceptibility and clonality of <i>Streptococcus pneumoniae</i> isolates recovered from invasive disease cases during a period with changes in pneumococcal childhood vaccination, Norway, 2004–2016. <i>Vaccine</i> , 2020, 38, 5454-5463.	3.8	18
12	Meningococcal disease surveillance in the Asia–Pacific region (2020): The global meningococcal initiative. <i>Journal of Infection</i> , 2020, 81, 698-711.	3.3	51
13	Geographically widespread invasive meningococcal disease caused by a ciprofloxacin resistant non-groupable strain of the ST-175 clonal complex. <i>Journal of Infection</i> , 2020, 81, 575-584.	3.3	9
14	The global meningitis genome partnership. <i>Journal of Infection</i> , 2020, 81, 510-520.	3.3	13
15	Genomic epidemiology and population structure of <i>Neisseria gonorrhoeae</i> in Norway, 2016–2017. <i>Microbial Genomics</i> , 2020, 6, .	2.0	20
16	Sudden emergence of a <i>Neisseria gonorrhoeae</i> clade with reduced susceptibility to extended-spectrum cephalosporins, Norway. <i>Microbial Genomics</i> , 2020, 6, .	2.0	11
17	Country Data for Action: The MenAfriNet Experience in Strengthening Meningitis Surveillance in Africa. <i>Journal of Infectious Diseases</i> , 2019, 220, S137-S139.	4.0	2
18	Prevention and control of meningococcal disease: Updates from the Global Meningococcal Initiative in Eastern Europe. <i>Journal of Infection</i> , 2019, 79, 528-541.	3.3	29

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19	Genetic Meningococcal Antigen Typing System (gMATS): A genotyping tool that predicts 4CMenB strain coverage worldwide. <i>Vaccine</i> , 2019, 37, 991-1000.	3.8	64
20	Molecular studies of meningococcal and pneumococcal meningitis patients in Ethiopia. <i>Innate Immunity</i> , 2019, 25, 158-167.	2.4	2
21	Phylogenetic relationships and regional spread of meningococcal strains in the meningitis belt, 2011–2016. <i>EBioMedicine</i> , 2019, 41, 488-496.	6.1	17
22	Integrated analysis of population genomics, transcriptomics and virulence provides novel insights into <i>Streptococcus pyogenes</i> pathogenesis. <i>Nature Genetics</i> , 2019, 51, 548-559.	21.4	58
23	Narrative review of methods and findings of recent studies on the carriage of meningococci and other <i>Neisseria</i> species in the African Meningitis Belt. <i>Tropical Medicine and International Health</i> , 2019, 24, 143-154.	2.3	8
24	The Global Meningococcal Initiative meeting on prevention of meningococcal disease worldwide: Epidemiology, surveillance, hypervirulent strains, antibiotic resistance and high-risk populations. <i>Expert Review of Vaccines</i> , 2019, 18, 15-30.	4.4	136
25	Gauging the epidemic potential of a widely circulating non-invasive meningococcal strain in Africa. <i>Microbial Genomics</i> , 2019, 5, .	2.0	5
26	Meningococcal Meningitis: A Multicentric Hospital-based Study in Kathmandu, Nepal. <i>Open Microbiology Journal</i> , 2019, 13, 273-278.	0.7	3
27	Increase of invasive meningococcal serogroup W disease in Europe, 2013 to 2017. <i>Eurosurveillance</i> , 2019, 24, .	7.0	59
28	Predicting the Susceptibility of Meningococcal Serogroup B Isolates to Bactericidal Antibodies Elicited by Bivalent rLP2086, a Novel Prophylactic Vaccine. <i>MBio</i> , 2018, 9, .	4.1	53
29	Genotypic and Phenotypic Characterization of the <i>O</i> -Linked Protein Glycosylation System Reveals High Glycan Diversity in Paired Meningococcal Carriage Isolates. <i>Journal of Bacteriology</i> , 2018, 200, .	2.2	13
30	New molecular tools for meningitis diagnostics in Ethiopia – a necessary step towards improving antimicrobial prescription. <i>BMC Infectious Diseases</i> , 2018, 18, 684.	2.9	28
31	Metagenomics for investigation of an unusual meningococcal outbreak. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 1295-1296.	9.1	2
32	Acquisition of virulence genes by a carrier strain gave rise to the ongoing epidemics of meningococcal disease in West Africa. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 5510-5515.	7.1	45
33	Establishment of the European meningococcal strain collection genome library (EMSC-GL) for the 2011 to 2012 epidemiological year. <i>Eurosurveillance</i> , 2018, 23, .	7.0	8
34	Meningococcal disease in the Middle East and Africa: Findings and updates from the Global Meningococcal Initiative. <i>Journal of Infection</i> , 2017, 75, 1-11.	3.3	63
35	The epidemiology of invasive meningococcal disease in EU/EEA countries, 2004–2014. <i>Vaccine</i> , 2017, 35, 2034-2041.	3.8	156
36	Diagnosis, management, and causes of meningitis in the Gaza Strip: an analysis of guidelines, field assessment, and microbiological study. <i>Lancet</i> , The, 2017, 390, S17.	13.7	0

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37	Four years of caseâ€based surveillance of meningitis following the introduction of MenAfriVac in Moissala, Chad: lessons learned. <i>Tropical Medicine and International Health</i> , 2017, 22, 1561-1568.	2.3	4
38	Hierarchical genomic analysis of carried and invasive serogroup A <i>Neisseria meningitidis</i> during the 2011 epidemic in Chad. <i>BMC Genomics</i> , 2017, 18, 398.	2.8	15
39	Whole genome sequencing reveals within-host genetic changes in paired meningococcal carriage isolates from Ethiopia. <i>BMC Genomics</i> , 2017, 18, 407.	2.8	25
40	Outer membrane phospholipase Aâ€™s roles in <i>Helicobacter pylori</i> acid adaptation. <i>Gut Pathogens</i> , 2017, 9, 36.	3.4	9
41	The Global Meningococcal Initiative: global epidemiology, the impact of vaccines on meningococcal disease and the importance of herd protection. <i>Expert Review of Vaccines</i> , 2017, 16, 313-328.	4.4	194
42	Surveillance of Bacterial Meningitis, Ethiopia, 2012â€2013. <i>Emerging Infectious Diseases</i> , 2016, 22, 75-78.	4.3	15
43	Evaluation of Pastorex meningitis kit performance for the rapid identification of <i>Neisseria meningitidis</i> serogroup C in Nigeria. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2016, 110, 381-385.	1.8	8
44	Whole-Genome Characterization of Epidemic <i>Neisseria meningitidis</i> Serogroup C and Resurgence of Serogroup W, Niger, 2015. <i>Emerging Infectious Diseases</i> , 2016, 22, 1762-1768.	4.3	53
45	Salivary and Serum Antibody Response Against <i>Neisseria meningitidis</i> After Vaccination With Conjugate Polysaccharide Vaccines in Ethiopian Volunteers. <i>Scandinavian Journal of Immunology</i> , 2016, 84, 118-129.	2.7	13
46	Prevalence and epidemiology of meningococcal carriage in Southern Ethiopia prior to implementation of MenAfriVac, a conjugate vaccine. <i>BMC Infectious Diseases</i> , 2016, 16, 639.	2.9	20
47	Invasive Meningococcal Meningitis Serogroup C Outbreak in Northwest Nigeria, 2015 â€ Third Consecutive Outbreak of a New Strain. <i>PLOS Currents</i> , 2016, 8, .	1.4	28
48	Decreased Carriage and Genetic Shifts in the <i>Streptococcus pneumoniae</i> Population After Changing the Seven-valent to the Thirteen-valent Pneumococcal Vaccine in Norway. <i>Pediatric Infectious Disease Journal</i> , 2015, 34, 875-883.	2.0	38
49	Public Health Impact After the Introduction of PsA-TT: The First 4 Years. <i>Clinical Infectious Diseases</i> , 2015, 61, S467-S472.	5.8	29
50	Genomic Analysis of the Evolution and Global Spread of Hyper-invasive Meningococcal Lineage 5. <i>EBioMedicine</i> , 2015, 2, 234-243.	6.1	20
51	Development and Evaluation of a Multiplex Microsphere Assay for Quantitation of IgG and IgA Antibodies against <i>Neisseria meningitidis</i> Serogroup A, C, W, and Y Polysaccharides. <i>Vaccine Journal</i> , 2015, 22, 697-705.	3.1	4
52	Continuing Effectiveness of Serogroup A Meningococcal Conjugate Vaccine, Chad, 2013. <i>Emerging Infectious Diseases</i> , 2015, 21, 115-118.	4.3	32
53	Serogroup A meningococcal conjugate vaccines in Africa. <i>Expert Review of Vaccines</i> , 2015, 14, 1441-1458.	4.4	10
54	Evolutionary pathway to increased virulence and epidemic group A <i>Streptococcus</i> disease derived from 3,615 genome sequences. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E1768-76.	7.1	215

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55	Persistent low carriage of serogroup A <i>Neisseria meningitidis</i> two years after mass vaccination with the meningococcal conjugate vaccine, MenAfriVac. <i>BMC Infectious Diseases</i> , 2014, 14, 663.	2.9	52
56	Evolution of extensively drug-resistant <i>Mycobacterium tuberculosis</i> from a susceptible ancestor in a single patient. <i>Genome Biology</i> , 2014, 15, 490.	8.8	150
57	Detection of <i>Legionella</i> by cultivation and quantitative real-time polymerase chain reaction in biological waste water treatment plants in Norway. <i>Journal of Water and Health</i> , 2014, 12, 543-554.	2.6	16
58	Implications of Differential Age Distribution of Disease-Associated Meningococcal Lineages for Vaccine Development. <i>Vaccine Journal</i> , 2014, 21, 847-853.	3.1	19
59	Human antibody responses to pneumococcal surface protein A and capsular polysaccharides during acute and convalescent stages of invasive disease in adult patients. <i>Pathogens and Disease</i> , 2014, 70, 40-50.	2.0	2
60	Variability of genes encoding surface proteins used as vaccine antigens in meningococcal endemic and epidemic strain panels from Norway. <i>Vaccine</i> , 2014, 32, 2722-2731.	3.8	9
61	A broadly-protective vaccine against meningococcal disease in sub-Saharan Africa based on Generalized Modules for Membrane Antigens (GMMA). <i>Vaccine</i> , 2014, 32, 2688-2695.	3.8	55
62	Effect of a serogroup A meningococcal conjugate vaccine (PsAâ€“TT) on serogroup A meningococcal meningitis and carriage in Chad: a community study. <i>Lancet, The</i> , 2014, 383, 40-47.	13.7	230
63	Multilocus sequence typing and <i>ftsI</i> sequencing: a powerful tool for surveillance of penicillin-binding protein 3-mediated beta-lactam resistance in nontypeable <i>Haemophilus influenzae</i> . <i>BMC Microbiology</i> , 2014, 14, 131.	3.3	49
64	Sequential Outbreaks Due to a New Strain of <i>Neisseria Meningitidis</i> Serogroup C in Northern Nigeria, 2013-14. <i>PLOS Currents</i> , 2014, 6, .	1.4	50
65	Priorities for research on meningococcal disease and the impact of serogroup A vaccination in the African meningitis belt. <i>Vaccine</i> , 2013, 31, 1453-1457.	3.8	35
66	Phenotypic and genotypic characterization of meningococcal carriage and disease isolates in Burkina Faso after mass vaccination with a serogroup A conjugate vaccine. <i>BMC Infectious Diseases</i> , 2013, 13, 363.	2.9	25
67	Molecular characterization of clinical and environmental isolates of <i>Legionella pneumophila</i> in Norway, 2001â€“2008. <i>Scandinavian Journal of Infectious Diseases</i> , 2013, 45, 59-64.	1.5	8
68	Predicted strain coverage of a meningococcal multicomponent vaccine (4CMenB) in Europe: a qualitative and quantitative assessment. <i>Lancet Infectious Diseases, The</i> , 2013, 13, 416-425.	9.1	261
69	Impact of the Serogroup A Meningococcal Conjugate Vaccine, MenAfriVac, on Carriage and Herd Immunity. <i>Clinical Infectious Diseases</i> , 2013, 56, 354-363.	5.8	188
70	A Multi-country Evaluation of <i>Neisseria meningitidis</i> Serogroup B Factor Hâ€“Binding Proteins and Implications for Vaccine Coverage in Different Age Groups. <i>Pediatric Infectious Disease Journal</i> , 2013, 32, 1096-1101.	2.0	36
71	Interlaboratory Standardization of the Sandwich Enzyme-Linked Immunosorbent Assay Designed for MATS, a Rapid, Reproducible Method for Estimating the Strain Coverage of Investigational Vaccines. <i>Vaccine Journal</i> , 2012, 19, 1609-1617.	3.1	59
72	Carriage of <i>Neisseria lactamica</i> in 1- to 29-Year-Old People in Burkina Faso: Epidemiology and Molecular Characterization. <i>Journal of Clinical Microbiology</i> , 2012, 50, 4020-4027.	3.9	23

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73	Postvaccination Increase in Serotype 19A Pneumococcal Disease in Norway Is Driven by Expansion of Penicillin-Susceptible Strains of the ST199 Complex. <i>Vaccine Journal</i> , 2012, 19, 443-445.	3.1	25
74	Decline in Early Childhood Respiratory Tract Infections in the Norwegian Mother and Child Cohort Study After Introduction of Pneumococcal Conjugate Vaccination. <i>Pediatric Infectious Disease Journal</i> , 2012, 31, 951-955.	2.0	33
75	Effectively introducing a new meningococcal A conjugate vaccine in Africa: The Burkina Faso experience. <i>Vaccine</i> , 2012, 30, B40-B45.	3.8	84
76	Laboratory quality control in a multicentre meningococcal carriage study in Burkina Faso. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2012, 106, 289-297.	1.8	9
77	An Outer Membrane Vesicle Vaccine for Prevention of Serogroup A and W135 Meningococcal Disease in the African Meningitis Belt. <i>Scandinavian Journal of Immunology</i> , 2012, 76, 99-107.	2.7	24
78	Molecular Characterization of Invasive Meningococcal Isolates from Countries in the African Meningitis Belt before Introduction of a Serogroup A Conjugate Vaccine. <i>PLoS ONE</i> , 2012, 7, e46019.	2.5	46
79	Molecular typing methods for outbreak detection and surveillance of invasive disease caused by <i>Neisseria meningitidis</i> , <i>Haemophilus influenzae</i> and <i>Streptococcus pneumoniae</i> , a review. <i>Microbiology (United Kingdom)</i> , 2011, 157, 2181-2195.	1.8	32
80	Epidemic Meningococcal Meningitis, Cameroon. <i>Emerging Infectious Diseases</i> , 2011, 17, 2070-2.	4.3	6
81	Avidity of Serogroup A Meningococcal IgG Antibodies after Immunization with Different Doses of a Tetravalent A/C/Y/W135 Polysaccharide Vaccine. <i>Scandinavian Journal of Immunology</i> , 2011, 74, 87-94.	2.7	6
82	Similar Superantigen Gene Profiles and Superantigen Activity in Norwegian Isolates of Invasive and Noninvasive Group A Streptococci. <i>Scandinavian Journal of Immunology</i> , 2011, 74, 423-429.	2.7	12
83	Phenotypes of pneumococcal strains co-existing in healthy children. <i>Infection, Genetics and Evolution</i> , 2011, 11, 1703-1708.	2.3	11
84	Baseline Meningococcal Carriage in Burkina Faso before the Introduction of a Meningococcal Serogroup A Conjugate Vaccine. <i>Vaccine Journal</i> , 2011, 18, 435-443.	3.1	70
85	Meningococcal Factor H Binding Proteins in Epidemic Strains from Africa: Implications for Vaccine Development. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1302.	3.0	49
86	Molecular characteristics of pharyngeal and invasive emm3 <i>Streptococcus pyogenes</i> strains from Norway, 1988-2003. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2010, 29, 31-43.	2.9	7
87	Meningitis Dipstick Rapid Test: Evaluating Diagnostic Performance during an Urban <i>Neisseria meningitidis</i> Serogroup A Outbreak, Burkina Faso, 2007. <i>PLoS ONE</i> , 2010, 5, e11086.	2.5	21
88	Impact of a Pneumococcal Conjugate Vaccination Program on Carriage among Children in Norway. <i>Vaccine Journal</i> , 2010, 17, 325-334.	3.1	99
89	Phenotypic and genomic characterization of pneumococcus-like streptococci isolated from HIV-seropositive patients. <i>Microbiology (United Kingdom)</i> , 2010, 156, 838-848.	1.8	17
90	<i>Streptococcus pyogenes</i> Isolates Causing Severe Infections in Norway in 2006 to 2007: emm Types, Multilocus Sequence Types, and Superantigen Profiles. <i>Journal of Clinical Microbiology</i> , 2010, 48, 842-851.	3.9	50

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91	Indirect effect of conjugate pneumococcal vaccination in a 2 + 1 dose schedule. <i>Vaccine</i> , 2010, 28, 2214-2221.	3.8	69
92	Field Evaluation of Two Rapid Diagnostic Tests for <i>Neisseria meningitidis</i> Serogroup A during the 2006 Outbreak in Niger. <i>PLoS ONE</i> , 2009, 4, e7326.	2.5	14
93	Antibiotic Susceptibility and Characteristics of <i>Neisseria meningitidis</i> Isolates from the African Meningitis Belt, 2000 to 2006: Phenotypic and Genotypic Perspectives. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 1561-1566.	3.2	32
94	Seroepidemiological Study after a Long-Distance Industrial Outbreak of Legionnaires' Disease. <i>Vaccine Journal</i> , 2009, 16, 528-534.	3.1	12
95	Sequence Diversity of the Factor H Binding Protein Vaccine Candidate in Epidemiologically Relevant Strains of Serogroup B <i>Neisseria meningitidis</i> . <i>Journal of Infectious Diseases</i> , 2009, 200, 379-389.	4.0	180
96	Meningococcal carriage and disease—Population biology and evolution. <i>Vaccine</i> , 2009, 27, B64-B70.	3.8	302
97	Genetics and evolution of <i>Neisseria meningitidis</i> : Importance for the epidemiology of meningococcal disease. <i>Infection, Genetics and Evolution</i> , 2008, 8, 558-565.	2.3	96
98	Clinical isolates of <i>Staphylococcus aureus</i> from the Arkhangelsk region, Russia: antimicrobial susceptibility, molecular epidemiology, and distribution of Pantóná€Valentine leucocidin genes. <i>Apmis</i> , 2008, 116, 877-887.	2.0	34
99	Effectiveness of a 2+1 dose schedule pneumococcal conjugate vaccination programme on invasive pneumococcal disease among children in Norway. <i>Vaccine</i> , 2008, 26, 3277-3281.	3.8	140
100	Brucellar mastitis: presentation of a case and review of the literature. <i>International Journal of Infectious Diseases</i> , 2008, 12, 98-100.	3.3	5
101	Tracking Airborne Legionella and Legionella pneumophila at a Biological Treatment Plant. <i>Environmental Science &amp; Technology</i> , 2008, 42, 7360-7367.	10.0	65
102	Phenotypic and Genotypic Characterization of <i>Streptococcus pneumoniae</i> Strains Colonizing Children Attending Day-Care Centers in Norway. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2508-2518.	3.9	35
103	An Outbreak of Legionnaires Disease Caused by Long-Distance Spread from an Industrial Air Scrubber in Sarpborg, Norway. <i>Clinical Infectious Diseases</i> , 2008, 46, 61-69.	5.8	195
104	Specificity of Subcapsular Antibody Responses in Ethiopian Patients following Disease Caused by Serogroup A Meningococci. <i>Vaccine Journal</i> , 2008, 15, 863-871.	3.1	15
105	Sequence Type and <i>emm</i> Type Diversity in <i>Streptococcus pyogenes</i> Isolates Causing Invasive Disease in Norway between 1988 and 2003. <i>Journal of Clinical Microbiology</i> , 2008, 46, 2102-2105.	3.9	14
106	fimA Genotypes and Multilocus Sequence Types of <i>Porphyromonas gingivalis</i> from Patients with Periodontitis. <i>Journal of Clinical Microbiology</i> , 2008, 46, 31-42.	3.9	61
107	Genetic Diversity of <i>Porphyromonas gingivalis</i> Isolates Recovered from Single á€œRefractoryá€• Periodontitis Sites. <i>Applied and Environmental Microbiology</i> , 2008, 74, 5817-5821.	3.1	11
108	Immunogenicity of Fractional Doses of Tetravalent A/C/Y/W135 Meningococcal Polysaccharide Vaccine: Results from a Randomized Non-Inferiority Controlled Trial in Uganda. <i>PLoS Neglected Tropical Diseases</i> , 2008, 2, e342.	3.0	16



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109	Serum Antibody Responses in Ethiopian Meningitis Patients Infected with <i>Neisseria meningitidis</i> Serogroup A Sequence Type 7. <i>Vaccine Journal</i> , 2007, 14, 451-463.	3.1	10
110	Molecular surveillance of meningococcal meningitis in Africa. <i>Vaccine</i> , 2007, 25, A8-A11.	3.8	26
111	The first large epidemic of meningococcal disease caused by serogroup W135, Burkina Faso, 2002. <i>Vaccine</i> , 2007, 25, A37-A41.	3.8	53
112	Rapid Spread in Norway of an Erythromycin-Resistant Pneumococcal Clone, Despite Low Usage of Macrolides. <i>Microbial Drug Resistance</i> , 2007, 13, 29-36.	2.0	19
113	Meningitis Serogroup W135 Outbreak, Burkina Faso, 2002. <i>Emerging Infectious Diseases</i> , 2007, 13, 920-923.	4.3	46
114	Serogroup X in Meningococcal Disease, Western Kenya. <i>Emerging Infectious Diseases</i> , 2007, 13, 944-945.	4.3	35
115	Lessons from meningococcal carriage studies. <i>FEMS Microbiology Reviews</i> , 2007, 31, 52-63.	8.6	158
116	Bacterial diversity in aortic aneurysms determined by 16S ribosomal RNA gene analysis. <i>Journal of Vascular Surgery</i> , 2006, 44, 1055-1060.	1.1	104
117	High case-fatality rates of meningococcal disease in Western Norway caused by serogroup C strains belonging to both sequence type (ST)-32 and ST-11 complexes, 1985-2002. <i>Epidemiology and Infection</i> , 2006, 134, 1195-1202.	2.1	22
118	The Population Biology of <i>Neisseria meningitidis</i> : Implications for Meningococcal Disease, <i>Epidemiology and Control.</i> , 2006, , 17-35.		11
119	Conventional and molecular investigation of meningococcal isolates in relation to two outbreaks in the area of Athens, Greece. <i>Clinical Microbiology and Infection</i> , 2006, 12, 1024-1026.	6.0	13
120	Pharyngeal carriage of <i>Neisseria meningitidis</i> in 19-year-old individuals in Uganda. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2006, 100, 1159-1163.	1.8	16
121	Carriage of <i>Streptococcus pneumoniae</i> in healthy Norwegian children attending day-care centres. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2006, 25, 510-514.	2.9	7
122	Epidemic of tuberculosis in the former Soviet Union: Social and biological reasons. <i>Tuberculosis</i> , 2006, 86, 1-10.	1.9	35
123	Septicaemia due to <i>Actinobaculum schalii</i> . <i>Scandinavian Journal of Infectious Diseases</i> , 2006, 38, 735-737.	1.5	13
124	Molecular Characterization of Non-Penicillin-Susceptible <i>Streptococcus pneumoniae</i> in Norway. <i>Journal of Clinical Microbiology</i> , 2006, 44, 3225-3230.	3.9	15
125	Macrolide-Resistant <i>Streptococcus pyogenes</i> in Norway: Population Structure and Resistance Determinants. <i>Antimicrobial Agents and Chemotherapy</i> , 2006, 50, 1896-1899.	3.2	31
126	Multilocus Sequence Typing of <i>Porphyromonas gingivalis</i> Strains from Different Geographic Origins. <i>Journal of Clinical Microbiology</i> , 2006, 44, 35-41.	3.9	37



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127	Characterization of <i>Neisseria meningitidis</i> Isolates from Recent Outbreaks in Ethiopia and Comparison with Those Recovered during the Epidemic of 1988 to 1989. <i>Journal of Clinical Microbiology</i> , 2006, 44, 861-871.	3.9	30
128	Detection and characterization of $\beta$ -lactamase genes in subgingival bacteria from patients with refractory periodontitis. <i>FEMS Microbiology Letters</i> , 2005, 242, 319-324.	1.8	42
129	Rapid molecular identification of <i>Neisseria meningitidis</i> isolates using the polymerase chain reaction followed by single-stranded conformation polymorphism analysis. <i>FEMS Immunology and Medical Microbiology</i> , 2005, 45, 143-149.	2.7	3
130	Pharyngeal carriage of serogroup W135 <i>Neisseria meningitidis</i> in Hajjees and their family contacts in Morocco, Oman and Sudan. <i>Apmis</i> , 2005, 113, 182-186.	2.0	22
131	Variable-Number Tandem Repeat Analysis of Meningococcal Isolates Belonging to the Sequence Type 162 Complex. <i>Journal of Clinical Microbiology</i> , 2005, 43, 4865-4867.	3.9	11
132	Molecular Epidemiology of <i>Neisseria meningitidis</i> Isolated in the African Meningitis Belt between 1988 and 2003 Shows Dominance of Sequence Type 5 (ST-5) and ST-11 Complexes. <i>Journal of Clinical Microbiology</i> , 2005, 43, 5129-5135.	3.9	91
133	Tuberculosis in contacts need not indicate disease transmission. <i>Thorax</i> , 2005, 60, 136-137.	5.6	10
134	Use of Variable-Number Tandem Repeats To Examine Genetic Diversity of <i>Neisseria meningitidis</i> . <i>Journal of Clinical Microbiology</i> , 2005, 43, 1699-1705.	3.9	53
135	Interlaboratory Comparison of PCR-Based Identification and Genogrouping of <i>Neisseria meningitidis</i> . <i>Journal of Clinical Microbiology</i> , 2005, 43, 144-149.	3.9	89
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