Michael Kemp

List of Publications by Year in descending order

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105 2,714 29 45 papers citations h-index g-index

117 117 2862 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Whole-genome sequence analyses by a new easy-to-use software solution support the suspicion of a neonatal ward outbreak of methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) and transmission between hospitals. Infection Control and Hospital Epidemiology, 2022, 43, 947-949.	1.8	3
2	One Day in Denmark: Nationwide point-prevalence survey of human bacterial isolates and comparison of classical and whole-genome sequence-based species identification methods. PLoS ONE, 2022, 17, e0261999.	2.5	5
3	Absence of N-Acetylglucosamine Glycosylation on Listeria monocytogenes Wall Teichoic Acids Promotes Fatty Acid Tolerance by Repulsion From the Bacterial Surface. Frontiers in Microbiology, 2022, 13, .	3.5	1
4	The Global Regulator CcpA of Listeria monocytogenes Confers Sensitivity to Antimicrobial Fatty Acids. Frontiers in Microbiology, 2022, 13, 895942.	3.5	3
5	Using core genome multilocus sequence typing (cgMLST) for vancomycin-resistant Enterococcus faecium isolates to guide infection control interventions and end an outbreak. Journal of Global Antimicrobial Resistance, 2021, 24, 418-423.	2.2	12
6	A New Tool for Analyses of Whole Genome Sequences Reveals Dissemination of Specific Strains of Vancomycin-Resistant Enterococcus faecium in a Hospital. Frontiers in Medicine, 2021, 8, 733676.	2.6	1
7	Danish Whole-Genome-Sequenced Candida albicans and Candida glabrata Samples Fit into Globally Prevalent Clades. Journal of Fungi (Basel, Switzerland), 2021, 7, 962.	3.5	3
8	Core genome multi-locus sequence typing as an essential tool in a high-cost livestock-associated meticillin-resistant Staphylococcus aureus CC398 hospital outbreak. Journal of Hospital Infection, 2020, 104, 574-581.	2.9	14
9	<i>Plasmodium cynomolgi</i> as Cause of Malaria in Tourist to Southeast Asia, 2018. Emerging Infectious Diseases, 2019, 25, 1936-1939.	4.3	39
10	Complete hybrid genome assembly of clinical multidrug-resistant Bacteroides fragilis isolates enables comprehensive identification of antimicrobial-resistance genes and plasmids. Microbial Genomics, 2019, 5, .	2.0	16
11	Surveillance of vancomycin-resistant enterococci reveals shift in dominating clones and national spread of a vancomycin-variable vanA Enterococcus faecium ST1421-CT1134 clone, Denmark, 2015 to March 2019. Eurosurveillance, 2019, 24, .	7.0	40
12	Use of Loop-Mediated Isothermal Amplification in a Resource-Saving Strategy for Primary Malaria Screening in a Non-Endemic Setting. American Journal of Tropical Medicine and Hygiene, 2019, 100, 566-571.	1.4	13
13	False-Positive Diagnostics of Bordetella Pertussis using IS481 PCR is Limited in Danish Patients. Open Microbiology Journal, 2019, 13, 51-54.	0.7	1
14	Cryptosporidium Species are Frequently Present But Rarely Detected in Clinical Samples From Children with Diarrhea in a Developed Country. Pediatric Infectious Disease Journal, 2018, 37, e138-e140.	2.0	5
15	Pleural infection: a retrospective study of clinical outcome and the correlation to known etiology, co-morbidity and treatment factors. BMC Pulmonary Medicine, 2018, 18, 160.	2.0	19
16	Atypical Hand, Foot, and Mouth Disease Caused by Coxsackievirus A6 in Denmark: A Diagnostic Mimicker. Acta Dermato-Venereologica, 2018, 98, 350-354.	1.3	32
17	2017 European guideline for the management of chancroid. International Journal of STD and AIDS, 2017, 28, 324-329.	1.1	29
18	Whole-genome sequencing for identification of the source in hospital-acquired Legionnaires' disease. Journal of Hospital Infection, 2017, 96, 392-395.	2.9	10

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19	Whole Genome Sequencing of Danish Staphylococcus argenteus Reveals a Genetically Diverse Collection with Clear Separation from Staphylococcus aureus. Frontiers in Microbiology, 2017, 8, 1512.	3.5	59
20	Selecting PCR for the Diagnosis of Intestinal Parasitosis: Choice of Targets, Evaluation of In-House Assays, and Comparison with Commercial Kits. Journal of Parasitology Research, 2017, 2017, 1-6.	1.2	5
21	Virulence Factors Associated with Enterococcus Faecalis Infective Endocarditis: A Mini Review. Open Microbiology Journal, 2017, 11, 1-11.	0.7	33
22	Molecular Typing and Epidemiology of Human Listeriosis Cases, Denmark, 2002–20121. Emerging Infectious Diseases, 2016, 22, 625-633.	4.3	57
23	Fatal Septicemia Linked to Transmission of MRSA Clonal Complex 398 in Hospital and Nursing Home, Denmark. Emerging Infectious Diseases, 2016, 22, 900-902.	4.3	18
24	Bacteremia with the bovis group streptococci: species identification and association with infective endocarditis and with gastrointestinal disease. Diagnostic Microbiology and Infectious Disease, 2016, 85, 239-242.	1.8	35
25	Proteome-wide antigen discovery of novel protective vaccine candidates against Staphylococcus aureus infection. Vaccine, 2016, 34, 4602-4609.	3.8	6
26	Draft Genome Sequence of " Terrisporobacter othiniensis ―Isolated from a Blood Culture from a Human Patient. Genome Announcements, 2015, 3, .	0.8	7
27	The incidence and clinical symptomatology of Clostridium difficile infections in a community setting in a cohort of Danish patients attending general practice. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 957-967.	2.9	14
28	Ribosomal PCR and DNA sequencing for detection and identification of bacteria: experience from 6Âyears of routine analyses of patient samples. Apmis, 2014, 122, 248-255.	2.0	10
29	Variations in the Staphylococcus aureus-specific nuc gene can potentially lead to misidentification of meticillin-susceptible and -resistant S. aureus. Journal of Medical Microbiology, 2014, 63, 1020-1022.	1.8	18
30	Risk factors for <i>Clostridium difficile</i> infection in the community: a case-control study in patients in general practice, Denmark, 2009–2011. Epidemiology and Infection, 2014, 142, 1437-1448.	2.1	24
31	Risk factors for <i>Clostridium difficile</i> infection in the community: a case-control study in patients in general practice, Denmark, 2009–2011 – CORRIGENDUM. Epidemiology and Infection, 2014, 1449-1449.	2.1	0
32	Infective endocarditis caused by Bartonella quintana in Greenland. JMM Case Reports, 2014, 1, .	1.3	0
33	Osteitis in the dens of axis caused by Treponema pallidum. BMC Infectious Diseases, 2013, 13, 347.	2.9	12
34	Need for species-specific detection for the diagnosis of amoebiasis in a non-endemic setting. Scandinavian Journal of Infectious Diseases, 2013, 45, 868-871.	1.5	4
35	Performance of matrixâ€assisted laser desorptionâ€time of flight mass spectrometry for identification of clinical yeast isolates. Mycoses, 2013, 56, 229-235.	4.0	48
36	Advantages and Limitations of Ribosomal RNA PCR and DNA Sequenc-ing for Identification of Bacteria in Cardiac Valves of Danish Patients. Open Microbiology Journal, 2013, 7, 146-151.	0.7	13

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37	Species Identification of Clinical Isolates of Anaerobic Bacteria: a Comparison of Two Matrix-Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry Systems. Journal of Clinical Microbiology, 2012, 50, 542-542.	3.9	1
38	A Program Against Bacterial Bioterrorism: Improved Patient Management and Acquisition of New Knowledge on Infectious Diseases. Biosecurity and Bioterrorism, 2012, 10, 203-207.	1,2	2
39	Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry Analysis of Gram-Positive, Catalase-Negative Cocci Not Belonging to the Streptococcus or Enterococcus Genus and Benefits of Database Extension. Journal of Clinical Microbiology, 2012, 50, 1787-1791.	3.9	64
40	Clinical features of Clostridium difficile infection and molecular characterization of the isolated strains in a cohort of Danish hospitalized patients. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 185-192.	2.9	15
41	Emergence of extended-spectrum \hat{I}^2 -lactamase (ESBL)-producing Klebsiella pneumoniae in Danish hospitals; this is in part explained by spread of two CTX-M-15 clones with multilocus sequence types 15 and 16 in Zealand. International Journal of Antimicrobial Agents, 2011, 38, 180-182.	2.5	28
42	Imported brucellosis in Denmark: Molecular identification and multiple-locus variable number tandem repeat analysis (MLVA) genotyping of the bacteria. Scandinavian Journal of Infectious Diseases, 2011, 43, 536-538.	1.5	15
43	Outbreak of listeriosis caused by infected beef meat from a meals-on-wheels delivery in Denmark 2009. Clinical Microbiology and Infection, 2011, 17, 50-52.	6.0	38
44	European guideline for the management of chancroid, 2011. International Journal of STD and AIDS, 2011, 22, 241-244.	1,1	25
45	Cardiobacterium valvarum infective endocarditis and phenotypic/molecular characterization of 11 Cardiobacterium species strains. Journal of Medical Microbiology, 2011 , 60 , $522-528$.	1.8	20
46	Species Identification of Clinical Isolates of Anaerobic Bacteria: a Comparison of Two Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry Systems. Journal of Clinical Microbiology, 2011, 49, 4314-4318.	3.9	94
47	Routine ribosomal PCR and DNA sequencing for detection and identification of bacteria. Future Microbiology, 2010, 5, 1101-1107.	2.0	14
48	Q Fever in Greenland. Emerging Infectious Diseases, 2010, 16, 511-513.	4.3	25
49	Typing of vancomycin-resistant enterococci obtained from patients at Danish hospitals and detection of a genomic island specific to CC17 Enterococcus faecium. International Journal of Antimicrobial Agents, 2010, 35, 312-314.	2.5	11
50	Imported melioidosis in Danish travellers: A diagnostic challenge. Scandinavian Journal of Infectious Diseases, 2010, 42, 445-449.	1.5	12
51	Mass spectrometry: Pneumococcal meningitis verified and Brucella species identified in less than half an hour. Scandinavian Journal of Infectious Diseases, 2010, 42, 716-718.	1.5	42
52	Infective Endocarditis: Identification of Catalase-Negative, Gram-Positive Cocci from Blood Cultures by Partial 16S rRNA Gene Analysis and by Vitek 2 Examination. Open Microbiology Journal, 2010, 4, 116-122.	0.7	5
53	Substantial increase in listeriosis, Denmark 2009. Eurosurveillance, 2010, 15, .	7.0	23
54	Actinomyces species: A Danish Survey on Human Infections and Microbiological Characteristics. Open Microbiology Journal, 2009, 3, 113-120.	0.7	40

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55	Consequences of increased antibacterial consumption and change in pattern of antibacterial use in Danish hospitals. Journal of Antimicrobial Chemotherapy, 2009, 63, 812-815.	3.0	17
56	Identification of Clinically Relevant Nonhemolytic Streptococci on the Basis of Sequence Analysis of 16S-23S Intergenic Spacer Region and Partial gdh Gene. Journal of Clinical Microbiology, 2009, 47, 932-939.	3.9	32
57	Listeria monocytogenes: Maternal-foetal infections in Denmark 1994–2005. Scandinavian Journal of Infectious Diseases, 2009, 41, 21-25.	1.5	60
58	Detection of Burkholderia pseudomallei by SYBR Green Real Time PCR. The Open Pathology Journal, 2009, 3, 30-32.	1.0	7
59	PCR and DNA sequencing in establishing the aetiology of bacterial infections in children. Apmis, 2008, 116, 811-815.	2.0	9
60	Dichotomy of the human T cell response to <i>Leishmania</i> antigens. I. Th1-like response to <i>Leishmania major</i> promastigote antigens in individuals recovered from cutaneous leishmaniasis. Clinical and Experimental Immunology, 2008, 96, 410-415.	2.6	75
61	Dichotomy of the human T cell response to Leishmania antigens. II. Absent or Th2-like response to gp63 and ThI-like response to lipophosphoglycan- associated protein in cells from cured visceral leishmaniasis patients. Clinical and Experimental Immunology, 2008, 96, 416-421.	2.6	58
62	Dichotomy of the T cell response to Leishmania antigens in patients suffering from cutaneous leishmaniasis; absence or scarcity of Th1 activity is associated with severe infections. Clinical and Experimental Immunology, 2008, 100, 239-245.	2.6	39
63	Detection of anaerobic prosthetic joint infection by PCR and DNA sequencing—a case report. Monthly Notices of the Royal Astronomical Society: Letters, 2008, 79, 568-570.	3.3	12
64	Infections with beta-haemolytic streptococci: Detection by a universal PCR for bacterial DNA and DNA sequencing. Scandinavian Journal of Infectious Diseases, 2008, 40, 547-550.	1.5	5
65	Six cases of Aerococcus sanguinicola infection: Clinical relevance and bacterial identification. Scandinavian Journal of Infectious Diseases, 2008, 40, 761-765.	1.5	32
66	A Case of Helicobacter cinaedi Bacteraemia in a Previously Healthy Person with Cellulitis. Open Microbiology Journal, 2008, 2, 29-31.	0.7	36
67	Infective Arthritis: Bacterial 23S rRNA Gene Sequencing as a Supplementary Diagnostic Method. Open Microbiology Journal, 2008, 2, 85-88.	0.7	5
68	Report of the First Human Case of Caulobacter sp. Infection. Journal of Clinical Microbiology, 2007, 45, 1366-1369.	3.9	15
69	Globicatella sanguinis bacteraemia identified by partial 16S rRNA gene sequencing. Scandinavian Journal of Infectious Diseases, 2007, 39, 745-748.	1.5	10
70	Granulicatella elegans bacteraemia in patients with abdominal infections. Scandinavian Journal of Infectious Diseases, 2007, 39, 830-833.	1.5	14
71	Ribosomal DNA sequencing of streptococci: Usefulness in species identification?. International Congress Series, 2006, 1289, 155-158.	0.2	7
72	An integrated modelling system for management of the Patuxent River estuary and basin, Maryland, USA. International Journal of Remote Sensing, 2006, 27, 3705-3726.	2.9	14

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73	Ribosomal DNA sequencing: experiences from use in the Danish National Reference Laboratory for Identification of Bacteria. Apmis, 2005, 113, 621-628.	2.0	33
74	Demonstration by PCR and DNA sequencing of Corynebacterium pseudodiphtheriticum as a cause of joint infection and isolation of the same organism from a surface swab specimen from the patient. Journal of Medical Microbiology, 2005, 54, 689-691.	1.8	21
75	Ten Cases of Actinobaculum schaalii Infection: Clinical Relevance, Bacterial Identification, and Antibiotic Susceptibility. Journal of Clinical Microbiology, 2005, 43, 5305-5308.	3.9	80
76	Serodiagnosis of Leishmania donovani infections: assessment of enzyme-linked immunosorbent assays using recombinant L. donovani gene B protein (GBP) and a peptide sequence of L. donovani GBP. Transactions of the Royal Society of Tropical Medicine and Hygiene, 1999, 93, 157-160.	1.8	33
77	T-cell response in human leishmaniasis. Immunology Letters, 1999, 65, 105-108.	2.5	81
78	Leishmania-specific T cells expressing interferon-gamma (IFN- \hat{l}^3) and IL-10 upon activation are expanded in individuals cured of visceral leishmaniasis. Clinical and Experimental Immunology, 1999, 116, 500-504.	2.6	59
79	Interferon-Î ³ - and Tumour Necrosis Factor-α-Producing Cells in Humans who are Immune to Cutaneous Leishmaniasis. Scandinavian Journal of Immunology, 1999, 49, 655-659.	2.7	36
80	TheLeishmaniapromastigote surface antigen-2 (PSA-2) is specifically recognised by Th1 cells in humans with naturally acquired immunity toL. major. FEMS Immunology and Medical Microbiology, 1998, 20, 209-218.	2.7	23
81	Humoral and Cellular Immune Responses to Synthetic Peptides of theLeishmania donovaniKinetoplastid Membrane Proteinâ€11. Scandinavian Journal of Immunology, 1998, 48, 103-109.	2.7	33
82	The Leishmania promastigote surface antigen-2 (PSA-2) is specifically recognised by Th1 cells in humans with naturally acquired immunity to L. major. FEMS Immunology and Medical Microbiology, 1998, 20, 209-218.	2.7	2
83	Excretion of ciprofloxacin in sweat and multiresistant Staphylococcus epidermidis. Lancet, The, 1997, 349, 167-169.	13.7	135
84	Regulator and effector functions of Tâ€cell subsets in human <i>Leishmanaia</i> infections. Apmis, 1997, 105, 5-33.	2.0	14
85	Interferonâ $\in \hat{\mathbb{I}}^3$ Production by Human T Cells and Natural Killer Cells In Vitro in Response to Antigens from the Two Intracellular Pathogens Mycobacterium tuberculosis and Leishmania major. Scandinavian Journal of Immunology, 1997, 46, 495-499.	2.7	24
86	The contrasting roles of CD4+ T cells in intracellular infections in humans: leishmaniasis as an example. Trends in Immunology, 1996, 17, 13-16.	7.5	95
87	Serodiagnosis of Cutaneous Leishmaniasis: Assessment of an Enzyme-Linked Immunosorbent Assay Using A Peptide Sequence from Gene B Protein. American Journal of Tropical Medicine and Hygiene, 1996, 55, 490-495.	1.4	29
88	Interferonâ€Î³ and interleukinâ€4 production by human T cells recognizing <i>Leishmania donovani</i> antigens separated by SDSâ€PAGE. Apmis, 1995, 103, 131-139.	2.0	8
89	Interleukin-4 and Interferon-Gamma Production by Leishmania Stimulated Peripheral Blood Mononuclear Cells from Nonexposed Individuals. Scandinavian Journal of Immunology, 1995, 41, 343-349.	2.7	35
90	Ciprofloxacin in sweat and antibiotic resistance. Lancet, The, 1995, 346, 1235.	13.7	15

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91	Sporotrichoid cutaneous leishmaniasis due to Leishmania major of different zymodemes in the Sudan and Saudi Arabia: a comparative study. Transactions of the Royal Society of Tropical Medicine and Hygiene, 1994, 88, 552-554.	1.8	29
92	Thl-Like Human T-Cell Clones Recognizing Leishmania gp63 Inhibit Leishmania major in Human Macrophages. Scandinavian Journal of Immunology, 1994, 40, 629-635.	2.7	9
93	Dichotomy in the human CD4 ⁺ Tâ€cell response to <i>Leishmania</i> parasites. Apmis, 1994, 102, 81-88.	2.0	12
94	Longâ€ŧerm cyclosporin A nephrotoxicity in the rat. Apmis, 1994, 102, 347-355.	2.0	4
95	The major surface glycoprotein (gp63) from Leishmania major and Leishmania donovani cleaves CD4 molecules on human T cells. Journal of Immunology, 1994, 152, 4542-8.	0.8	26
96	Production of interferon-gamma and interleukin-4 by human T cells recognizing Leishmania lipophosphoglycan-associated protein. Immunology Letters, 1993, 38, 137-144.	2.5	20
97	Interferonâ€gamma and interleukinâ€4 in human <i>Leishmania donovani</i> infections. Immunology and Cell Biology, 1993, 71, 583-587.	2.3	32
98	Leishmania donovani-reactive Th1- and Th2-like T-cell clones from individuals who have recovered from visceral leishmaniasis. Infection and Immunity, 1993, 61, 1069-1073.	2.2	141
99	Prevalence of Cutaneous Leishmaniasis along the Nile River North of Khartoum (Sudan) in the Aftermath of an Epidemic in 1985. American Journal of Tropical Medicine and Hygiene, 1993, 48, 44-49.	1.4	24
100	Recognition of Leishmania antigens by T lymphocytes from nonexposed individuals. Infection and Immunity, 1992, 60, 2246-2251.	2.2	67
101	Activation of Human T Lymphocytes by Leishmania Lipophosphoglycan. Scandinavian Journal of Immunology, 1991, 33, 219-224.	2.7	37
102	ELISA Analysis of IgA Subclass Antibodies to Dietary Antigens. International Archives of Allergy and Immunology, 1988, 87, 247-253.	2.1	12
103	ANALYSIS OF RHEUMATOID FACTORS BY A BIOTINâ€AVIDIN BASED ISOTYPEâ€SPECIFIC ELISA. Acta Pathologica, Microbiologica, Et Immunologica Scandinavica Section C, Immunology, 1985, 93C, 217-223.	0.2	2
104	One Day in Denmark: Comparison of Phenotypic and Genotypic Antimicrobial Susceptibility Testing in Bacterial Isolates From Clinical Settings. Frontiers in Microbiology, 0, 13, .	3.5	11
105	Free online genome analyses reveal multiple strains in the beginning of a hospital outbreak of <i>Enterobacter hormaechei</i> carrying <i>bla</i> csub>OXA-436 carbapenemase gene. Journal of Infection Prevention, 0, , 175717742211072.	0.9	0