# Michael Kemp

List of Publications by Year
in descending order

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| 105 | 2,714 | 172457 | 233421 |
|---|---|---|---|
| papers | citations | 29 | 45 |
| | | h-index | g-index |

| 117 | 117 | 117 | 2862 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | 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 |
| 2 | Excretion of ciprofloxacin in sweat and multiresistant Staphylococcus epidermidis. Lancet, The, 1997, 349, 167-169. | 13.7 | 135 |
| 3 | 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 |
| 4 | Species Identification of Clinical Isolates of Anaerobic Bacteria: a Comparison of Two Matrix-Assisted Laser Desorption Ionizationâ€“Time of Flight Mass Spectrometry Systems. Journal of Clinical Microbiology, 2011, 49, 4314-4318. | 3.9 | 94 |
| 5 | T-cell response in human leishmaniasis. Immunology Letters, 1999, 65, 105-108. | 2.5 | 81 |
| 6 | Ten Cases of Actinobaculum schaalii Infection: Clinical Relevance, Bacterial Identification, and Antibiotic Susceptibility. Journal of Clinical Microbiology, 2005, 43, 5305-5308. | 3.9 | 80 |
| 7 | Dichotomy of the human T cell response to Leishmania antigens. I. Th1-like response to Leishmania major promastigote antigens in individuals recovered from cutaneous leishmaniasis. Clinical and Experimental Immunology, 2008, 96, 410-415. | 2.6 | 75 |
| 8 | Recognition of Leishmania antigens by T lymphocytes from nonexposed individuals. Infection and Immunity, 1992, 60, 2246-2251. | 2.2 | 67 |
| 9 | Matrix-Assisted Laser Desorption Ionizationâ€“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 |
| 10 | Listeria monocytogenes: Maternal-foetal infections in Denmark 1994â€“2005. Scandinavian Journal of Infectious Diseases, 2009, 41, 21-25. | 1.5 | 60 |
| 11 | Leishmania-specific T cells expressing interferon-gamma (IFN-Î³) and IL-10 upon activation are expanded in individuals cured of visceral leishmaniasis. Clinical and Experimental Immunology, 1999, 116, 500-504. | 2.6 | 59 |
| 12 | 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 |
| 13 | Dichotomy of the human T cell response to Leishmania antigens. II. Absent or Th2-like response to gp63 and Th1-like response to lipophosphoglycan- associated protein in cells from cured visceral leishmaniasis patients. Clinical and Experimental Immunology, 2008, 96, 416-421. | 2.6 | 58 |
| 14 | Molecular Typing and Epidemiology of Human Listeriosis Cases, Denmark, 2002â€“20121. Emerging Infectious Diseases, 2016, 22, 625-633. | 4.3 | 57 |
| 15 | 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 |
| 16 | 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 |
| 17 | Actinomyces species: A Danish Survey on Human Infections and Microbiological Characteristics. Open Microbiology Journal, 2009, 3, 113-120. | 0.7 | 40 |
| 18 | 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 |

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|---|---|---|---|
| 19 | 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 |
| 20 | Plasmodium cynomolgi as Cause of Malaria in Tourist to Southeast Asia, 2018. Emerging Infectious Diseases, 2019, 25, 1936-1939. | 4.3 | 39 |
| 21 | 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 |
| 22 | Activation of Human T Lymphocytes by Leishmania Lipophosphoglycan. Scandinavian Journal of Immunology, 1991, 33, 219-224. | 2.7 | 37 |
| 23 | Interferon-Î³- and Tumour Necrosis Factor-Î±-Producing Cells in Humans who are Immune to Cutaneous Leishmaniasis. Scandinavian Journal of Immunology, 1999, 49, 655-659. | 2.7 | 36 |
| 24 | A Case of Helicobacter cinaedi Bacteraemia in a Previously Healthy Person with Cellulitis. Open Microbiology Journal, 2008, 2, 29-31. | 0.7 | 36 |
| 25 | 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 |
| 26 | 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 |
| 27 | Humoral and Cellular Immune Responses to Synthetic Peptides of theLeishmania donovaniKinetoplastid Membrane Proteinâ€1. Scandinavian Journal of Immunology, 1998, 48, 103-109. | 2.7 | 33 |
| 28 | 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 |
| 29 | Ribosomal DNA sequencing: experiences from use in the Danish National Reference Laboratory for Identification of Bacteria. Apmis, 2005, 113, 621-628. | 2.0 | 33 |
| 30 | Virulence Factors Associated with Enterococcus Faecalis Infective Endocarditis: A Mini Review. Open Microbiology Journal, 2017, 11, 1-11. | 0.7 | 33 |
| 31 | Interferonâ€gamma and interleukinâ€4 in human Leishmania donovani infections. Immunology and Cell Biology, 1993, 71, 583-587. | 2.3 | 32 |
| 32 | Six cases of Aerococcus sanguinicola infection: Clinical relevance and bacterial identification. Scandinavian Journal of Infectious Diseases, 2008, 40, 761-765. | 1.5 | 32 |
| 33 | 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 |
| 34 | 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 |
| 35 | 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 |
| 36 | 2017 European guideline for the management of chancroid. International Journal of STD and AIDS, 2017, 28, 324-329. | 1.1 | 29 |

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|---|---|---|---|
| 37 | 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 |
| 38 | Emergence of extended-spectrum Î²-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 |
| 39 | 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 |
| 40 | Q Fever in Greenland. Emerging Infectious Diseases, 2010, 16, 511-513. | 4.3 | 25 |
| 41 | European guideline for the management of chancroid, 2011. International Journal of STD and AIDS, 2011, 22, 241-244. | 1.1 | 25 |
| 42 | Interferonâ€Î³ 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 |
| 43 | Risk factors forClostridium difficileinfection in the community: a case-control study in patients in general practice, Denmark, 2009â€“2011. Epidemiology and Infection, 2014, 142, 1437-1448. | 2.1 | 24 |
| 44 | 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 |
| 45 | 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 |
| 46 | Substantial increase in listeriosis, Denmark 2009. Eurosurveillance, 2010, 15, . | 7.0 | 23 |
| 47 | 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 |
| 48 | 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 |
| 49 | Cardiobacterium valvarum infective endocarditis and phenotypic/molecular characterization of 11 Cardiobacterium species strains. Journal of Medical Microbiology, 2011, 60, 522-528. | 1.8 | 20 |
| 50 | 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 |
| 51 | 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 |
| 52 | 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 |
| 53 | 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 |
| 54 | 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 |

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|---|---|---|---|
| 55 | Ciprofloxacin in sweat and antibiotic resistance. Lancet, The, 1995, 346, 1235. | 13.7 | 15 |
| 56 | Report of the First Human Case of Caulobacter sp. Infection. Journal of Clinical Microbiology, 2007, 45, 1366-1369. | 3.9 | 15 |
| 57 | 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 |
| 58 | 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 |
| 59 | Regulator and effector functions of Tâ€¢cell subsets in human Leishmanaia infections. Apmis, 1997, 105, 5-33. | 2.0 | 14 |
| 60 | 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 |
| 61 | Granulicatella elegans bacteraemia in patients with abdominal infections. Scandinavian Journal of Infectious Diseases, 2007, 39, 830-833. | 1.5 | 14 |
| 62 | Routine ribosomal PCR and DNA sequencing for detection and identification of bacteria. Future Microbiology, 2010, 5, 1101-1107. | 2.0 | 14 |
| 63 | 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 |
| 64 | 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 |
| 65 | 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 |
| 66 | 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 |
| 67 | ELISA Analysis of IgA Subclass Antibodies to Dietary Antigens. International Archives of Allergy and Immunology, 1988, 87, 247-253. | 2.1 | 12 |
| 68 | Dichotomy in the human CD4<sup>+</sup> Tâ€¢cell response to Leishmania parasites. Apmis, 1994, 102, 81-88. | 2.0 | 12 |
| 69 | Detection of anaerobic prosthetic joint infection by PCR and DNA sequencingâ€"a case report. Monthly Notices of the Royal Astronomical Society: Letters, 2008, 79, 568-570. | 3.3 | 12 |
| 70 | Imported melioidosis in Danish travellers: A diagnostic challenge. Scandinavian Journal of Infectious Diseases, 2010, 42, 445-449. | 1.5 | 12 |
| 71 | Osteitis in the dens of axis caused by Treponema pallidum. BMC Infectious Diseases, 2013, 13, 347. | 2.9 | 12 |
| 72 | 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 |

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|---|---|---|---|
| 73 | 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 |
| 74 | 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 |
| 75 | Globicatella sanguinis bacteraemia identified by partial 16S rRNA gene sequencing. Scandinavian Journal of Infectious Diseases, 2007, 39, 745-748. | 1.5 | 10 |
| 76 | 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 |
| 77 | Whole-genome sequencing for identification of the source in hospital-acquired Legionnaires' disease. Journal of Hospital Infection, 2017, 96, 392-395. | 2.9 | 10 |
| 78 | 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 |
| 79 | PCR and DNA sequencing in establishing the aetiology of bacterial infections in children. Apmis, 2008, 116, 811-815. | 2.0 | 9 |
| 80 | Interferonâ€Î³ and interleukinâ€4 production by human T cells recognizing Leishmania donovani antigens separated by SDSâ€PAGE. Apmis, 1995, 103, 131-139. | 2.0 | 8 |
| 81 | Ribosomal DNA sequencing of streptococci: Usefulness in species identification?. International Congress Series, 2006, 1289, 155-158. | 0.2 | 7 |
| 82 | Draft Genome Sequence of â€œ Terrisporobacter othiniensis â€•Isolated from a Blood Culture from a Human Patient. Genome Announcements, 2015, 3, . | 0.8 | 7 |
| 83 | Detection of Burkholderia pseudomallei by SYBR Green Real Time PCR. The Open Pathology Journal, 2009, 3, 30-32. | 1.0 | 7 |
| 84 | Proteome-wide antigen discovery of novel protective vaccine candidates against Staphylococcus aureus infection. Vaccine, 2016, 34, 4602-4609. | 3.8 | 6 |
| 85 | 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 |
| 86 | 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 |
| 87 | 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 |
| 88 | Infective Arthritis: Bacterial 23S rRNA Gene Sequencing as a Supplementary Diagnostic Method. Open Microbiology Journal, 2008, 2, 85-88. | 0.7 | 5 |
| 89 | 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 |
| 90 | 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 |

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|---|---|---|---|
| 91 | Longâ€term cyclosporin A nephrotoxicity in the rat. Apmis, 1994, 102, 347-355. | 2.0 | 4 |
| 92 | 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 |
| 93 | Whole-genome sequence analyses by a new easy-to-use software solution support the suspicion of a neonatal ward outbreak of methicillin-resistant Staphylococcus aureus (MRSA) and transmission between hospitals. Infection Control and Hospital Epidemiology, 2022, 43, 947-949. | 1.8 | 3 |
| 94 | 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 |
| 95 | The Global Regulator CcpA of Listeria monocytogenes Confers Sensitivity to Antimicrobial Fatty Acids. Frontiers in Microbiology, 2022, 13, 895942. | 3.5 | 3 |
| 96 | 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 |
| 97 | 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 |
| 98 | 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 |
| 99 | 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 |
| 100 | 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 |
| 101 | False-Positive Diagnostics of Bordetella Pertussis using IS481 PCR is Limited in Danish Patients. Open Microbiology Journal, 2019, 13, 51-54. | 0.7 | 1 |
| 102 | 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 |
| 103 | Risk factors for Clostridium difficile infection in the community: a case-control study in patients in general practice, Denmark, 2009â€“2011 â€“ CORRIGENDUM. Epidemiology and Infection, 2014, 142, 1449-1449. | 2.1 | 0 |
| 104 | Infective endocarditis caused by Bartonella quintana in Greenland. JMM Case Reports, 2014, 1, . | 1.3 | 0 |
| 105 | Free online genome analyses reveal multiple strains in the beginning of a hospital outbreak of Enterobacter hormaechei carrying bla<sub>OXA-436</sub> carbapenemase gene. Journal of Infection Prevention, 0, , 175717742211072. | 0.9 | 0 |