## Ulrik Stenz Justesen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Factors associated with C-reactive protein testing when prescribing antibiotics in general practice: a register-based study. , 2022, 23, 17.		3
2	Microbiological diagnosis in cardiac implantable electronic device infections detected by sonication and next-generation sequencing. Heart Rhythm, 2022, 19, 901-908.	0.3	8
3	Socioeconomic functioning in patients with brain abscess – a nationwide, population-based cohort study in Denmark. Journal of Infection, 2022, 84, 621-627.	1.7	3
4	Bacteremia With Anaerobic Bacteria and Association With Colorectal Cancer: A Population-based Cohort Study. Clinical Infectious Diseases, 2022, 75, 1747-1753.	2.9	19
5	Determinants of Antibody Response to a Third SARS-CoV-2 mRNA Vaccine Dose in Solid Organ Transplant Recipients: Results from the Prospective Cohort Study COVAC-Tx. Vaccines, 2022, 10, 565.	2.1	17
6	Dentist's Visits and Risk of Brain Abscess: A Nationwide, Population-Based Case-Control Study. Clinical Infectious Diseases, 2022, 75, 824-829.	2.9	9
7	A case report describing <i>Candida albicans</i> endophthalmitis demonstrated by 16S/18S microbiome sequencing. Acta Ophthalmologica, 2021, 99, e1536-e1537.	0.6	0
8	<i>Mycoplasma hominis</i> septic arthritis in a patient with hypogammaglobinaemia and rheumatoid arthritis. BMJ Case Reports, 2021, 14, e237798.	0.2	7
9	Preâ€hospital antibiotic therapy preceded by blood cultures in a physicianâ€manned mobile emergency care unit. Acta Anaesthesiologica Scandinavica, 2021, 65, 540-548.	0.7	2
10	Comparison of six commercially available SARS-CoV-2 antibody assays—Choice of assay depends on intended use. International Journal of Infectious Diseases, 2021, 103, 381-388.	1.5	34
11	Development of a EUCAST disk diffusion method for the susceptibility testing of rapidly growing anaerobic bacteria using Fastidious Anaerobe Agar (FAA): a development study using Bacteroides species. Clinical Microbiology and Infection, 2021, 27, 1695.e1-1695.e6.	2.8	12
12	Classification of Salmonella enterica of the (Para-)Typhoid Fever Group by Fourier-Transform Infrared (FTIR) Spectroscopy. Microorganisms, 2021, 9, 853.	1.6	17
13	Immunogenicity of SARSâ€CoVâ€2 mRNA vaccine in solid organ transplant recipients. Journal of Internal Medicine, 2021, 290, 1264-1267.	2.7	28
14	Antibody and TÂcell immune responses following mRNA COVID-19 vaccination in patients with cancer. Cancer Cell, 2021, 39, 1034-1036.	7.7	132
15	Negative SARS-CoV-2 antibodies, T-cell response and virus neutralization following full vaccination in a renal transplant recipient: a call for vigilance. Clinical Microbiology and Infection, 2021, 27, 1371-1373.	2.8	9
16	Diagnostics with clinical microbiomeâ€based identification of microorganisms in patients with brain abscesse—a prospective cohort study. Apmis, 2021, 129, 641-652.	0.9	6
17	SARS-CoV-2 seroprevalence among 7950 healthcare workers in the Region of Southern Denmark. International Journal of Infectious Diseases, 2021, 112, 96-102.	1.5	8
18	The SARS-CoV-2–neutralizing capacity of kidney transplant recipients 4 weeks after receiving a second dose of the BNT162b2 vaccine. Kidney International, 2021, 100, 1129-1131.	2.6	14

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19	Prescribing antibiotics: the use of diagnostic tests in general practice. A register-based study. Scandinavian Journal of Primary Health Care, 2021, 39, 466-475.	0.6	6
20	Detection of beta-lactamase production in clinical Prevotella species by MALDI-TOF MS method. Anaerobe, 2020, 65, 102240.	1.0	8
21	Molecular characterization of Danish ESBL/AmpC-producing Klebsiella pneumoniae from bloodstream infections, 2018. Journal of Global Antimicrobial Resistance, 2020, 22, 562-567.	0.9	10
22	Ceftriaxone treatment of spondylodiscitis and other serious infections with Cutibacterium acnes. Journal of Antimicrobial Chemotherapy, 2020, 75, 3046-3048.	1.3	5
23	Similar genomic patterns of clinical infective endocarditis and oral isolates of Streptococcus sanguinis and Streptococcus gordonii. Scientific Reports, 2020, 10, 2728.	1.6	8
24	Investigation of possible clonal transmission of carbapenemase-producing Klebsiella pneumoniae complex member isolates in Denmark using core genome MLST and National Patient Registry Data. International Journal of Antimicrobial Agents, 2020, 55, 105931.	1.1	8
25	Bacteraemia with Moryella indoligenes and Fastidiosipila sanguinis: a case report. Access Microbiology, 2020, 2, acmi000108.	0.2	4
26	Arcobacter butzleri is an opportunistic pathogen: recurrent bacteraemia in an immunocompromised patient without diarrhoea. Access Microbiology, 2020, 2, acmi000145.	0.2	5
27	Surveillance of OXA-244-producing Escherichia coli and epidemiologic investigation of cases, Denmark, January 2016 to August 2019. Eurosurveillance, 2020, 25, .	3.9	19
28	Meropenem to Children With Febrile Neutropenia Induces Monoresistant Pseudomonas aeruginosa. Journal of Pediatric Hematology/Oncology, 2020, 42, e783-e787.	0.3	2
29	Community-acquired meningitis caused by beta-haemolytic streptococci in adults: a nationwide population-based cohort study. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 2305-2310.	1.3	6
30	Physician administered antibiotics in severe infections in a prehospital setting. Resuscitation, 2019, 142, e105.	1.3	0
31	Complete hybrid genome assembly of clinical multidrug-resistant Bacteroides fragilis isolates enables comprehensive identification of antimicrobial-resistance genes and plasmids. Microbial Genomics, 2019, 5, .	1.0	16
32	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, .	3.9	40
33	Comparing identification of clinically relevant Prevotella species by VITEK MS and MALDI biotyper. Acta Microbiologica Et Immunologica Hungarica, 2019, 67, 6-13.	0.4	0
34	Antimicrobial susceptibility testing of Bacteroides fragilis using the MALDI Biotyper antibiotic susceptibility test rapid assay (MBT-ASTRA). Anaerobe, 2018, 54, 236-239.	1.0	20
35	How to isolate, identify and determine antimicrobial susceptibility of anaerobic bacteria in routine laboratories. Clinical Microbiology and Infection, 2018, 24, 1139-1148.	2.8	84
36	Typing of vancomycin-resistant enterococci with MALDI-TOF mass spectrometry in a nosocomial outbreak setting. Clinical Microbiology and Infection, 2018, 24, 1104.e1-1104.e4.	2.8	13

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37	Validation of MALDI-TOF MS Biotyper database optimized for anaerobic bacteria: The ENRIA project. Anaerobe, 2018, 54, 224-230.	1.0	47
38	A multicenter survey of antimicrobial susceptibility of Prevotella species as determined by Etest methodology. Anaerobe, 2018, 52, 9-15.	1.0	24
39	Complete Nucleotide Sequence of an <i>Escherichia coli</i> Sequence Type 410 Strain Carrying <i>bla</i> <sub>NDM-5</sub> on an IncF Multidrug Resistance Plasmid and <i>bla</i> <sub>OXA-181</sub> on an IncX3 Plasmid. Genome Announcements, 2018, 6, .	0.8	31
40	Performance of the EUCAST disc diffusion method and two MIC methods in detection of Enterobacteriaceae with reduced susceptibility to meropenem: the NordicAST CPE study. Journal of Antimicrobial Chemotherapy, 2018, 73, 2738-2747.	1.3	13
41	<i>Escherichia coli</i> Sequence Type 410 Is Causing New International High-Risk Clones. MSphere, 2018, 3, .	1.3	183
42	An overview of the data obtained during the validation of an optimized MALDI-TOF MS Biotyper database for the identification of anaerobic bacteria. Data in Brief, 2018, 18, 1484-1496.	0.5	8
43	Performance of mass spectrometric identification of clinical Prevotella species using the VITEK MS system: A prospective multi-center study. Anaerobe, 2018, 54, 205-209.	1.0	8
44	Species identification of Streptococcus bovis group isolates causing bacteremia: a comparison of two MALDI-TOF MS systems. Diagnostic Microbiology and Infectious Disease, 2017, 88, 23-25.	0.8	8
45	WGS-based surveillance of third-generation cephalosporin-resistant Escherichia coli from bloodstream infections in Denmark. Journal of Antimicrobial Chemotherapy, 2017, 72, 1922-1929.	1.3	73
46	Antimicrobial resistance in the Bacteroides fragilis group in faecal samples from patients receiving broad-spectrum antibiotics. Anaerobe, 2017, 47, 79-85.	1.0	32
47	Detection of the optrA gene in a clinical ST16 Enterococcus faecalis isolate in Denmark. Journal of Global Antimicrobial Resistance, 2017, 10, 12-13.	0.9	19
48	Antimicrobial resistance in the Bacteroides fragilis group in faecal microbiota from healthy Danish children. International Journal of Antimicrobial Agents, 2017, 49, 573-578.	1.1	8
49	A multi-center ring trial for the identification of anaerobic bacteria using MALDI-TOF MS. Anaerobe, 2017, 48, 94-97.	1.0	10
50	Prevalence of antimicrobial resistance and the cfiA resistance gene in Danish Bacteroides fragilis group isolates since 1973. International Journal of Antimicrobial Agents, 2017, 50, 552-556.	1.1	27
51	Phenotypic detection of the <i>cfiA</i> metallo-β-lactamase in <i>Bacteroides fragilis</i> with the meropenem–EDTA double-ended Etest and the ROSCO KPC/MBL Confirm Kit. Journal of Antimicrobial Chemotherapy, 2017, 72, 437-440.	1.3	11
52	Characterization of a novel blaIMP gene, blaIMP-58, using whole genome sequencing in a Pseudomonas putida isolate detected in Denmark. Diagnostic Microbiology and Infectious Disease, 2017, 87, 68-70.	0.8	2
53	Emergence of vanA Enterococcus faecium in Denmark, 2005–15. Journal of Antimicrobial Chemotherapy, 2017, 72, 2184-2190.	1.3	47
54	In silico assessment of virulence factors in strains of Streptococcus oralis and Streptococcus mitis isolated from patients with Infective Endocarditis. Journal of Medical Microbiology, 2017, 66, 1316-1323.	0.7	41

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55	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.	0.8	35
56	Molecular characterisation of the clonal emergence of high-level ciprofloxacin-monoresistant Haemophilus influenzae in the Region of Southern Denmark. Journal of Global Antimicrobial Resistance, 2016, 5, 67-70.	0.9	11
57	Incidence of Propionibacterium acnes in initially culture-negative thioglycollate broths—a prospective cohort study at a Danish University Hospital. Clinical Microbiology and Infection, 2016, 22, 941-945.	2.8	11
58	The optimization and validation of the Biotyper MALDI-TOF MS database for the identification of Gram-positive anaerobic cocci. Clinical Microbiology and Infection, 2016, 22, 793-798.	2.8	27
59	Two Serious Cases of Infection with Clostridium celatum after 40 Years in Hiding?. Journal of Clinical Microbiology, 2016, 54, 236-238.	1.8	12
60	Draft Genome Sequence of Parabacteroides goldsteinii with Putative Novel Metallo-β-Lactamases Isolated from a Blood Culture from a Human Patient. Genome Announcements, 2015, 3, .	0.8	4
61	Multiple hospital outbreaks of <i>vanA Enterococcus faecium</i> in Denmark, 2012–13, investigated by WCS, MLST and PFGE. Journal of Antimicrobial Chemotherapy, 2015, 70, 2474-2482.	1.3	93
62	First Report of Sphingomonas koreensis as a Human Pathogen in a Patient with Meningitis. Journal of Clinical Microbiology, 2015, 53, 1028-1030.	1.8	19
63	Draft Genome Sequences of <i>Sanguibacteroides justesenii</i> , gen. nov., sp. nov., Strains OUH 308042 <sup>T</sup> (= ATCC BAA-2681 <sup>T</sup> ) and OUH 334697 (= ATCC BAA-2682), Isolated from Blood Cultures from Two Different Patients. Genome Announcements, 2015, 3, .	0.8	1
64	Draft Genome Sequence of " Terrisporobacter othiniensis ―Isolated from a Blood Culture from a Human Patient. Genome Announcements, 2015, 3, .	0.8	7
65	Development of EUCAST disk diffusion method for susceptibility testing of the Bacteroides fragilis group isolates. Anaerobe, 2015, 31, 65-71.	1.0	46
66	Characterisation of a multidrug-resistant Bacteroides fragilis isolate recovered from blood of a patient in Denmark using whole-genome sequencing. International Journal of Antimicrobial Agents, 2015, 46, 117-120.	1.1	29
67	Investigation of a possible outbreak of carbapenem-resistant <i>Acinetobacter baumannii</i> in Odense, Denmark using PFGE, MLST and whole-genome-based SNPs. Journal of Antimicrobial Chemotherapy, 2015, 70, 1965-1968.	1.3	54
68	Routine disc diffusion antimicrobial susceptibility testing of Clostridium difficile and association with PCR ribotype 027. European Journal of Clinical Microbiology and Infectious Diseases, 2015, 34, 2243-2246.	1.3	8
69	Identification of antimicrobial resistance genes in multidrug-resistant clinical Bacteroides fragilis isolates by whole genome shotgun sequencing. Anaerobe, 2015, 31, 59-64.	1.0	42
70	Four Cases of Bacteremia Caused by Oscillibacter ruminantium, a Newly Described Species. Journal of Clinical Microbiology, 2014, 52, 1304-1307.	1.8	31
71	Characterization of Third-Generation Cephalosporin-ResistantEscherichia colifrom Bloodstream Infections in Denmark. Microbial Drug Resistance, 2014, 20, 316-324.	0.9	26
72	Characterization of Carbapenem Nonsusceptible Pseudomonas aeruginosa in Denmark: A Nationwide, Prospective Study. Microbial Drug Resistance, 2014, 20, 22-29.	0.9	11

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73	Echinocandin Failure Case Due to a Previously Unreported <i>FKS1</i> Mutation in Candida krusei. Antimicrobial Agents and Chemotherapy, 2014, 58, 3550-3552.	1.4	26
74	Species differentiation of Bacteroides dorei from Bacteroides vulgatus and Bacteroides ovatus from Bacteroides xylanisolvens – Back to basics. Anaerobe, 2013, 24, 1-3.	1.0	25
75	A simple and sensitive quality control method of the anaerobic atmosphere for identification and antimicrobial susceptibility testing of anaerobic bacteria. Diagnostic Microbiology and Infectious Disease, 2013, 76, 138-140.	0.8	9
76	High rates of reduced susceptibility in the Bacteroides fragilis group isolated from blood cultures – The first national survey in Denmark. International Journal of Antimicrobial Agents, 2013, 42, 188-190.	1.1	17
77	Comparison of Rosco Neo-Sensitabs with Oxoid paper disks in EUCAST disk diffusion antimicrobial susceptibility testing on Mueller–Hinton agar. European Journal of Clinical Microbiology and Infectious Diseases, 2013, 32, 621-625.	1.3	8
78	Two Cases of Ruminococcus gnavus Bacteremia Associated with Diverticulitis. Journal of Clinical Microbiology, 2013, 51, 1334-1336.	1.8	57
79	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.	1.8	1
80	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.	1.8	64
81	Multidrug-resistant Bacteroides fragilis group on the rise in Europe?. Journal of Medical Microbiology, 2012, 61, 1784-1788.	0.7	68
82	Extended-spectrum β-lactamase (ESBL) in Danish clinical isolates of Escherichia coli and Klebsiella pneumoniae: Prevalence, β-lactamase distribution, phylogroups, and co-resistance. Scandinavian Journal of Infectious Diseases, 2012, 44, 174-181.	1.5	43
83	Antimicrobial susceptibility testing of Clostridium difficile using EUCAST epidemiological cut-off values and disk diffusion correlates. Clinical Microbiology and Infection, 2012, 18, E266-E272.	2.8	47
84	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
85	Emergence of extended-spectrum β-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.	1.1	28
86	Solobacterium moorei Bacteremia: Identification, Antimicrobial Susceptibility, and Clinical Characteristics. Journal of Clinical Microbiology, 2011, 49, 2766-2768.	1.8	50
87	Bacteremia with Bacteroides pyogenes after a Cat Bite. Journal of Clinical Microbiology, 2011, 49, 3092-3093.	1.8	21
88	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.	1.8	94
89	Direct 16S rRNA gene sequencing of polymicrobial culture-negative sampleswith analysis of mixed chromatograms. Journal of Medical Microbiology, 2010, 59, 486-488.	0.7	12
90	16S rRNA Gene Sequencing in Routine Identification of Anaerobic Bacteria Isolated from Blood Cultures. Journal of Clinical Microbiology, 2010, 48, 946-948.	1.8	54

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91	Monomicrobial necrotizing fasciitis in a white male caused by hypermucoviscous Klebsiella pneumoniae. Journal of Medical Microbiology, 2009, 58, 1519-1521.	0.7	23
92	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.	1.8	32
93	Protease inhibitor plasma concentrations in HIV antiretroviral therapy. Danish Medical Bulletin, 2008, 55, 165-85.	0.3	8
94	Report of the First Human Case of Caulobacter sp. Infection. Journal of Clinical Microbiology, 2007, 45, 1366-1369.	1.8	15
95	Pharmacokinetics of Two Randomized Trials Evaluating the Safety and Efficacy of Indinavir, Saquinavir and Lopinavir in Combination with Low-Dose Ritonavir: The MaxCmin1 and 2 Trials. Basic and Clinical Pharmacology and Toxicology, 2007, 101, 339-344.	1.2	11
96	Therapeutic Drug Monitoring and Human Immunodeficiency Virus (HIV) Antiretroviral Therapy. Basic and Clinical Pharmacology and Toxicology, 2006, 98, 20-31.	1.2	20
97	Genotyping of CYP2B6 and therapeutic drug monitoring in an HIV-infected patient with high efavirenz plasma concentrations and severe CNS side-effects. Scandinavian Journal of Infectious Diseases, 2006, 38, 733-735.	1.5	7
98	The long-term pharmacokinetics and safety of adding low-dose ritonavir to a nelfinavir 1250 mg twice-daily regimen in HIV-infected patients. HIV Medicine, 2005, 6, 334-340.	1.0	7
99	Dose-dependent pharmacokinetics of delavirdine in combination with amprenavir in healthy volunteers. Journal of Antimicrobial Chemotherapy, 2004, 54, 206-210.	1.3	3
100	Pharmacokinetic Interaction between Rifampin and the Combination of Indinavir and Lowâ€Dose Ritonavir in HIVâ€Infected Patients. Clinical Infectious Diseases, 2004, 38, 426-429.	2.9	49
101	Changing incidence of central nervous system diseases in the EuroSIDA cohort. Annals of Neurology, 2004, 55, 320-328.	2.8	273
102	Simultaneous quantitative determination of the HIV protease inhibitors indinavir, amprenavir, ritonavir, lopinavir, saquinavir, nelfinavir and the nelfinavir active metabolite M8 in plasma by liquid chromatography. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2003, 783, 491-500.	1.2	47
103	The use of calcium carbonate in nelfinavir-associated diarrhoea in HIV-1-infected patients. HIV Medicine, 2003, 4, 48-52.	1.0	14
104	Low-dose indinavir in combination with low-dose ritonavir: steady-state pharmacokinetics and long-term clinical outcome follow-up. HIV Medicine, 2003, 4, 250-254.	1.0	22
105	Pharmacokinetic interaction between amprenavir and delavirdine after multiple-dose administration in healthy volunteers. British Journal of Clinical Pharmacology, 2003, 55, 100-106.	1.1	16
106	Household Transmission of Invasive Group A Streptococcus with Necrotizing Fasciitis. Scandinavian Journal of Infectious Diseases, 2003, 35, 414-415.	1.5	3
107	Diurnal variation of plasma protease inhibitor concentrations. Aids, 2002, 16, 2487-2489.	1.0	21