

# Kristian SchÄ, nning

## List of Publications by Year in descending order

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46  
papers

1,387  
citations

394390

19  
h-index

361001

35  
g-index

46  
all docs

46  
docs citations

46  
times ranked

2400  
citing authors

#	ARTICLE	IF	CITATIONS
1	HCV genome-wide analysis for development of efficient culture systems and unravelling of antiviral resistance in genotype 4. <i>Gut</i> , 2022, 71, 627-642.	12.1	7
2	Efficacy of piperacillin-tazobactam and cefotaxime against <i>Escherichia coli</i> hyperproducing TEM-1 in a mouse peritonitis infection model. <i>International Journal of Antimicrobial Agents</i> , 2022, 59, 106543.	2.5	4
3	Performance of PCR-based syndromic testing compared to bacterial culture in patients with suspected pneumonia applying microscopy for quality assessment. <i>Apmis</i> , 2022, 130, 417-426.	2.0	4
4	Incidence and severity of SARS-CoV-2 infections in liver and kidney transplant recipients in the post-vaccination era: Real-life data from Denmark. <i>American Journal of Transplantation</i> , 2022, 22, 2637-2650.	4.7	22
5	Global evolutionary analysis of chronic hepatitis C patients revealed significant effect of baseline viral resistance, including novel non-target sites, for DAA-based treatment and retreatment outcome. <i>Journal of Viral Hepatitis</i> , 2021, 28, 302-316.	2.0	7
6	An in-well direct lysis method for rapid detection of SARS-CoV-2 by real time RT-PCR in eSwab specimens. <i>Journal of Virological Methods</i> , 2021, 289, 114062.	2.1	11
7	Comparison of 16 Serological SARS-CoV-2 Immunoassays in 16 Clinical Laboratories. <i>Journal of Clinical Microbiology</i> , 2021, 59, .	3.9	97
8	Evaluation of the Hologic Panther Fusion MRSA Assay for the detection of MRSA in ESwab specimens obtained from nose, throat, and perineum. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 2177-2183.	2.9	2
9	COVID-19 versus influenza A/B superinfection in the Intensive care unit (CRISIS): Protocol for a Danish nationwide cohort study. <i>Acta Anaesthesiologica Scandinavica</i> , 2021, 65, 1345-1350.	1.6	2
10	Electronic reporting of diagnostic laboratory test results from all healthcare sectors is a cornerstone of national preparedness and control of COVID-19 in Denmark. <i>Apmis</i> , 2021, 129, 438-451.	2.0	54
11	Investigation of the introduction and dissemination of <i>vanB</i> Enterococcus faecium in the Capital Region of Denmark and development of a rapid and accurate clone-specific <i>vanB</i> E. faecium PCR. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2260-2267.	3.0	5
12	Randomised clinical trial: a 12-strain bacterial mixture versus faecal microbiota transplantation versus vancomycin for recurrent <i>Clostridioides difficile</i> infections. <i>Alimentary Pharmacology and Therapeutics</i> , 2021, 53, 999-1009.	3.7	25
13	Resistance to piperacillin/tazobactam in <i>Escherichia coli</i> resulting from extensive IS26-associated gene amplification of blaTEM-1. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 3179-3183.	3.0	48
14	Mutational change of CTX $\beta$ 15 to CTX $\beta$ 127 resulting in mecillinam resistant <i>Escherichia coli</i> during pivmecillinam treatment of a patient. <i>MicrobiologyOpen</i> , 2019, 8, e941.	3.0	9
15	Evolutionary Pathways to Persistence of Highly Fit and Resistant Hepatitis C Virus Protease Inhibitor Escape Variants. <i>Hepatology</i> , 2019, 70, 771-787.	7.3	46
16	Children Attending Day Care Centers are a Year-round Reservoir of Gastrointestinal Viruses. <i>Scientific Reports</i> , 2019, 9, 3286.	3.3	31
17	Genome Sequence of an Unknown Subtype of Hepatitis C Virus Genotype 6: Another Piece for the Taxonomic Puzzle. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	3
18	HCV genotype 1-6 NS3 residue 80 substitutions impact protease inhibitor activity and promote viral escape. <i>Journal of Hepatology</i> , 2019, 70, 388-397.	3.7	34

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19	Surveillance of vancomycin-resistant enterococci reveals shift in dominating clones and national spread of a vancomycin-variable vanA <i>Enterococcus faecium</i> ST1421-CT1134 clone, Denmark, 2015 to March 2019. <i>Eurosurveillance</i> , 2019, 24, .	7.0	40
20	Direct acting antiviral treatment of chronic hepatitis C in Denmark: factors associated with and barriers to treatment initiation. <i>Scandinavian Journal of Gastroenterology</i> , 2018, 53, 849-856.	1.5	16
21	Evaluation of dual target-specific real-time PCR for the detection of <i>Kingella kingae</i> in a Danish paediatric population. <i>Infectious Diseases</i> , 2018, 50, 200-206.	2.8	2
22	Outcome and adverse events in patients with chronic hepatitis C treated with direct-acting antivirals: a clinical randomized study. <i>European Journal of Gastroenterology and Hepatology</i> , 2018, 30, 1177-1186.	1.6	7
23	Analytical performance of the Hologic Aptima HBV Quant Assay and the COBAS Ampliprep/COBAS TaqMan HBV test v2.0 for the quantification of HBV DNA in plasma samples. <i>Journal of Clinical Virology</i> , 2018, 104, 83-88.	3.1	11
24	<i>Escherichia coli</i> Sequence Type 410 Is Causing New International High-Risk Clones. <i>MSphere</i> , 2018, 3, .	2.9	183
25	Emergence of a vancomycin-variable <i>Enterococcus faecium</i> ST1421 strain containing a deletion in vanX. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 2936-2940.	3.0	25
26	Use of prophylactic <i>Saccharomyces boulardii</i> to prevent <i>Clostridium difficile</i> infection in hospitalized patients: a controlled prospective intervention study. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2018, 37, 1431-1439.	2.9	13
27	A near full-length open reading frame next generation sequencing assay for genotyping and identification of resistance-associated variants in hepatitis C virus. <i>Journal of Clinical Virology</i> , 2018, 105, 49-56.	3.1	9
28	Incidence and seasonality of respiratory syncytial virus hospitalisations in young children in Denmark, 2010 to 2015. <i>Eurosurveillance</i> , 2018, 23, .	7.0	38
29	Asymptomatic Carriers Contribute to Nosocomial <i>Clostridium difficile</i> Infection: A Cohort Study of 4508 Patients. <i>Gastroenterology</i> , 2017, 152, 1031-1041.e2.	1.3	65
30	Comparison of the Hologic Aptima HIV-1 Quant Dx Assay to the Roche COBAS Ampliprep/COBAS TaqMan HIV-1 Test v2.0 for the quantification of HIV-1 RNA in plasma samples.. <i>Journal of Clinical Virology</i> , 2017, 92, 14-19.	3.1	13
31	Draft genome sequence of a <i>Kluyvera intermedia</i> isolate from a patient with a pancreatic abscess. <i>Journal of Global Antimicrobial Resistance</i> , 2017, 10, 1-2.	2.2	13
32	Analytical and clinical performance of the Hologic Aptima HCV Quant Dx Assay for the quantification of HCV RNA in plasma samples. <i>Journal of Virological Methods</i> , 2017, 248, 159-165.	2.1	12
33	Survival and Evolution of a Large Multidrug Resistance Plasmid in New Clinical Bacterial Hosts. <i>Molecular Biology and Evolution</i> , 2016, 33, 2860-2873.	8.9	212
34	Enterogastric <i>Escherichia coli</i> in Daycare – A 1-Year Dynamic Cohort Study. <i>Frontiers in Cellular and Infection Microbiology</i> , 2016, 6, 75.	3.9	13
35	Host-Specific Patterns of Genetic Diversity among Inc11-III <sup>3</sup> and IncK Plasmids Encoding CMY-2 $\beta$ -Lactamase in <i>Escherichia coli</i> Isolates from Humans, Poultry Meat, Poultry, and Dogs in Denmark. <i>Applied and Environmental Microbiology</i> , 2016, 82, 4705-4714.	3.1	72
36	A $\beta$ -Nuclease Genotyping Assay for Identification of Macrolide-Resistant <i>Mycoplasma genitalium</i> in Clinical Specimens. <i>Journal of Clinical Microbiology</i> , 2016, 54, 1593-1597.	3.9	28

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37	Insights into Alpha-Hemolysin (Hla) Evolution and Expression among Staphylococcus aureus Clones with Hospital and Community Origin. PLoS ONE, 2014, 9, e98634.	2.5	43
38	Influence of Hepatitis C Virus and IL28B Genotypes on Liver Stiffness. PLoS ONE, 2014, 9, e115882.	2.5	6
39	Comparison of the QIAGEN artus HCV QS-RGQ test with the Roche COBAS Ampliprep/COBAS TaqMan HCV test v2.0 for the quantification of HCV-RNA in plasma samples. Journal of Clinical Virology, 2014, 60, 323-327.	3.1	9
40	An NDM-1-producing Escherichia coli obtained in Denmark has a genetic profile similar to an NDM-1-producing E. coli isolate from the UK. Journal of Antimicrobial Chemotherapy, 2012, 67, 2049-2051.	3.0	26
41	Prevalence and molecular characterization of clinical isolates of Escherichia coli expressing an AmpC phenotype. Journal of Antimicrobial Chemotherapy, 2010, 65, 460-464.	3.0	56
42	Antibody neutralization of human immunodeficiency virus type 1 (HIV-1). Acta Pathologica Microbiologica Et Immunologica Scandinavica - Supplementum, 2003, , 1-42.	0.2	2
43	The N-linked glycan of the V3 region of HIV-1 gp120 and CXCR4-dependent multiplication of a human immunodeficiency virus type 1 lymphocyte-tropic variant. FEBS Letters, 1999, 454, 47-52.	2.8	19
44	Complement-Mediated Enhancement of HIV-1 Infection in Peripheral Blood Mononuclear Cells. Scandinavian Journal of Infectious Diseases, 1997, 29, 447-452.	1.5	13
45	Gene therapy of T helper cells in HIV infection: Mathematical model of the criteria for clinical effect. Bulletin of Mathematical Biology, 1997, 59, 725-745.	1.9	25
46	Gene therapy of T helper cells in HIV infection: Mathematical model of the criteria for clinical effect. Bulletin of Mathematical Biology, 1997, 59, 725-745.	1.9	5