

# Jannik Fonager

## List of Publications by Year in descending order

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Version: 2024-02-01

40  
papers

2,173  
citations

304743

22  
h-index

302126

39  
g-index

52  
all docs

52  
docs citations

52  
times ranked

3517  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular epidemiology of the SARS-CoV-2 variant Omicron BA.2 sub-lineage in Denmark, 29 November 2021 to 2 January 2022. <i>Eurosurveillance</i> , 2022, 27, .	7.0	70
2	Risk of hospitalisation associated with infection with SARS-CoV-2 omicron variant versus delta variant in Denmark: an observational cohort study. <i>Lancet Infectious Diseases</i> , The, 2022, 22, 967-976.	9.1	140
3	Introduction and transmission of SARS-CoV-2 lineage B.1.1.7, Alpha variant, in Denmark. <i>Genome Medicine</i> , 2022, 14, 47.	8.2	14
4	Emergence of the G118R Pan-Integrase Resistance Mutation as a Result of Low Compliance to a Dolutegravir-Based cART. <i>Infectious Disease Reports</i> , 2022, 14, 501-504.	3.1	3
5	Effect of vaccination on household transmission of SARS-CoV-2 Delta variant of concern. <i>Nature Communications</i> , 2022, 13, .	12.8	28
6	Possible Involvement of Central Nervous System in COVID-19 and Sequence Variability of SARS-CoV-2 Revealed in Autopsy Tissue Samples: A Case Report. <i>BMC Clinical Pathology</i> , 2021, 14, 2632010X2110060.	1.7	4
7	Identification of risk factors associated with national transmission and late presentation of HIV-1, Denmark, 2009 to 2017. <i>Eurosurveillance</i> , 2021, 26, .	7.0	3
8	Neutralisation of the SARS-CoV-2 Delta variant sub-lineages AY.4.2 and B.1.617.2 with the mutation E484K by Comirnaty (BNT162b2 mRNA) vaccine-elicited sera, Denmark, 1 to 26 November 2021. <i>Eurosurveillance</i> , 2021, 26, .	7.0	16
9	Epidemiological characterisation of the first 785 SARS-CoV-2 Omicron variant cases in Denmark, December 2021. <i>Eurosurveillance</i> , 2021, 26, .	7.0	163
10	Increased transmissibility of SARS-CoV-2 lineage B.1.1.7 by age and viral load. <i>Nature Communications</i> , 2021, 12, 7251.	12.8	67
11	Taking gastro-surveillance into the 21st century. <i>Journal of Clinical Virology</i> , 2019, 117, 43-48.	3.1	4
12	Limited anti-HIV neutralizing antibody breadth and potency before and after HIV superinfection in Danish men who have sex with men. <i>Infectious Diseases</i> , 2019, 51, 56-61.	2.8	1
13	Piloting a surveillance system for HIV drug resistance in the European Union. <i>Eurosurveillance</i> , 2019, 24, .	7.0	10
14	Molecular epidemiology of human adenovirus infections in Denmark, 2011â€“2016. <i>Journal of Clinical Virology</i> , 2018, 104, 16-22.	3.1	23
15	Characterisation of HIV-1 transmission clusters and drug-resistant mutations in Denmark, 2004 to 2016. <i>Eurosurveillance</i> , 2018, 23, .	7.0	10
16	A universal primer-independent next-generation sequencing approach for investigations of norovirus outbreaks and novel variants. <i>Scientific Reports</i> , 2017, 7, 813.	3.3	19
17	Variant Exported Blood-Stage Proteins Encoded by Plasmodium Multigene Families Are Expressed in Liver Stages Where They Are Exported into the Parasitophorous Vacuole. <i>PLoS Pathogens</i> , 2016, 12, e1005917.	4.7	56
18	HIV-1 transmission between MSM and heterosexuals, and increasing proportions of circulating recombinant forms in the Nordic Countries. <i>Virus Evolution</i> , 2016, 2, vew010.	4.9	68

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19	Letter to the editor: A norovirus intervariant GII.4 recombinant in Victoria, Australia, June 2016: the next epidemic variant? Reflections and a note of caution. <i>Eurosurveillance</i> , 2016, 21, .	7.0	0
20	Investigation of Human Cancers for Retrovirus by Low-Stringency Target Enrichment and High-Throughput Sequencing. <i>Scientific Reports</i> , 2015, 5, 13201.	3.3	34
21	Sources of Calicivirus Contamination in Foodborne Outbreaks in Denmark, 2005â€“2011â€”The Role of the Asymptomatic Food Handler. <i>Journal of Infectious Diseases</i> , 2015, 211, 563-570.	4.0	56
22	Identification of minority resistance mutations in the HIV-1 integrase coding region using next generation sequencing. <i>Journal of Clinical Virology</i> , 2015, 73, 95-100.	3.1	15
23	Sequence Analysis of the Capsid Gene during a Genotype II.4 Dominated Norovirus Season in One University Hospital: Identification of Possible Transmission Routes. <i>PLoS ONE</i> , 2015, 10, e0115331.	2.5	14
24	Phylogenetic and epidemiological analysis of measles outbreaks in Denmark, 2013 to 2014. <i>Eurosurveillance</i> , 2015, 20, .	7.0	14
25	Mumps resurgence in Denmark. <i>Journal of Clinical Virology</i> , 2014, 61, 435-438.	3.1	12
26	Drug resistance in HIV patients with virological failure or slow virological response to antiretroviral therapy in Ethiopia. <i>BMC Infectious Diseases</i> , 2014, 14, 181.	2.9	34
27	The Application of New Molecular Methods in the Investigation of a Waterborne Outbreak of Norovirus in Denmark, 2012. <i>PLoS ONE</i> , 2014, 9, e105053.	2.5	18
28	Norovirus Epidemiology in Community and Health Care Settings and Association with Patient Age, Denmark. <i>Emerging Infectious Diseases</i> , 2014, 20, 1123-1131.	4.3	27
29	Proteomic and Genetic Analyses Demonstrate that <i>Plasmodium berghei</i> Blood Stages Export a Large and Diverse Repertoire of Proteins. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 426-448.	3.8	65
30	Reduced CD36-dependent tissue sequestration of <i>Plasmodium</i> -infected erythrocytes is detrimental to malaria parasite growth in vivo. <i>Journal of Experimental Medicine</i> , 2012, 209, 93-107.	8.5	97
31	A genotype and phenotype database of genetically modified malaria-parasites. <i>Trends in Parasitology</i> , 2011, 27, 31-39.	3.3	51
32	Development of the piggyBac transposable system for <i>Plasmodium berghei</i> and its application for random mutagenesis in malaria parasites. <i>BMC Genomics</i> , 2011, 12, 155.	2.8	30
33	Sequestration and Tissue Accumulation of Human Malaria Parasites: Can We Learn Anything from Rodent Models of Malaria?. <i>PLoS Pathogens</i> , 2010, 6, e1001032.	4.7	99
34	Rapid Changes in Transcription Profiles of the <i>Plasmodium yoelii</i> yir Multigene Family in Clonal Populations: Lack of Epigenetic Memory?. <i>PLoS ONE</i> , 2009, 4, e4285.	2.5	25
35	Visualisation and Quantitative Analysis of the Rodent Malaria Liver Stage by Real Time Imaging. <i>PLoS ONE</i> , 2009, 4, e7881.	2.5	205
36	Transcription and alternative splicing in the yir multigene family of the malaria parasite <i>Plasmodium yoelii</i> : Identification of motifs suggesting epigenetic and post-transcriptional control of RNA expression. <i>Molecular and Biochemical Parasitology</i> , 2007, 156, 1-11.	1.1	23

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37	Host immunity modulates transcriptional changes in a multigene family (yir) of rodent malaria. <i>Molecular Microbiology</i> , 2005, 58, 636-647.	2.5	41
38	Mild stress-induced stimulation of heat-shock protein synthesis and improved functional ability of human fibroblasts undergoing aging in vitro. <i>Experimental Gerontology</i> , 2002, 37, 1223-1228.	2.8	106
39	HEAT SHOCK RESPONSE AND AGEING: MECHANISMS AND APPLICATIONS. <i>Cell Biology International</i> , 2001, 25, 845-857.	3.0	199
40	Increased Risk of Hospitalisation Associated with Infection with SARS-CoV-2 Lineage B.1.1.7 in Denmark. <i>SSRN Electronic Journal</i> , 0, , .	0.4	34