

# Lorenzo Subissi

## List of Publications by Year in descending order

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Version: 2024-02-01

33  
papers

2,279  
citations

516561

16  
h-index

414303

32  
g-index

36  
all docs

36  
docs citations

36  
times ranked

4109  
citing authors

#	ARTICLE	IF	CITATIONS
1	One severe acute respiratory syndrome coronavirus protein complex integrates processive RNA polymerase and exonuclease activities. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E3900-9.	3.3	482
2	Structural and molecular basis of mismatch correction and ribavirin excision from coronavirus RNA. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E162-E171.	3.3	331
3	RNA 3'-end mismatch excision by the severe acute respiratory syndrome coronavirus nonstructural protein nsp10/nsp14 exoribonuclease complex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 9372-9377.	3.3	297
4	SARS-CoV-2 Variants of Interest and Concern naming scheme conducive for global discourse. <i>Nature Microbiology</i> , 2021, 6, 821-823.	5.9	221
5	Insights into RNA synthesis, capping, and proofreading mechanisms of SARS-coronavirus. <i>Virus Research</i> , 2014, 194, 90-99.	1.1	191
6	SARS-CoV ORF1b-encoded nonstructural proteins 12-16: Replicative enzymes as antiviral targets. <i>Antiviral Research</i> , 2014, 101, 122-130.	1.9	153
7	Evaluation of post-introduction COVID-19 vaccine effectiveness: Summary of interim guidance of the World Health Organization. <i>Vaccine</i> , 2021, 39, 4013-4024.	1.7	110
8	Ebola Virus Transmission Caused by Persistently Infected Survivors of the 2014-2016 Outbreak in West Africa. <i>Journal of Infectious Diseases</i> , 2018, 218, S287-S291.	1.9	58
9	Early epidemiological investigations: World Health Organization UNITY protocols provide a standardized and timely international investigation framework during the COVID-19 pandemic. <i>Influenza and Other Respiratory Viruses</i> , 2022, 16, 7-13.	1.5	51
10	An early warning system for emerging SARS-CoV-2 variants. <i>Nature Medicine</i> , 2022, 28, 1110-1115.	15.2	47
11	Zika Virus Infection during Pregnancy and Effects on Early Childhood Development, French Polynesia, 2013-2016. <i>Emerging Infectious Diseases</i> , 2018, 24, 1850-1858.	2.0	36
12	Epidemiology and genotype 3 subtype dynamics of hepatitis E virus in Belgium, 2010 to 2017. <i>Eurosurveillance</i> , 2019, 24, .	3.9	33
13	Chikungunya fever in Africa: a systematic review. <i>Pathogens and Global Health</i> , 2020, 114, 111-119.	1.0	29
14	Subtype-specific differences in the risk of hospitalisation among patients infected with hepatitis E virus genotype 3 in Belgium, 2010-2018. <i>Epidemiology and Infection</i> , 2019, 147, e224.	1.0	26
15	Unusual Ebola Virus Chain of Transmission, Conakry, Guinea, 2014-2015. <i>Emerging Infectious Diseases</i> , 2016, 22, 2149-2152.	2.0	21
16	Impact of infection prevention and control training on health facilities during the Ebola virus disease outbreak in Guinea. <i>BMC Public Health</i> , 2018, 18, 547.	1.2	20
17	Monitoring of human coronaviruses in Belgian primary care and hospitals, 2015-20: a surveillance study. <i>Lancet Microbe</i> , The, 2021, 2, e105-e114.	3.4	20
18	Revising rates of asymptomatic Zika virus infection based on sentinel surveillance data from French Overseas Territories. <i>International Journal of Infectious Diseases</i> , 2017, 65, 116-118.	1.5	18

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19	Evaluation of Chimerism Dynamics after Allogeneic Hematopoietic Stem Cell Transplantation in Children with Nonmalignant Diseases. <i>Biology of Blood and Marrow Transplantation</i> , 2018, 24, 1088-1093.	2.0	17
20	Subsequent mortality in survivors of Ebola virus disease in Guinea: a nationwide retrospective cohort study. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 1202-1208.	4.6	16
21	Sex practices and awareness of Ebola virus disease among male survivors and their partners in Guinea. <i>BMJ Global Health</i> , 2017, 2, e000412.	2.0	9
22	Investigation of a cross-border case of Lassa fever in West Africa. <i>BMC Infectious Diseases</i> , 2019, 19, 606.	1.3	8
23	Capturing respiratory syncytial virus season in Belgium using the influenza severe acute respiratory infection surveillance network, season 2018/19. <i>Eurosurveillance</i> , 2020, 25, .	3.9	8
24	Transmission of SARS-CoV-2 in standardised first few X cases and household transmission investigations: A systematic review and meta-analysis. <i>Influenza and Other Respiratory Viruses</i> , 2022, 16, 803-819.	1.5	6
25	Lessons learned by surveillance during the tail-end of the Ebola outbreak in Guinea, June-October 2015: a case series. <i>BMC Infectious Diseases</i> , 2017, 17, 304.	1.3	4
26	Public Health Program for Decreasing Risk for Ebola Virus Disease Resurgence from Survivors of the 2013-2016 Outbreak, Guinea. <i>Emerging Infectious Diseases</i> , 2020, 26, 206-211.	2.0	4
27	Spotlight influenza: Extending influenza surveillance to detect non-influenza respiratory viruses of public health relevance: analysis of surveillance data, Belgium, 2015 to 2019. <i>Eurosurveillance</i> , 2021, 26, .	3.9	4
28	Can Ebola virus re-emerge from survivors' body fluids other than semen?. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 933-934.	4.6	3
29	Assessment of a health facility based active case finding system for Ebola virus disease in Mbandaka, Democratic Republic of the Congo, June-July 2018. <i>BMC Infectious Diseases</i> , 2019, 19, 981.	1.3	3
30	Strong increase of true and false positive mycobacterial cultures sent to the National Reference Centre in Belgium, 2007 to 2016. <i>Eurosurveillance</i> , 2019, 24, .	3.9	3
31	<i>Plasmodium malariae</i> and <i>Plasmodium ovale</i> infections and their association with common red blood cell polymorphisms in a highly endemic area of Uganda. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2019, 113, 370-378.	0.7	2
32	A closed-handed affair: positive-strand RNA virus polymerases. <i>Future Virology</i> , 2014, 9, 769-784.	0.9	1
33	Les enzymes de la r�plication/transcription chez les coronavirus. <i>Virologie</i> , 2012, 16, 199-209.	0.1	1