

# Divya Venkatesh

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/211218/publications.pdf>

Version: 2024-02-01

14  
papers

525  
citations

1040056

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h-index

1125743

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19  
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19  
docs citations

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times ranked

704  
citing authors

#	ARTICLE	IF	CITATIONS
1	Antigenic Distance between North American Swine and Human Seasonal H3N2 Influenza A Viruses as an Indication of Zoonotic Risk to Humans. <i>Journal of Virology</i> , 2022, 96, JVI0137421.	3.4	10
2	Swine Influenza A Viruses and the Tangled Relationship with Humans. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2021, 11, a038737.	6.2	128
3	Regional Transmission and Reassortment of 2.3.4.4b Highly Pathogenic Avian Influenza (HPAI) Viruses in Bulgarian Poultry 2017/18. <i>Viruses</i> , 2020, 12, 605.	3.3	8
4	Detection of H3N8 influenza A virus with multiple mammalian-adaptive mutations in a rescued Grey seal ( <i>Halichoerus grypus</i> ) pup. <i>Virus Evolution</i> , 2020, 6, veaa016.	4.9	13
5	A30â€fAvian influenza viruses in wild birds: Virus evolution in a multi-host ecosystem. <i>Virus Evolution</i> , 2019, 5, .	4.9	0
6	Co-circulation of genetically distinct highly pathogenic avian influenza A clade 2.3.4.4 (H5N6) viruses in wild waterfowl and poultry in Europe and East Asia, 2017â€“18. <i>Virus Evolution</i> , 2019, 5, vez004.	4.9	63
7	Antigenic evolution of H3N2 influenza A viruses in swine in the United States from 2012 to 2016. <i>Influenza and Other Respiratory Viruses</i> , 2019, 13, 83-90.	3.4	29
8	Comparison of 2016â€“17 and Previous Epizootics of Highly Pathogenic Avian Influenza H5 Guangdong Lineage in Europe. <i>Emerging Infectious Diseases</i> , 2018, 24, 2270-2283.	4.3	60
9	Avian Influenza Viruses in Wild Birds: Virus Evolution in a Multihost Ecosystem. <i>Journal of Virology</i> , 2018, 92, .	3.4	83
10	Evolution of protein trafficking in kinetoplastid parasites: Complexity and pathogenesis. <i>Traffic</i> , 2018, 19, 803-812.	2.7	8
11	Evolution of the endomembrane systems of trypanosomatids: conservation and specialisation. <i>Journal of Cell Science</i> , 2017, 130, 1421-1434.	2.0	23
12	Genome of <i>Leptomonas pyrrhocoris</i> : a high-quality reference for monoxenous trypanosomatids and new insights into evolution of <i>Leishmania</i> . <i>Scientific Reports</i> , 2016, 6, 23704.	3.3	74
13	Unconventional Tâ€cell recognition of an arthritogenic epitope of proteoglycan aggrecan released from degrading cartilage. <i>Immunology</i> , 2016, 147, 389-398.	4.4	2
14	A comparative analysis of trypanosomatid SNARE proteins. <i>Parasitology International</i> , 2014, 63, 341-348.	1.3	17