Nicola S Lewis

List of Publications by Year in descending order

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#	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. Journal of Virology, 2022, 96, JVI0137421.	3.4	10
2	JMM Profile: Avian influenza: a veterinary pathogen with zoonotic potential. Journal of Medical Microbiology, 2022, 71, .	1.8	1
3	Swine Influenza A Viruses and the Tangled Relationship with Humans. Cold Spring Harbor Perspectives in Medicine, 2021, 11, a038737.	6.2	128
4	Antigenic evolution of H3N2 influenza A viruses in swine in the United States from 2012 to 2016. Influenza and Other Respiratory Viruses, 2019, 13, 83-90.	3.4	29
5	Antigenic and genetic evolution of contemporary swine H1 influenza viruses in the United States. Virology, 2018, 518, 45-54.	2.4	64
6	The type of adjuvant in whole inactivated influenza a virus vaccines impacts vaccine-associated enhanced respiratory disease. Vaccine, 2018, 36, 6103-6110.	3.8	20
7	Avian Influenza Viruses in Wild Birds: Virus Evolution in a Multihost Ecosystem. Journal of Virology, 2018, 92, .	3.4	83
8	Influenza A virus vaccines for swine. Veterinary Microbiology, 2017, 206, 35-44.	1.9	85
9	Drivers of airborne human-to-human pathogen transmission. Current Opinion in Virology, 2017, 22, 22-29.	5.4	81
10	A Phylogeny-Based Global Nomenclature System and Automated Annotation Tool for H1 Hemagglutinin Genes from Swine Influenza A Viruses. MSphere, 2016, 1, .	2.9	151
11	The Molecular Determinants of Antibody Recognition and Antigenic Drift in the H3 Hemagglutinin of Swine Influenza A Virus. Journal of Virology, 2016, 90, 8266-8280.	3.4	54
12	The global antigenic diversity of swine influenza A viruses. ELife, 2016, 5, e12217.	6.0	146
13	Global circulation patterns of seasonal influenza viruses vary with antigenic drift. Nature, 2015, 523, 217-220.	27.8	445
14	Comparative virulence of wild-type H1N1pdm09 influenza A isolates in swine. Veterinary Microbiology, 2015, 176, 40-49.	1.9	13
15	Characterization of co-circulating swine influenza A viruses in North America and the identification of a novel H1 genetic clade with antigenic significance. Virus Research, 2015, 201, 24-31.	2.2	48
16	Influenza A virus evolution and spatio-temporal dynamics in Eurasian wild birds: a phylogenetic and phylogeographical study of whole-genome sequence data. Journal of General Virology, 2015, 96, 2050-2060.	2.9	23
17	Novel Reassortant Human-Like H3N2 and H3N1 Influenza A Viruses Detected in Pigs Are Virulent and Antigenically Distinct from Swine Viruses Endemic to the United States. Journal of Virology, 2015, 89, 11213-11222.	3.4	84
18	Improving pandemic influenza risk assessment. ELife, 2014, 3, e03883.	6.0	53

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19	Substitutions near the Hemagglutinin Receptor-Binding Site Determine the Antigenic Evolution of Influenza A H3N2 Viruses in U.S. Swine. Journal of Virology, 2014, 88, 4752-4763.	3.4	86
20	Development of a surveillance scheme for equine influenza in the UK and characterisation of viruses isolated in Europe, Dubai and the USA from 2010–2012. Veterinary Microbiology, 2014, 169, 113-127.	1.9	55
21	Serological Evidence for Non-Lethal Exposures of Mongolian Wild Birds to Highly Pathogenic Avian Influenza H5N1 Virus. PLoS ONE, 2014, 9, e113569.	2.5	18
22	Substitutions Near the Receptor Binding Site Determine Major Antigenic Change During Influenza Virus Evolution. Science, 2013, 342, 976-979.	12.6	500
23	Avian Influenza Virus Surveillance in Wild Birds in Georgia: 2009–2011. PLoS ONE, 2013, 8, e58534.	2.5	42
24	Characterization of H1N1 Swine Influenza Viruses Circulating in Canadian Pigs in 2009. Journal of Virology, 2011, 85, 8667-8679.	3.4	41
25	Genetic and antigenic characterization of H1 influenza viruses from United States swine from 2008. Journal of General Virology, 2011, 92, 919-930.	2.9	123