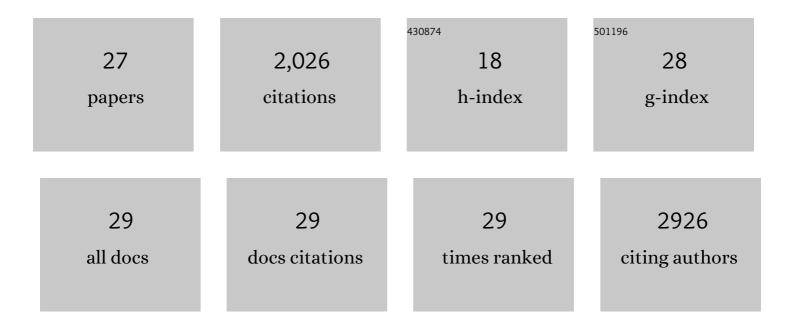
Ralf Dürrwald

List of Publications by Year in descending order

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RALE DÃ14 PRIMALD

#	Article	IF	CITATIONS
1	Taxonomy of the order Mononegavirales: update 2016. Archives of Virology, 2016, 161, 2351-2360.	2.1	407
2	Taxonomy of the order Mononegavirales: update 2019. Archives of Virology, 2019, 164, 1967-1980.	2.1	224
3	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2020, 165, 3023-3072.	2.1	184
4	Taxonomy of the order Mononegavirales: update 2017. Archives of Virology, 2017, 162, 2493-2504.	2.1	173
5	Taxonomy of the order Mononegavirales: update 2018. Archives of Virology, 2018, 163, 2283-2294.	2.1	153
6	Molecular Epidemiology and Evolution of Influenza Viruses Circulating within European Swine between 2009 and 2013. Journal of Virology, 2015, 89, 9920-9931.	3.4	148
7	European Surveillance Network for Influenza in Pigs: Surveillance Programs, Diagnostic Tools and Swine Influenza Virus Subtypes Identified in 14 European Countries from 2010 to 2013. PLoS ONE, 2014, 9, e115815.	2.5	107
8	Trends in respiratory virus circulation following COVID-19-targeted nonpharmaceutical interventions in Germany, January - September 2020: Analysis of national surveillance data. Lancet Regional Health - Europe, The, 2021, 6, 100112.	5.6	95
9	High prevalence of amantadine resistance among circulating European porcine influenza A viruses. Journal of General Virology, 2009, 90, 900-908.	2.9	77
10	Taxonomy of the order Mononegavirales: second update 2018. Archives of Virology, 2019, 164, 1233-1244.	2.1	70
11	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Archives of Virology, 2021, 166, 3513-3566.	2.1	62
12	Novel reassortant of swine influenza H1N2 virus in Germany. Journal of General Virology, 2008, 89, 271-276.	2.9	48
13	Reassortants of the pandemic (H1N1) 2009 virus and establishment of a novel porcine H1N2 influenza virus, lineage in Germany. Veterinary Microbiology, 2013, 167, 345-356.	1.9	46
14	Swine Influenza A Vaccines, Pandemic (H1N1) 2009 Virus, and Cross-Reactivity. Emerging Infectious Diseases, 2010, 16, 1029-1030.	4.3	37
15	Severe impairment of T-cell responses to BNT162b2 immunization in patients with multiple myeloma. Blood, 2022, 139, 137-142.	1.4	29
16	Prevalence of antibodies to European porcine influenza viruses in humans living in high pig density areas of Germany. Medical Microbiology and Immunology, 2014, 203, 13-24.	4.8	22
17	Origin of the European avian-like swine influenza viruses. Journal of General Virology, 2014, 95, 2372-2376.	2.9	21
18	Ongoing evolution of swine influenza viruses: a novel reassortant. Archives of Virology, 2008, 153, 2085-2092.	2.1	20

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#	Article	IF	CITATIONS
19	Displacement of the Gent/1999 human-like swine H1N2 influenza A virus lineage by novel H1N2 reassortants in Germany. Archives of Virology, 2020, 165, 55-67.	2.1	13
20	Cocirculation of Swine H1N1 Influenza A Virus Lineages in Germany. Viruses, 2020, 12, 762.	3.3	12
21	Infection Studies in Pigs and Porcine Airway Epithelial Cells Reveal an Evolution of A(H1N1)pdm09 Influenza A Viruses Toward Lower Virulence. Journal of Infectious Diseases, 2019, 219, 1596-1604.	4.0	11
22	Strengthening the Interaction of the Virology Community with the International Committee on Taxonomy of Viruses (ICTV) by Linking Virus Names and Their Abbreviations to Virus Species. Systematic Biology, 2019, 68, 828-839.	5.6	11
23	Zoonotic infection with swine A/H1avN1 influenza virus in a child, Germany, June 2020. Eurosurveillance, 2020, 25, .	7.0	11
24	Determination of respiratory syncytial virus epidemic seasons by using 95% confidence interval of positivity rates, 2011–2021, Germany. Influenza and Other Respiratory Viruses, 2022, 16, 854-857.	3.4	11
25	Novel reassortant swine H3N2 influenza A viruses in Germany. Scientific Reports, 2020, 10, 14296.	3.3	10
26	Circulation of classical swine influenza virus in Europe between the wars?. Archives of Virology, 2014, 159, 1467-1473.	2.1	9
27	Virological Surveillance and Molecular Characterization of Human Parainfluenzavirus Infection in Children with Acute Respiratory Illness: Germany, 2015–2019. Microorganisms, 2021, 9, 1508.	3.6	6