## Andres Velasco-Villa

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7917726/publications.pdf

Version: 2024-02-01

49 papers

2,141 citations

201674 27 h-index 233421 45 g-index

49 all docs 49 docs citations

times ranked

49

1976 citing authors

#	Article	IF	CITATIONS
1	Severe Acute Respiratory Syndrome Coronavirus 2 Transmission in a Georgia School District—United States, December 2020–January 2021. Clinical Infectious Diseases, 2022, 74, 319-326.	5.8	19
2	The Importance of Accurate Host Species Identification in the Framework of Rabies Surveillance, Control and Elimination. Viruses, 2022, 14, 492.	3.3	6
3	A Cross Sectional Sampling Reveals Novel Coronaviruses in Bat Populations of Georgia. Viruses, 2022, 14, 72.	3.3	3
4	Divergent Rabies Virus Variant of Probable Bat Origin in 2 Gray Foxes, New Mexico, USA. Emerging Infectious Diseases, 2022, 28, .	4.3	2
5	Clusters of SARS-CoV-2 Infection Among Elementary School Educators and Students in One School District â€" Georgia, December 2020â€"January 2021. Morbidity and Mortality Weekly Report, 2021, 70, 289-292.	15.1	68
6	Use of partial N-gene sequences as a tool to monitor progress on rabies control and elimination efforts in Ethiopia. Acta Tropica, 2021, 221, 106022.	2.0	5
7	Evaluation of rabies virus characterization to enhance early detection of important rabies epizootic events in the United States. Journal of the American Veterinary Medical Association, 2020, 256, 66-76.	0.5	19
8	Evaluation of species identification and rabies virus characterization among bat rabies cases in the United States. Journal of the American Veterinary Medical Association, 2020, 256, 77-84.	0.5	21
9	Defining New Pathways to Manage the Ongoing Emergence of Bat Rabies in Latin America. Viruses, 2020, 12, 1002.	3.3	38
10	Unexpected rabies variant identified in kinkajou ( <i>Potos flavus</i> ), Mato Grosso, Brazil. Emerging Microbes and Infections, 2020, 9, 851-854.	6.5	9
11	Clinical Presentation and Serologic Response during a Rabies Epizootic in Captive Common Vampire Bats (Desmodus rotundus). Tropical Medicine and Infectious Disease, 2020, 5, 34.	2.3	17
12	Abortive vampire bat rabies infections in Peruvian peridomestic livestock. PLoS Neglected Tropical Diseases, 2020, 14, e0008194.	3.0	18
13	Rabies surveillance in the United States during 2015. Journal of the American Veterinary Medical Association, 2017, 250, 1117-1130.	0.5	55
14	The history of rabies in the Western Hemisphere. Antiviral Research, 2017, 146, 221-232.	4.1	77
15	Successful strategies implemented towards the elimination of canine rabies in the Western Hemisphere. Antiviral Research, 2017, 143, 1-12.	4.1	94
16	A Pan-Lyssavirus Taqman Real-Time RT-PCR Assay for the Detection of Highly Variable Rabies virus and Other Lyssaviruses. PLoS Neglected Tropical Diseases, 2017, 11, e0005258.	3.0	58
17	An inter- laboratory proficiency testing exercise for rabies diagnosis in Latin America and the Caribbean. PLoS Neglected Tropical Diseases, 2017, 11, e0005427.	3.0	13
18	Infectivity of attenuated poxvirus vaccine vectors and immunogenicity of a raccoonpox vectored rabies vaccine in the Brazilian Free-tailed bat (Tadarida brasiliensis). Vaccine, 2016, 34, 5352-5358.	3.8	27

#	Article	IF	CITATIONS
19	Host–pathogen evolutionary signatures reveal dynamics and future invasions of vampire bat rabies. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10926-10931.	7.1	108
20	Virology, Immunology and Pathology of Human Rabies During Treatment. Pediatric Infectious Disease Journal, 2015, 34, 520-528.	2.0	23
21	Phylodynamics of vampire batâ€transmitted rabies in <scp>A</scp> rgentina. Molecular Ecology, 2014, 23, 2340-2352.	3.9	27
22	Rabies death attributed to exposure in Central America with symptom onset in a U.S. detention facility - Texas, 2013. Morbidity and Mortality Weekly Report, 2014, 63, 446-9.	15.1	6
23	Raccoon Rabies Virus Variant Transmission Through Solid Organ Transplantation. JAMA - Journal of the American Medical Association, 2013, 310, 398.	7.4	107
24	Metabolomics of Cerebrospinal Fluid from Humans Treated for Rabies. Journal of Proteome Research, 2013, 12, 481-490.	3.7	48
25	Enzootic and Epizootic Rabies Associated with Vampire Bats, Peru. Emerging Infectious Diseases, 2013, 19, 1463-69.	4.3	48
26	Evolutionary History and Phylogeography of Rabies Viruses Associated with Outbreaks in Trinidad. PLoS Neglected Tropical Diseases, 2013, 7, e2365.	3.0	24
27	Overview of Johne's disease immunology. Veterinary World, 2013, 6, 901-904.	1.7	2
28	Molecular Inferences Suggest Multiple Host Shifts of Rabies Viruses from Bats to Mesocarnivores in Arizona during 2001–2009. PLoS Pathogens, 2012, 8, e1002786.	4.7	160
29	Rates of Viral Evolution Are Linked to Host Geography in Bat Rabies. PLoS Pathogens, 2012, 8, e1002720.	4.7	79
30	Variable evolutionary routes to host establishment across repeated rabies virus host shifts among bats. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 19715-19720.	7.1	70
31	Evaluation of a Direct, Rapid Immunohistochemical Test for Rabies Diagnosis. Emerging Infectious Diseases, 2012, 12, 310-313.	4.3	162
32	Immune response after rabies vaccine in a kidney transplant recipient. Transplant Infectious Disease, 2011, 13, 492-495.	1.7	16
33	Detection of North American orthopoxviruses by real time-PCR. Virology Journal, 2011, 8, 313.	3.4	4
34	New rabies virus variant found during an epizootic in white-nosed coatis from the Yucatan Peninsula. Epidemiology and Infection, 2010, 138, 1586-1589.	2.1	32
35	Ferret badger rabies origin and its revisited importance as potential source of rabies transmission in Southeast China. BMC Infectious Diseases, 2010, 10, 234.	2.9	38
36	Human Rabies and Rabies in Vampire and Nonvampire Bat Species, Southeastern Peru, 2007. Emerging Infectious Diseases, 2009, 15, 1308-1310.	4.3	43

#	Article	IF	CITATIONS
37	Molecular Epidemiology of Rabies in Southern People's Republic of China. Emerging Infectious Diseases, 2009, 15, 1192-1198.	4.3	54
38	Rabies virus pathogenesis in relationship to intervention with inactivated and attenuated rabies vaccines. Vaccine, 2009, 27, 7149-7155.	3.8	35
39	Identification of New Rabies Virus Variant in Mexican Immigrant. Emerging Infectious Diseases, 2008, 14, 1906-1908.	4.3	28
40	Enzootic Rabies Elimination from Dogs and Reemergence in Wild Terrestrial Carnivores, United States. Emerging Infectious Diseases, 2008, 14, 1849-1854.	4.3	126
41	Molecular epidemiology identifies only a single rabies virus variant circulating in complex carnivore communities of the Serengeti. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 2123-2130.	2.6	45
42	Molecular epidemiology of rabies in Colombia 1994–2005 based on partial nucleoprotein gene sequences. Virus Research, 2007, 130, 172-181.	2.2	20
43	Are all lyssavirus genes equal for phylogenetic analyses?. Virus Research, 2007, 129, 91-103.	2.2	35
44	A new phylogenetic lineage of Rabies virus associated with western pipistrelle bats (Pipistrellus) Tj ETQq0 0 0 rgB	T <u>l</u> Overloc	k 10 Tf 50 46
45	Molecular Diversity of Rabies Viruses Associated with Bats in Mexico and Other Countries of the Americas. Journal of Clinical Microbiology, 2006, 44, 1697-1710.	3.9	87
46	Molecular epizootiology of rabies associated with terrestrial carnivores in Mexico. Virus Research, 2005, 111, 13-27.	2.2	55
47	Antigenic Diversity and Distribution of Rabies Virus in Mexico. Journal of Clinical Microbiology, 2002, 40, 951-958.	3.9	34
48	Infectious Diseases in Mexico. A Survey from 1995–2000. Archives of Medical Research, 2002, 33, 343-350.	3.3	32
49	VP4 and VP7 Genotyping by Reverse Transcription-PCR of Human Rotavirus in Mexican Children with Acute Diarrhea. Journal of Clinical Microbiology, 2000, 38, 3876-3878.	3.9	16