

Andres Velasco-Villa

List of Publications by Year in descending order

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Version: 2024-02-01

49
papers

2,141
citations

201658

27
h-index

233409

45
g-index

49
all docs

49
docs citations

49
times ranked

1976
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluation of a Direct, Rapid Immunohistochemical Test for Rabies Diagnosis. <i>Emerging Infectious Diseases</i> , 2012, 12, 310-313.	4.3	162
2	Molecular Inferences Suggest Multiple Host Shifts of Rabies Viruses from Bats to Mesocarnivores in Arizona during 2001â€“2009. <i>PLoS Pathogens</i> , 2012, 8, e1002786.	4.7	160
3	Enzootic Rabies Elimination from Dogs and Reemergence in Wild Terrestrial Carnivores, United States. <i>Emerging Infectious Diseases</i> , 2008, 14, 1849-1854.	4.3	126
4	Hostâ€“pathogen evolutionary signatures reveal dynamics and future invasions of vampire bat rabies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 10926-10931.	7.1	108
5	Raccoon Rabies Virus Variant Transmission Through Solid Organ Transplantation. <i>JAMA - Journal of the American Medical Association</i> , 2013, 310, 398.	7.4	107
6	Successful strategies implemented towards the elimination of canine rabies in the Western Hemisphere. <i>Antiviral Research</i> , 2017, 143, 1-12.	4.1	94
7	Molecular Diversity of Rabies Viruses Associated with Bats in Mexico and Other Countries of the Americas. <i>Journal of Clinical Microbiology</i> , 2006, 44, 1697-1710.	3.9	87
8	Rates of Viral Evolution Are Linked to Host Geography in Bat Rabies. <i>PLoS Pathogens</i> , 2012, 8, e1002720.	4.7	79
9	The history of rabies in the Western Hemisphere. <i>Antiviral Research</i> , 2017, 146, 221-232.	4.1	77
10	Variable evolutionary routes to host establishment across repeated rabies virus host shifts among bats. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 19715-19720.	7.1	70
11	Clusters of SARS-CoV-2 Infection Among Elementary School Educators and Students in One School District â€” Georgia, December 2020â€“January 2021. <i>Morbidity and Mortality Weekly Report</i> , 2021, 70, 289-292.	15.1	68
12	A Pan-Lyssavirus Taqman Real-Time RT-PCR Assay for the Detection of Highly Variable Rabies virus and Other Lyssaviruses. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005258.	3.0	58
13	Molecular epizootiology of rabies associated with terrestrial carnivores in Mexico. <i>Virus Research</i> , 2005, 111, 13-27.	2.2	55
14	Rabies surveillance in the United States during 2015. <i>Journal of the American Veterinary Medical Association</i> , 2017, 250, 1117-1130.	0.5	55
15	Molecular Epidemiology of Rabies in Southern Peopleâ€™s Republic of China. <i>Emerging Infectious Diseases</i> , 2009, 15, 1192-1198.	4.3	54
16	Metabolomics of Cerebrospinal Fluid from Humans Treated for Rabies. <i>Journal of Proteome Research</i> , 2013, 12, 481-490.	3.7	48
17	Enzootic and Epizootic Rabies Associated with Vampire Bats, Peru. <i>Emerging Infectious Diseases</i> , 2013, 19, 1463-69.	4.3	48
18	Molecular epidemiology identifies only a single rabies virus variant circulating in complex carnivore communities of the Serengeti. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2007, 274, 2123-2130.	2.6	45

#	ARTICLE	IF	CITATIONS
19	Human Rabies and Rabies in Vampire and Nonvampire Bat Species, Southeastern Peru, 2007. <i>Emerging Infectious Diseases</i> , 2009, 15, 1308-1310.	4.3	43
20	Ferret badger rabies origin and its revisited importance as potential source of rabies transmission in Southeast China. <i>BMC Infectious Diseases</i> , 2010, 10, 234.	2.9	38
21	Defining New Pathways to Manage the Ongoing Emergence of Bat Rabies in Latin America. <i>Viruses</i> , 2020, 12, 1002.	3.3	38
22	Are all lyssavirus genes equal for phylogenetic analyses?. <i>Virus Research</i> , 2007, 129, 91-103.	2.2	35
23	Rabies virus pathogenesis in relationship to intervention with inactivated and attenuated rabies vaccines. <i>Vaccine</i> , 2009, 27, 7149-7155.	3.8	35
24	Antigenic Diversity and Distribution of Rabies Virus in Mexico. <i>Journal of Clinical Microbiology</i> , 2002, 40, 951-958.	3.9	34
25	Infectious Diseases in Mexico. A Survey from 1995â€“2000. <i>Archives of Medical Research</i> , 2002, 33, 343-350.	3.3	32
26	New rabies virus variant found during an epizootic in white-nosed coatis from the Yucatan Peninsula. <i>Epidemiology and Infection</i> , 2010, 138, 1586-1589.	2.1	32
27	A new phylogenetic lineage of Rabies virus associated with western pipistrelle bats (<i>Pipistrellus</i>) Tj ETQq1 1 0.784314 rgBT / Overlock	2.9	28
28	Identification of New Rabies Virus Variant in Mexican Immigrant. <i>Emerging Infectious Diseases</i> , 2008, 14, 1906-1908.	4.3	28
29	Phylodynamics of vampire batâ€“transmitted rabies in Argentina. <i>Molecular Ecology</i> , 2014, 23, 2340-2352.	3.9	27
30	Infectivity of attenuated poxvirus vaccine vectors and immunogenicity of a raccoonpox vectored rabies vaccine in the Brazilian Free-tailed bat (<i>Tadarida brasiliensis</i>). <i>Vaccine</i> , 2016, 34, 5352-5358.	3.8	27
31	Evolutionary History and Phylogeography of Rabies Viruses Associated with Outbreaks in Trinidad. <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2365.	3.0	24
32	Virology, Immunology and Pathology of Human Rabies During Treatment. <i>Pediatric Infectious Disease Journal</i> , 2015, 34, 520-528.	2.0	23
33	Evaluation of species identification and rabies virus characterization among bat rabies cases in the United States. <i>Journal of the American Veterinary Medical Association</i> , 2020, 256, 77-84.	0.5	21
34	Molecular epidemiology of rabies in Colombia 1994â€“2005 based on partial nucleoprotein gene sequences. <i>Virus Research</i> , 2007, 130, 172-181.	2.2	20
35	Evaluation of rabies virus characterization to enhance early detection of important rabies epizootic events in the United States. <i>Journal of the American Veterinary Medical Association</i> , 2020, 256, 66-76.	0.5	19
36	Severe Acute Respiratory Syndrome Coronavirus 2 Transmission in a Georgia School Districtâ€”United States, December 2020â€“January 2021. <i>Clinical Infectious Diseases</i> , 2022, 74, 319-326.	5.8	19

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37	Abortive vampire bat rabies infections in Peruvian peridomestic livestock. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008194.	3.0	18
38	Clinical Presentation and Serologic Response during a Rabies Epizootic in Captive Common Vampire Bats (<i>Desmodus rotundus</i>). <i>Tropical Medicine and Infectious Disease</i> , 2020, 5, 34.	2.3	17
39	Immune response after rabies vaccine in a kidney transplant recipient. <i>Transplant Infectious Disease</i> , 2011, 13, 492-495.	1.7	16
40	VP4 and VP7 Genotyping by Reverse Transcription-PCR of Human Rotavirus in Mexican Children with Acute Diarrhea. <i>Journal of Clinical Microbiology</i> , 2000, 38, 3876-3878.	3.9	16
41	An inter-laboratory proficiency testing exercise for rabies diagnosis in Latin America and the Caribbean. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005427.	3.0	13
42	Unexpected rabies variant identified in kinkajou (<i>Proteles flavus</i>), Mato Grosso, Brazil. <i>Emerging Microbes and Infections</i> , 2020, 9, 851-854.	6.5	9
43	Rabies death attributed to exposure in Central America with symptom onset in a U.S. detention facility - Texas, 2013. <i>Morbidity and Mortality Weekly Report</i> , 2014, 63, 446-9.	15.1	6
44	The Importance of Accurate Host Species Identification in the Framework of Rabies Surveillance, Control and Elimination. <i>Viruses</i> , 2022, 14, 492.	3.3	6
45	Use of partial N-gene sequences as a tool to monitor progress on rabies control and elimination efforts in Ethiopia. <i>Acta Tropica</i> , 2021, 221, 106022.	2.0	5
46	Detection of North American orthopoxviruses by real time-PCR. <i>Virology Journal</i> , 2011, 8, 313.	3.4	4
47	A Cross Sectional Sampling Reveals Novel Coronaviruses in Bat Populations of Georgia. <i>Viruses</i> , 2022, 14, 72.	3.3	3
48	Overview of Johne's disease immunology. <i>Veterinary World</i> , 2013, 6, 901-904.	1.7	2
49	Divergent Rabies Virus Variant of Probable Bat Origin in 2 Gray Foxes, New Mexico, USA. <i>Emerging Infectious Diseases</i> , 2022, 28, .	4.3	2