

# Rafael Ojeda-Flores

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5804338/publications.pdf>

Version: 2024-02-01

19  
papers

1,182  
citations

1163117

8  
h-index

839539

18  
g-index

19  
all docs

19  
docs citations

19  
times ranked

2473  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular detection of <i>Rickettsia amblyommatis</i> and <i>Rickettsia parkeri</i> in ticks collected from wild pigs in Campeche, Mexico. <i>Ticks and Tick-borne Diseases</i> , 2022, 13, 101844.	2.7	6
2	Effects of landscape anthropization on sylvatic mosquito assemblages in a rainforest in Chiapas, Mexico. <i>Acta Tropica</i> , 2021, 216, 105849.	2.0	8
3	Global subtype diversity, spatial distribution patterns, and phylogenetic analysis of avian influenza virus in water. <i>Transboundary and Emerging Diseases</i> , 2021, , .	3.0	3
4	Molecular identification and phylogenetic characterization of influenza A virus at a wildlife–livestock interface in Mexico. <i>Transboundary and Emerging Diseases</i> , 2021, 68, 3563-3573.	3.0	3
5	Phylogenetic characterization of a reassortant H5N2 influenza A virus from a resident Mexican duck ( <i>Anas diazi</i> ). <i>Infection, Genetics and Evolution</i> , 2020, 84, 104475.	2.3	2
6	Assemblage variation of mosquitoes (Diptera: Culicidae) in different land use and activity periods within a lowland tropical forest matrix in Campeche, Mexico. <i>Journal of Vector Ecology</i> , 2020, 45, 188-196.	1.0	5
7	Eco-Epidemiological Evidence of the Transmission of Avian and Human Influenza A Viruses in Wild Pigs in Campeche, Mexico. <i>Viruses</i> , 2020, 12, 528.	3.3	6
8	Simulation modeling of influenza transmission through backyard pig trade networks in a wildlife/livestock interface area. <i>Tropical Animal Health and Production</i> , 2019, 51, 2019-2024.	1.4	3
9	Composición de comunidades y fiabilidad de ectoparásitos de murciélagos en paisajes agropecuarios de Veracruz, México. <i>Ecosistemas Y Recursos Agropecuarios</i> , 2019, 7, .	0.2	1
10	Potential Sympatric Vectors and Mammalian Hosts of Venezuelan Equine Encephalitis Virus in Southern Mexico. <i>Journal of Wildlife Diseases</i> , 2017, 53, 657.	0.8	16
11	One Health proof of concept: Bringing a transdisciplinary approach to surveillance for zoonotic viruses at the human-wild animal interface. <i>Preventive Veterinary Medicine</i> , 2017, 137, 112-118.	1.9	112
12	Global patterns in coronavirus diversity. <i>Virus Evolution</i> , 2017, 3, vex012.	4.9	310
13	Viral diversity of bat communities in human-dominated landscapes in Mexico. <i>Veterinaria México OA</i> , 2015, 2, .	0.2	7
14	Non-random patterns in viral diversity. <i>Nature Communications</i> , 2015, 6, 8147.	12.8	65
15	Dengue Virus in Bats from Southeastern Mexico. <i>American Journal of Tropical Medicine and Hygiene</i> , 2014, 91, 129-131.	1.4	40
16	A Strategy To Estimate Unknown Viral Diversity in Mammals. <i>MBio</i> , 2013, 4, e00598-13.	4.1	320
17	Bats are a major natural reservoir for hepaciviruses and pegiviruses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8194-8199.	7.1	251
18	Quantitative evaluation of MPTP-treated nonhuman parkinsonian primates in the HALLWAY task. <i>Journal of Neuroscience Methods</i> , 2009, 177, 361-368.	2.5	24

#	ARTICLE	IF	CITATIONS
19	A transdisciplinary approach to disease ecology: Emerging coronaviruses. <i>Veterinaria MÃ©xico OA</i> , 0, 8, .	0.2	0