

Bradley J Blitvich

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/7112732/bradley-j-blitvich-publications-by-year.pdf>

Version: 2024-04-19

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

78
papers

2,227
citations

25
h-index

45
g-index

82
ext. papers

2,538
ext. citations

4.2
avg, IF

4.82
L-index

#	Paper	IF	Citations
78	Evidence of Coinfections between SARS-CoV-2 and Select Arboviruses in Guerrero, Mexico, 2020-2021.. <i>American Journal of Tropical Medicine and Hygiene</i> , 2022 ,	3.2	2
77	Detection of Antibodies to Lokern, Main Drain, St. Louis Encephalitis, and West Nile Viruses in Vertebrate Animals in Chihuahua, Guerrero, and Michoacán, Mexico. <i>Vector-Borne and Zoonotic Diseases</i> , 2021 , 21, 884-891	2.4	2
76	Infection, dissemination, and transmission efficiencies of Zika virus in <i>Aedes aegypti</i> after serial passage in mosquito or mammalian cell lines or alternating passage in both cell types. <i>Parasites and Vectors</i> , 2021 , 14, 261	4	2
75	Co-Circulation of All Four Dengue Viruses and Zika Virus in Guerrero, Mexico, 2019. <i>Vector-Borne and Zoonotic Diseases</i> , 2021 , 21, 458-465	2.4	4
74	Chimeric Zika viruses containing structural protein genes of insect-specific flaviviruses cannot replicate in vertebrate cells due to entry and post-translational restrictions. <i>Virology</i> , 2021 , 559, 30-39	3.6	2
73	Chikungunya in Guerrero, Mexico, 2019 and Evidence of Gross Underreporting in the Region. <i>American Journal of Tropical Medicine and Hygiene</i> , 2021 ,	3.2	1
72	Sexual Transmission of Arboviruses: A Systematic Review. <i>Viruses</i> , 2020 , 12,	6.2	6
71	Entomological and virological surveillance for dengue virus in churches in Merida, Mexico. <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2019 , 61, e9	2.2	3
70	Infection and transmission of Cache Valley virus by <i>Aedes albopictus</i> and <i>Aedes aegypti</i> mosquitoes. <i>Parasites and Vectors</i> , 2019 , 12, 384	4	7
69	Skunk River virus, a novel orbivirus isolated from <i>Aedes trivittatus</i> in the United States. <i>Journal of General Virology</i> , 2019 , 100, 295-300	4.9	4
68	Discovery of a novel Tymoviridae-like virus in mosquitoes from Mexico. <i>Archives of Virology</i> , 2019 , 164, 649-652	2.6	4
67	Complete nucleotide sequences of the large RNA genome segments of Main Drain and Northway viruses (Family Peribunyaviridae). <i>Archives of Virology</i> , 2018 , 163, 2253-2255	2.6	
66	Molecular detection of in dogs and mosquitoes in Tabasco, Mexico. <i>Journal of Vector Borne Diseases</i> , 2018 , 55, 151-158	0.7	7
65	Bunyavirus Taxonomy: Limitations and Misconceptions Associated with the Current ICTV Criteria Used for Species Demarcation. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018 , 99, 11-16	3.2	14
64	Arbovirus Surveillance near the Mexico-U.S. Border: Isolation and Sequence Analysis of Chikungunya Virus from Patients with Dengue-like Symptoms in Reynosa, Tamaulipas. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018 , 99, 191-194	3.2	7
63	Surveillance for Flaviviruses Near the Mexico-U.S. Border: Co-circulation of Dengue Virus Serotypes 1, 2, and 3 and West Nile Virus in Tamaulipas, Northern Mexico, 2014-2016. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018 , 99, 1308-1317	3.2	5
62	Detection of novel and recognized RNA viruses in mosquitoes from the Yucatan Peninsula of Mexico using metagenomics and characterization of their in vitro host ranges. <i>Journal of General Virology</i> , 2018 , 99, 1729-1738	4.9	14

61	Complete Genome Sequence of Houston Virus, a Newly Discovered Mosquito-Specific Virus Isolated from <i>Culex quinquefasciatus</i> in Mexico. <i>Microbiology Resource Announcements</i> , 2018 , 7,	1.3	2
60	Restriction of Zika virus infection and transmission in <i>Aedes aegypti</i> mediated by an insect-specific flavivirus. <i>Emerging Microbes and Infections</i> , 2018 , 7, 181	18.9	35
59	<i>Culex tarsalis</i> is a competent vector species for Cache Valley virus. <i>Parasites and Vectors</i> , 2018 , 11, 519	4	8
58	Evidence that Lokern virus (family Peribunyaviridae) is a reassortant that acquired its small and large genome segments from Main Drain virus and its medium genome segment from an undiscovered virus. <i>Virology Journal</i> , 2018 , 15, 122	6.1	4
57	Maternal, Fetal, and Neonatal Outcomes in Pregnant Dengue Patients in Mexico. <i>BioMed Research International</i> , 2018 , 2018, 9643083	3	11
56	A Review of Flaviviruses that Have No Known Arthropod Vector. <i>Viruses</i> , 2017 , 9,	6.2	39
55	Complete genome sequences of two insect-specific flaviviruses. <i>Archives of Virology</i> , 2017 , 162, 3913-3917	1.6	3
54	Complete genome sequence of TWHo virus, a novel putative flavivirus from the Yucatan Peninsula of Mexico. <i>Virology Journal</i> , 2017 , 14, 110	6.1	2
53	Hematologic RIs for healthy water buffaloes (<i>Bubalus bubalis</i>) in southern Mexico. <i>Veterinary Clinical Pathology</i> , 2017 , 46, 436-441	1	1
52	Arboviruses: Molecular Biology, Evolution and Control. Nikos Vasilakis and Duane J. Gubler. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016 , 95, 488-489	3.2	3
51	Serologic Evidence of Flavivirus Infections in Peridomestic Rodents in Merida, Mexico. <i>Journal of Wildlife Diseases</i> , 2016 , 52, 168-72	1.3	9
50	Merida virus, a putative novel rhabdovirus discovered in <i>Culex</i> and <i>Ochlerotatus</i> spp. mosquitoes in the Yucatan Peninsula of Mexico. <i>Journal of General Virology</i> , 2016 , 97, 977-987	4.9	20
49	Chikungunya Virus in Febrile Humans and <i>Aedes aegypti</i> Mosquitoes, Yucatan, Mexico. <i>Emerging Infectious Diseases</i> , 2016 , 22, 1804-7	10.2	17
48	West Nile Virus Infection in Human and Mouse Cornea Tissue. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016 , 95, 1185-1191	3.2	3
47	Insect-specific flaviviruses: a systematic review of their discovery, host range, mode of transmission, superinfection exclusion potential and genomic organization. <i>Viruses</i> , 2015 , 7, 1927-59	6.2	188
46	Characterization of newly revealed sequences in the infectious myonecrosis virus genome in <i>Litopenaeus vannamei</i> . <i>Journal of General Virology</i> , 2015 , 96, 1821-9	4.9	6
45	Management Factors Associated with Operation-Level Prevalence of Antibodies to Cache Valley Virus and Other Bunyamwera Serogroup Viruses in Sheep in the United States. <i>Vector-Borne and Zoonotic Diseases</i> , 2015 , 15, 683-93	2.4	15
44	Evidence for West Nile virus spillover into the squirrel population in Atlanta, Georgia. <i>Vector-Borne and Zoonotic Diseases</i> , 2015 , 15, 303-10	2.4	7

43	Substitution of the premembrane and envelope protein genes of Modoc virus with the homologous sequences of West Nile virus generates a chimeric virus that replicates in vertebrate but not mosquito cells. <i>Virology Journal</i> , 2014 , 11, 150	6.1	9
42	Detection of hand, foot and mouth disease in the yucatan peninsula of Mexico. <i>Gastroenterology Insights</i> , 2014 , 6, 5627	2.1	4
41	Monitoring sheep and Culicoides midges in Montana for evidence of Bunyamwera serogroup virus infection. <i>Veterinary Record Open</i> , 2014 , 1, e000071	1.4	7
40	Antibodies to West Nile virus in wild and farmed crocodiles in southeastern Mexico. <i>Journal of Wildlife Diseases</i> , 2013 , 49, 690-3	1.3	13
39	A new insect-specific flavivirus from northern Australia suppresses replication of West Nile virus and Murray Valley encephalitis virus in co-infected mosquito cells. <i>PLoS ONE</i> , 2013 , 8, e56534	3.7	14 ⁰
38	Orthobunyaviruses, a common cause of infection of livestock in the Yucatan peninsula of Mexico. <i>American Journal of Tropical Medicine and Hygiene</i> , 2012 , 87, 1132-9	3.2	16
37	Orthobunyavirus antibodies in humans, Yucatan Peninsula, Mexico. <i>Emerging Infectious Diseases</i> , 2012 , 18, 1629-32	10.2	11
36	Nucleotide sequencing and serologic analysis of Cache Valley virus isolates from the Yucatan Peninsula of Mexico. <i>Virus Genes</i> , 2012 , 45, 176-80	2.3	9
35	Sequence and phylogenetic data indicate that an orthobunyavirus recently detected in the Yucatan Peninsula of Mexico is a novel reassortant of Potosi and Cache Valley viruses. <i>Archives of Virology</i> , 2012 , 157, 1199-204	2.6	14
34	Identification of a novel subtype of South River virus (family Bunyaviridae). <i>Archives of Virology</i> , 2012 , 157, 1205-9	2.6	6
33	Serological evidence of flaviviruses and alphaviruses in livestock and wildlife in Trinidad. <i>Vector-Borne and Zoonotic Diseases</i> , 2012 , 12, 969-78	2.4	22
32	Evidence of efficient transovarial transmission of Culex flavivirus by Culex pipiens (Diptera: Culicidae). <i>Journal of Medical Entomology</i> , 2011 , 48, 1031-8	2.2	77
31	Detection of antibodies to West Nile virus in horses, Costa Rica, 2004. <i>Vector-Borne and Zoonotic Diseases</i> , 2011 , 11, 1081-4	2.4	13
30	Serologic surveillance for West Nile virus and other flaviviruses in febrile patients, encephalitic patients, and asymptomatic blood donors in northern Mexico. <i>Vector-Borne and Zoonotic Diseases</i> , 2010 , 10, 151-7	2.4	21
29	NS1Vof flaviviruses in the Japanese encephalitis virus serogroup is a product of ribosomal frameshifting and plays a role in viral neuroinvasiveness. <i>Journal of Virology</i> , 2010 , 84, 1641-7	6.6	134
28	Detection of flaviviruses and orthobunyaviruses in mosquitoes in the Yucatan Peninsula of Mexico in 2008. <i>Vector-Borne and Zoonotic Diseases</i> , 2010 , 10, 777-83	2.4	46
27	Isolation and sequence analysis of Culex flavivirus from Culex interrogator and Culex quinquefasciatus in the Yucatan Peninsula of Mexico. <i>Archives of Virology</i> , 2010 , 155, 983-6	2.6	22
26	Evidence for ribosomal frameshifting and a novel overlapping gene in the genomes of insect-specific flaviviruses. <i>Virology</i> , 2010 , 399, 153-166	3.6	54

25	Antibodies to West Nile virus in raccoons and other wild peridomestic mammals in Iowa. <i>Journal of Wildlife Diseases</i> , 2009 , 45, 1163-8	1.3	14
24	Genomic sequence and phylogenetic analysis of Culex flavivirus, an insect-specific flavivirus, isolated from Culex pipiens (Diptera: Culicidae) in Iowa. <i>Journal of Medical Entomology</i> , 2009 , 46, 934-41	2.2	53
23	Arrival and establishment of Aedes japonicus japonicus (Diptera: Culicidae) in Iowa. <i>Journal of Medical Entomology</i> , 2009 , 46, 1282-9	2.2	12
22	Complete nucleotide sequences of the small and medium RNA genome segments of Kairi virus (family Bunyaviridae). <i>Archives of Virology</i> , 2009 , 154, 1555-8	2.6	5
21	Detection of RNA from a novel West Nile-like virus and high prevalence of an insect-specific flavivirus in mosquitoes in the Yucatan Peninsula of Mexico. <i>American Journal of Tropical Medicine and Hygiene</i> , 2009 , 80, 85-95	3.2	74
20	Detection of RNA from a Novel West Nile-like Virus and High Prevalence of an Insect-specific Flavivirus in Mosquitoes in the Yucatan Peninsula of Mexico. <i>American Journal of Tropical Medicine and Hygiene</i> , 2009 , 80, 85-95	3.2	96
19	Transmission dynamics and changing epidemiology of West Nile virus. <i>Animal Health Research Reviews</i> , 2008 , 9, 71-86	2.1	100
18	West Nile virus viremia in eastern chipmunks (<i>Tamias striatus</i>) sufficient for infecting different mosquitoes. <i>Emerging Infectious Diseases</i> , 2007 , 13, 831-7	10.2	49
17	West Nile virus in horses, Guatemala. <i>Emerging Infectious Diseases</i> , 2006 , 12, 1038-9	10.2	32
16	ANTIBODIES TO WEST NILE VIRUS IN ASYMPTOMATIC MAMMALS, BIRDS, AND REPTILES IN THE YUCATAN PENINSULA OF MEXICO. <i>American Journal of Tropical Medicine and Hygiene</i> , 2006 , 74, 908-914	3.2	42
15	West Nile Virus isolation in human and mosquitoes, Mexico. <i>Emerging Infectious Diseases</i> , 2005 , 11, 1449-52	10.2	47
14	Persistence of antibodies to West Nile virus in naturally infected rock pigeons (<i>Columba livia</i>). <i>Vaccine Journal</i> , 2005 , 12, 665-7		71
13	Longitudinal studies of West Nile virus infection in avians, Yucatán State, México. <i>Vector-Borne and Zoonotic Diseases</i> , 2004 , 4, 3-14	2.4	34
12	Detection of antibodies to West Nile and Saint Louis encephalitis viruses in horses. <i>Salud Publica De Mexico</i> , 2004 , 46, 373-5	1.7	13
11	Phylogenetic analysis of West Nile virus, Nuevo Leon State, Mexico. <i>Emerging Infectious Diseases</i> , 2004 , 10, 1314-7	10.2	23
10	Epitope-blocking enzyme-linked immunosorbent assays for detection of west nile virus antibodies in domestic mammals. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 2676-9	9.7	71
9	Serologic evidence of West Nile virus infection in horses, Coahuila State, Mexico. <i>Emerging Infectious Diseases</i> , 2003 , 9, 853-6	10.2	82
8	Serologic evidence of West Nile virus infection in horses, Yucatan State, Mexico. <i>Emerging Infectious Diseases</i> , 2003 , 9, 857-9	10.2	53

7	West Nile virus transmission in resident birds, Dominican Republic. <i>Emerging Infectious Diseases</i> , 2003 , 9, 1299-302	10.2	86
6	Serologic evidence of West Nile Virus infection in birds, Tamaulipas State, Mico. <i>Vector-Borne and Zoonotic Diseases</i> , 2003 , 3, 209-13	2.4	26
5	Epitope-blocking enzyme-linked immunosorbent assays for the detection of serum antibodies to west nile virus in multiple avian species. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 1041-7	9.7	115
4	Complete cDNA and deduced amino acid sequence of the chaperonin containing T-complex polypeptide 1 (CCT) delta subunit from <i>Aedes triseriatus</i> mosquitoes. <i>DNA Sequence</i> , 2001 , 12, 203-8		3
3	Identification and sequence determination of mRNAs detected in dormant (diapausing) <i>Aedes triseriatus</i> mosquito embryos. <i>DNA Sequence</i> , 2001 , 12, 197-202		20
2	Molecular cloning and complete cDNA sequences of the ribosomal proteins rpl34 and rpl44 from <i>Aedes triseriatus</i> mosquitoes. <i>DNA Sequence</i> , 2000 , 11, 451-5		
1	Identification and analysis of truncated and elongated species of the flavivirus NS1 protein. <i>Virus Research</i> , 1999 , 60, 67-79	6.4	29