

Houssam Attoui

List of Publications by Year in Descending Order

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

101
papers

5,343
citations

37
h-index

71
g-index

107
ext. papers

5,968
ext. citations

5.6
avg, IF

4.83
L-index

#	Paper	IF	Citations
101	Evaluation of two artificial infection methods of live ticks as tools for studying interactions between tick-borne viruses and their tick vectors.. <i>Scientific Reports</i> , 2022 , 12, 491	4.9	1
100	Evaluation of Vector Competence of Ixodes Ticks for Kemerovo Virus. <i>Viruses</i> , 2022 , 14, 1102	6.2	
99	Identification of the Genome Segments of Bluetongue Virus Type 26/Type 1 Reassortants Influencing Horizontal Transmission in a Mouse Model. <i>Viruses</i> , 2021 , 13,	6.2	3
98	Exploration of binary protein-protein interactions between tick-borne flaviviruses and Ixodes ricinus. <i>Parasites and Vectors</i> , 2021 , 14, 144	4	3
97	How relevant are in vitro culture models for study of tick-pathogen interactions?. <i>Pathogens and Global Health</i> , 2021 , 115, 437-455	3.1	2
96	Inhibition of Orbivirus Replication by Fluvastatin and Identification of the Key Elements of the Mevalonate Pathway Involved. <i>Viruses</i> , 2021 , 13,	6.2	3
95	Vector Transmission of Animal Viruses 2021 , 542-551		1
94	Continuous Cell Lines from the European Biting Midge (Meigen, 1830). <i>Microorganisms</i> , 2020 , 8,	4.9	3
93	Orbiviruses. <i>Livestock Diseases and Management</i> , 2020 , 161-214	0.5	0
92	Strategies for Assessing Arbovirus Genetic Variability in Vectors and/or Mammals. <i>Pathogens</i> , 2020 , 9,	4.5	6
91	Bluetongue virus outer-capsid protein VP2 expressed in raises neutralising antibodies and a protective immune response in IFNAR mice. <i>Vaccine: X</i> , 2019 , 2, 100026	3.8	4
90	ICTV virus taxonomy profile: Birnaviridae. <i>Journal of General Virology</i> , 2019 , 100, 5-6	4.9	33
89	ICTV virus taxonomy profile: Picobirnaviridae. <i>Journal of General Virology</i> , 2019 , 100, 133-134	4.9	23
88	Emerging Mosquito-Borne Threats and the Response from European and Eastern Mediterranean Countries. <i>International Journal of Environmental Research and Public Health</i> , 2018 , 15,	4.6	21
87	Bluetongue Virus: From BTV-1 to BTV-27. <i>Advances in Virus Research</i> , 2017 , 99, 161-197	10.7	35
86	Complete Genome Sequence of Bluetongue Virus Serotype 8, Which Reemerged in France in August 2015. <i>Genome Announcements</i> , 2016 , 4,		21
85	Virus Discovery Using Tick Cell Lines. <i>Evolutionary Bioinformatics</i> , 2016 , 12, 31-4	1.9	14

84	Identification of the Genome Segments of Bluetongue Virus Serotype 26 (Isolate KUW2010/02) that Restrict Replication in a <i>Culicoides sonorensis</i> Cell Line (KC Cells). <i>PLoS ONE</i> , 2016 , 11, e0149709	3.7	15
83	Genetic characterization of the tick-borne orbiviruses. <i>Viruses</i> , 2015 , 7, 2185-209	6.2	24
82	Association of vectors and environmental conditions during the emergence of Peruvian horse sickness orbivirus and Yunnan orbivirus in northern Peru. <i>Journal of Vector Ecology</i> , 2015 , 40, 355-63	1.5	9
81	Orbiviruses and Bluetongue Virus 2015 , 1-18		
80	Coltivirus and Seadornaviruses 2015 , 1-12		
79	Full-Genome Sequencing as a Basis for Molecular Epidemiology Studies of Bluetongue Virus in India. <i>PLoS ONE</i> , 2015 , 10, e0131257	3.7	42
78	Immunisation with bacterial expressed VP2 and VP5 of bluetongue virus (BTV) protect \square interferon-receptor knock-out (IFNAR(-/-)) mice from homologous lethal challenge. <i>Vaccine</i> , 2014 , 32, 4059-67	4.1	21
77	Full-genome characterisation of Orungo, Lebombo and Changuinola viruses provides evidence for co-evolution of orbiviruses with their arthropod vectors. <i>PLoS ONE</i> , 2014 , 9, e86392	3.7	27
76	Full genome characterization of the culicoides-borne marsupial orbiviruses: Wallal virus, Mudjinbarry virus and Warrego viruses. <i>PLoS ONE</i> , 2014 , 9, e108379	3.7	6
75	Implicating <i>Culicoides</i> biting midges as vectors of Schmallenberg virus using semi-quantitative RT-PCR. <i>PLoS ONE</i> , 2013 , 8, e57747	3.7	66
74	Acute Schmallenberg virus infections, France, 2012. <i>Emerging Infectious Diseases</i> , 2013 , 19, 321-2	10.2	14
73	Endogenous tick viruses and modulation of tick-borne pathogen growth. <i>Frontiers in Cellular and Infection Microbiology</i> , 2013 , 3, 25	5.9	24
72	Full genome sequencing of Corriparta virus, identifies California mosquito pool virus as a member of the Corriparta virus species. <i>PLoS ONE</i> , 2013 , 8, e70779	3.7	6
71	Full genome sequencing and genetic characterization of Eubenangee viruses identify Pata virus as a distinct species within the genus <i>Orbivirus</i> . <i>PLoS ONE</i> , 2012 , 7, e31911	3.7	26
70	Isolates of Liao ning virus from wild-caught mosquitoes in the Xinjiang province of China in 2005. <i>PLoS ONE</i> , 2012 , 7, e37732	3.7	13
69	Seroprevalence of Toscana virus in blood donors, France, 2007. <i>Emerging Infectious Diseases</i> , 2011 , 17, 941-3	10.2	17
68	Complete genome characterisation of a novel 26th bluetongue virus serotype from Kuwait. <i>PLoS ONE</i> , 2011 , 6, e26147	3.7	136
67	Uniformity of rotavirus strain nomenclature proposed by the Rotavirus Classification Working Group (RCWG). <i>Archives of Virology</i> , 2011 , 156, 1397-413	2.6	699

66	Umatilla virus genome sequencing and phylogenetic analysis: identification of stretch lagoon orbivirus as a new member of the Umatilla virus species. <i>PLoS ONE</i> , 2011 , 6, e23605	3.7	32
65	Detection of a fourth orbivirus non-structural protein. <i>PLoS ONE</i> , 2011 , 6, e25697	3.7	155
64	Complete sequence of Great Island virus and comparison with the T2 and outer-capsid proteins of Kemerovo, Lipovnik and Tribec viruses (genus Orbivirus, family Reoviridae). <i>Journal of General Virology</i> , 2010 , 91, 2985-93	4.9	48
63	Isolation and full-length sequence analysis of Armigeres subalbatus totivirus, the first totivirus isolate from mosquitoes representing a proposed novel genus (Artivirus) of the family Totiviridae. <i>Journal of General Virology</i> , 2010 , 91, 2836-45	4.9	67
62	Detection and prevalence of the nonsyncytial American grass carp reovirus Aquareovirus G by quantitative reverse transcriptase polymerase chain reaction. <i>Journal of Aquatic Animal Health</i> , 2010 , 22, 8-13	2.6	13
61	The evolution of two homologues of the core protein VP6 of epizootic haemorrhagic disease virus (EHDV), which correspond to the geographical origin of the virus. <i>Virus Genes</i> , 2010 , 40, 67-75	2.3	8
60	Analysis of hepatitis C virus strains circulating in Republic of the Congo. <i>Journal of Medical Virology</i> , 2010 , 82, 562-7	19.7	16
59	The structure of bluetongue virus core and proteins 2009 , 101-133		6
58	Peruvian horse sickness virus and Yunnan orbivirus, isolated from vertebrates and mosquitoes in Peru and Australia. <i>Virology</i> , 2009 , 394, 298-310	3.6	53
57	Genetic and phylogenetic analysis of the core proteins VP1, VP3, VP4, VP6 and VP7 of epizootic haemorrhagic disease virus (EHDV). <i>Virus Research</i> , 2009 , 145, 187-99	6.4	37
56	Genetic and phylogenetic analysis of the outer-coat proteins VP2 and VP5 of epizootic haemorrhagic disease virus (EHDV): comparison of genetic and serological data to characterise the EHDV serogroup. <i>Virus Research</i> , 2009 , 145, 200-10	6.4	68
55	Genetic and phylogenetic analysis of the non-structural proteins NS1, NS2 and NS3 of epizootic haemorrhagic disease virus (EHDV). <i>Virus Research</i> , 2009 , 145, 211-9	6.4	28
54	Bluetongue virus, other orbiviruses and other reoviruses: Their relationships and taxonomy 2009 , 23-52		28
53	Complete characterisation of the American grass carp reovirus genome (genus Aquareovirus: family Reoviridae) reveals an evolutionary link between aquareoviruses and coltviruses. <i>Virology</i> , 2008 , 373, 310-21	3.6	82
52	Sequence analysis of bluetongue virus serotype 8 from the Netherlands 2006 and comparison to other European strains. <i>Virology</i> , 2008 , 377, 308-18	3.6	152
51	Sequence analysis of a reovirus isolated from the winter moth Operophtera brumata (Lepidoptera: Geometridae) and its parasitoid wasp Phobocampe tempestiva (Hymenoptera: Ichneumonidae). <i>Virus Research</i> , 2008 , 135, 42-7	6.4	10
50	Complete sequence characterization of isolates of Getah virus (genus Alphavirus, family Togaviridae) from China. <i>Journal of General Virology</i> , 2008 , 89, 1446-1456	4.9	42
49	Isolation and characterization of the full coding sequence of a novel densovirus from the mosquito Culex pipiens pallens. <i>Journal of General Virology</i> , 2008 , 89, 195-199	4.9	41

48	Human parvovirus 4 in kidney transplant patients, France. <i>Emerging Infectious Diseases</i> , 2008 , 14, 1811-210.2	10.2	10
47	Recommendations for the classification of group A rotaviruses using all 11 genomic RNA segments. <i>Archives of Virology</i> , 2008 , 153, 1621-9	2.6	523
46	Molecular characterization of genotype 2 and 4 hepatitis C virus isolates in French blood donors. <i>Journal of Medical Virology</i> , 2008 , 80, 1732-9	19.7	22
45	Rapid cDNA synthesis and sequencing techniques for the genetic study of bluetongue and other dsRNA viruses. <i>Journal of Virological Methods</i> , 2007 , 143, 132-9	2.6	173
44	Analysis and phylogenetic comparisons of full-length VP2 genes of the 24 bluetongue virus serotypes. <i>Journal of General Virology</i> , 2007 , 88, 621-630	4.9	172
43	Circular genomes related to anelloviruses identified in human and animal samples by using a combined rolling-circle amplification/sequence-independent single primer amplification approach. <i>Journal of General Virology</i> , 2007 , 88, 2696-2701	4.9	55
42	Complete nucleotide sequence of Middelburg virus, isolated from the spleen of a horse with severe clinical disease in Zimbabwe. <i>Journal of General Virology</i> , 2007 , 88, 3078-3088	4.9	18
41	Distribution and genetic analysis of TTV and TTMV major phylogenetic groups in French blood donors. <i>Journal of Medical Virology</i> , 2006 , 78, 298-304	19.7	51
40	Micromonas pusilla reovirus: a new member of the family Reoviridae assigned to a novel proposed genus (Mimoreovirus). <i>Journal of General Virology</i> , 2006 , 87, 1375-1383	4.9	49
39	Liao ning virus, a new Chinese seadornavirus that replicates in transformed and embryonic mammalian cells. <i>Journal of General Virology</i> , 2006 , 87, 199-208	4.9	44
38	The structure and function of the outer coat protein VP9 of Banna virus. <i>Structure</i> , 2005 , 13, 17-28	5.2	27
37	Expansion of family Reoviridae to include nine-segmented dsRNA viruses: isolation and characterization of a new virus designated Aedes pseudoscutellaris reovirus assigned to a proposed genus (Dinovernavirus). <i>Virology</i> , 2005 , 343, 212-23	3.6	65
36	Coltivirus and seadornaviruses in North America, Europe, and Asia. <i>Emerging Infectious Diseases</i> , 2005 , 11, 1673-9	10.2	80
35	Low diversity of Alkhurma hemorrhagic fever virus, Saudi Arabia, 1994-1999. <i>Emerging Infectious Diseases</i> , 2005 , 11, 683-8	10.2	50
34	Identification and functional analysis of VP3, the guanylyltransferase of Banna virus (genus Seadornavirus, family Reoviridae). <i>Journal of General Virology</i> , 2005 , 86, 1141-1146	4.9	19
33	Yunnan orbivirus, a new orbivirus species isolated from Culex tritaeniorhynchus mosquitoes in China. <i>Journal of General Virology</i> , 2005 , 86, 3409-3417	4.9	62
32	Structural organization of an encephalitic human isolate of Banna virus (genus Seadornavirus, family Reoviridae). <i>Journal of General Virology</i> , 2005 , 86, 1147-1157	4.9	40
31	Genotype distribution and molecular epidemiology of hepatitis C virus in blood donors from southeast France. <i>Journal of Clinical Microbiology</i> , 2005 , 43, 3624-9	9.7	37

30	Termination and read-through proteins encoded by genome segment 9 of Colorado tick fever virus. <i>Journal of General Virology</i> , 2004 , 85, 2237-2244	4.9	14
29	Sequences of flavivirus-related RNA viruses persist in DNA form integrated in the genome of <i>Aedes</i> spp. mosquitoes. <i>Journal of General Virology</i> , 2004 , 85, 1971-1980	4.9	224
28	Tick-borne virus diseases of human interest in Europe. <i>Clinical Microbiology and Infection</i> , 2004 , 10, 1040-55	4.9	210
27	Recombinant VP9-based enzyme-linked immunosorbent assay for detection of immunoglobulin G antibodies to Banna virus (genus Seadornavirus). <i>Journal of Virological Methods</i> , 2004 , 116, 55-61	2.6	10
26	Recombinant VP6-based enzyme-linked immunosorbent assay for detection of immunoglobulin G antibodies to Eyach virus (genus Coltivirus). <i>Journal of Clinical Virology</i> , 2004 , 30, 248-53	14.5	9
25	Recombinant VP7-based enzyme-linked immunosorbent assay for detection of immunoglobulin G antibodies to Colorado tick fever virus. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 2102-5	9.7	18
24	Evolution of hepatitis C virus in blood donors and their respective recipients. <i>Journal of General Virology</i> , 2003 , 84, 441-446	4.9	13
23	High genetic diversity revealed by the study of TLMV infection in French hemodialysis patients. <i>Journal of Medical Virology</i> , 2002 , 67, 630-5	19.7	17
22	Genus Coltivirus (family Reoviridae): genomic and morphologic characterization of Old World and New World viruses. <i>Archives of Virology</i> , 2002 , 147, 533-61	2.6	69
21	Common evolutionary origin of aquareoviruses and orthoreoviruses revealed by genome characterization of Golden shiner reovirus, Grass carp reovirus, Striped bass reovirus and golden ide reovirus (genus Aquareovirus, family Reoviridae). <i>Journal of General Virology</i> , 2002 , 83, 1941-1951	4.9	179
20	Erroneous HCV genotype assignment by a hybridization typing assay in a case of posttransfusion HCV infection. <i>Transfusion</i> , 2001 , 41, 429-30	2.9	3
19	Prevalence of antibody against West Nile virus in volunteer blood donors living in southeastern France. <i>Transfusion</i> , 2001 , 41, 1320-1	2.9	21
18	Complete sequence characterization of the genome of the St Croix River virus, a new orbivirus isolated from cells of <i>Ixodes scapularis</i> . <i>Journal of General Virology</i> , 2001 , 82, 795-804	4.9	64
17	Complete coding sequence of the Alkhurma virus, a tick-borne flavivirus causing severe hemorrhagic fever in humans in Saudi Arabia. <i>Biochemical and Biophysical Research Communications</i> , 2001 , 287, 455-61	3.4	111
16	Sequence characterization of Ndelle virus genome segments 1, 5, 7, 8, and 10: evidence for reassignment to the genus Orthoreovirus, family Reoviridae. <i>Biochemical and Biophysical Research Communications</i> , 2001 , 287, 583-8	3.4	35
15	TT virus infection: prevalence of elevated viraemia and arguments for the immune control of viral load. <i>Journal of Clinical Virology</i> , 2001 , 21, 135-41	14.5	68
14	Comparison of systems performance for TT virus detection using PCR primer sets located in non-coding and coding regions of the viral genome. <i>Journal of Clinical Virology</i> , 2001 , 22, 91-9	14.5	21
13	Genetic analysis of full-length genomes and subgenomic sequences of TT virus-like mini virus human isolates. <i>Journal of General Virology</i> , 2001 , 82, 379-383	4.9	53

12	Strategies for the sequence determination of viral dsRNA genomes. <i>Journal of Virological Methods</i> , 2000 , 89, 147-58	2.6	155
11	Complete sequences of two highly divergent european isolates of TT virus. <i>Biochemical and Biophysical Research Communications</i> , 2000 , 271, 837-41	3.4	30
10	Sequence determination and analysis of the full-length genome of colorado tick fever virus, the type species of genus Coltivirus (Family Reoviridae). <i>Biochemical and Biophysical Research Communications</i> , 2000 , 273, 1121-5	3.4	25
9	Sequence of genome segments 1, 2, and 3 of the grass carp reovirus (Genus Aquareovirus, family Reoviridae). <i>Biochemical and Biophysical Research Communications</i> , 2000 , 274, 762-6	3.4	73
8	TT virus: a study of molecular epidemiology and transmission of genotypes 1, 2 and 3. <i>Journal of Clinical Virology</i> , 2000 , 17, 43-9	14.5	44
7	Complete sequence determination and genetic analysis of Banna virus and Kadipiro virus: proposal for assignment to a new genus (Seadornavirus) within the family Reoviridae. <i>Journal of General Virology</i> , 2000 , 81, 1507-15	4.9	91
6	Prevalence of GB virus type C/hepatitis G virus RNA and anti-E2 among blood donors in Southeastern France. <i>Transfusion</i> , 1999 , 39, 95-102	2.9	17
5	Molecular diagnosis of group B coltivirus infections. <i>Journal of Virological Methods</i> , 1999 , 81, 39-45	2.6	13
4	The complete coding sequence of a European isolate of GB-C/hepatitis G virus. <i>Biochemical and Biophysical Research Communications</i> , 1999 , 255, 432-7	3.4	2
3	Serologic and molecular diagnosis of Colorado tick fever viral infections. <i>American Journal of Tropical Medicine and Hygiene</i> , 1998 , 59, 763-8	3.2	29
2	Evaluation of four PCR systems amplifying different genomic regions for molecular diagnosis of GB virus C infections. <i>Journal of Virological Methods</i> , 1997 , 64, 131-5	2.6	21
1	Coltiviruses and Seadornaviruses		1