Richard ThiÃ©ry

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8162496/publications.pdf

Version: 2024-02-01

70	2,568	172457	206112
papers	citations	29 h-index	48 g-index

71	71	71	2512
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Mastitis impact on technological properties of milk and quality of milk productsâ€"a review. Dairy Science and Technology, 2011, 91, 247-282.	2.2	140
2	Sea bream Sparus aurata, an asymptomatic contagious fish host for nodavirus. Diseases of Aquatic Organisms, 2001, 47, 33-38.	1.0	132
3	Comparative analysis of both genomic segments of betanodaviruses isolated from epizootic outbreaks in farmed fish species provides evidence for genetic reassortment. Journal of General Virology, 2009, 90, 2940-2951.	2.9	119
4	Induction of a Protective Immune Response against Viral Nervous Necrosis in the European Sea Bass Dicentrarchus labrax by Using Betanodavirus Virus-Like Particles. Journal of Virology, 2006, 80, 10201-10207.	3.4	107
5	Molecular characterization of Porcine circovirus type 2 isolates from post-weaning multisystemic wasting syndrome-affected and non-affected pigs. Journal of General Virology, 2004, 85, 293-304.	2.9	105
6	Genomic classification of new betanodavirus isolates by phylogenetic analysis of the coat protein gene suggests a low host-fish species specificity. Journal of General Virology, 2004, 85, 3079-3087.	2.9	98
7	Experimental vertical transmission of nodavirus from broodfish to eggs and larvae of the sea bass, Dicentrarchus labrax (L.). Journal of Fish Diseases, 2002, 25, 697-702.	1.9	94
8	Molecular characterisation and phylogenetic analysis of Chronic bee paralysis virus, a honey bee virus. Virus Research, 2008, 132, 59-68.	2.2	93
9	Comparative study of viral encephalopathy and retinopathy in juvenile sea bass Dicentrarchus labrax infected in different ways. Diseases of Aquatic Organisms, 1999, 36, 11-20.	1.0	80
10	Molecular Basis of Virulence in Staphylococcus aureus Mastitis. PLoS ONE, 2011, 6, e27354.	2.5	77
11	Generation of Replication-Defective Virus-Based Vaccines That Confer Full Protection in Sheep against Virulent Bluetongue Virus Challenge. Journal of Virology, 2011, 85, 10213-10221.	3.4	75
12	Emergence of pathogenic betanodaviruses belonging to the SJNNV genogroup in farmed fish species from the Iberian Peninsula. Journal of Fish Diseases, 2007, 30, 225-232.	1.9	71
13	Recombinant capripoxviruses expressing proteins of bluetongue virus: Evaluation of immune responses and protection in small ruminants. Vaccine, 2007, 25, 6774-6783.	3.8	70
14	Redescription of Gyrodactylus teuchis Lautraite, Blanc, Thiery, Daniel & Vigneulle, 1999 (Monogenea:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 141-150.	1.1	63
15	First detection of Israeli acute paralysis virus (IAPV) in France, a dicistrovirus affecting honeybees (Apis mellifera). Journal of Invertebrate Pathology, 2008, 99, 348-350.	3.2	58
16	Monitoring SARS-CoV-2 variants alterations in Nice neighborhoods by wastewater nanopore sequencing. Lancet Regional Health - Europe, The, 2021, 10, 100202.	5.6	56
17	Genetic characterization of ovine pestiviruses isolated in France, between 1985 and 2006. Veterinary Microbiology, 2008, 130, 69-79.	1.9	54
18	Natural outbreak of viral encephalopathy and retinopathy in juvenile sea bass, Dicentrarchus labrax: study by nested reverse transcriptaseâ€"polymerase chain reaction. Virus Research, 1999, 63, 11-17.	2.2	50

#	Article	IF	Citations
19	Dictyopterin, 6-(d-threo-1,2-dihydroxypropyl)-pterin, a new natural isomer of l-biopterin. Isolation from vegetative cells of Dictyostelium discoideum and identification. FEBS Journal, 1990, 187, 665-669.	0.2	48
20	Metabolisation of thiamethoxam (a neonicotinoid pesticide) and interaction with the Chronic bee paralysis virus in honeybees. Pesticide Biochemistry and Physiology, 2018, 144, 10-18.	3.6	47
21	Detection ofAethina tumidaMurray (Coleoptera: Nitidulidae.) in Italy: outbreaks and early reaction measures. Journal of Apicultural Research, 2014, 53, 569-575.	1.5	46
22	Two isolates of sea bass,Dicentrarchus labraxL., nervous necrosis virus with distinct genomes. Journal of Fish Diseases, 1999, 22, 201-207.	1.9	45
23	Staphylococcus aureus seroproteomes discriminate ruminant isolates causing mild or severe mastitis. Veterinary Research, 2011, 42, 35.	3.0	43
24	Development and validation of a real-time two-step RT-qPCR TaqManÂ® assay for quantitation of Sacbrood virus (SBV) and its application to a field survey of symptomatic honey bee colonies. Journal of Virological Methods, 2014, 197, 7-13.	2.1	38
25	Evolution of infectious hematopoietic necrosis virus (IHNV), a fish rhabdovirus, in Europe over 20 years: implications for control. Diseases of Aquatic Organisms, 2010, 89, 9-15.	1.0	37
26	Phylogenetic analysis of viral haemorrhagic septicaemia virus (VHSV) isolates from France (1971-1999). Diseases of Aquatic Organisms, 2002, 52, 29-37.	1.0	37
27	Viral encephalopathy and retinopathy of Dicentrarchus labrax and Sparus aurata farmed in Tunisia. Veterinary Research Communications, 2009, 33, 345-353.	1.6	35
28	Genome Sequences of Two Staphylococcus aureus Ovine Strains That Induce Severe (Strain O11) and Mild (Strain O46) Mastitis. Journal of Bacteriology, 2011, 193, 2353-2354.	2.2	30
29	Molecular epidemiology of Q fever in Poland. Polish Journal of Microbiology, 2009, 58, 9-13.	1.7	30
30	Bluetongue virus serotype 8 virus-like particles protect sheep against virulent virus infection as a single or multi-serotype cocktail immunogen. Vaccine, 2013, 31, 553-558.	3.8	28
31	Validation of quantitative real-time RT-PCR assays for the detection of six honeybee viruses. Journal of Virological Methods, 2019, 270, 70-78.	2.1	28
32	Influence of chronic exposure to thiamethoxam and chronic bee paralysis virus on winter honey bees. PLoS ONE, 2019, 14, e0220703.	2.5	27
33	Interactions Between Thiamethoxam and Deformed Wing Virus Can Drastically Impair Flight Behavior of Honey Bees. Frontiers in Microbiology, 2020, 11, 766.	3.5	27
34	An rt-pcr-based method for the diagnosis of the sleeping disease virus in experimentally and naturally infected salmonids. Diseases of Aquatic Organisms, 2000, 40, 19-27.	1.0	27
35	Difference in virulence betweenStaphylococcus aureusisolates causing gangrenous mastitis versus subclinical mastitis in a dairy sheep flock. Veterinary Research, 2009, 40, 56.	3.0	26
36	Serological and molecular evidence of Q fever among small ruminant flocks in Algeria. Comparative Immunology, Microbiology and Infectious Diseases, 2016, 47, 19-25.	1.6	26

#	Article	IF	Citations
37	Characterization of Coxiella burnetii strains from ruminants in a Galleria mellonella host-based model. New Microbes and New Infections, 2018, 24, 8-13.	1.6	25
38	Molecular typing of Coxiella burnetii from sheep in Egypt. Comparative Immunology, Microbiology and Infectious Diseases, 2019, 67, 101353.	1.6	25
39	Outcomes of honeybee pupae inoculated with deformed wing virus genotypes A and B. Apidologie, 2020, 51, 18-34.	2.0	22
40	Canine adenoviruses elicit both humoral and cell-mediated immune responses against rabies following immunisation of sheep. Vaccine, 2011, 29, 1304-1310.	3.8	21
41	Phylogenetic analysis of the RNA-dependent RNA polymerase (RdRp) and a predicted structural protein (pSP) of the Chronic bee paralysis virus (CBPV) isolated from various geographic regions. Virus Research, 2009, 144, 334-338.	2.2	20
42	Pestiviruses infections at the wild and domestic ruminants interface in the French Southern Alps. Veterinary Microbiology, 2015, 175, 341-348.	1.9	20
43	Impact of IS1111 insertion on the MLVA genotyping of Coxiella burnetii. Microbes and Infection, 2015, 17, 789-794.	1.9	19
44	First Description of Infection of Caprine Herpesvirus 1 (CpHV-1) in Goats in Mainland France. Pathogens, 2016, 5, 17.	2.8	18
45	Expression of VP7, a Bluetongue Virus Group Specific Antigen by Viral Vectors: Analysis of the Induced Immune Responses and Evaluation of Protective Potential in Sheep. PLoS ONE, 2014, 9, e111605.	2.5	15
46	Evaluation of the recombinant Heat shock protein B (HspB) of Coxiella burnetii as a potential antigen for immunodiagnostic of Q fever in goats. Veterinary Microbiology, 2009, 134, 300-304.	1.9	13
47	Draft Genome Sequences of Six Ruminant Coxiella burnetii Isolates of European Origin. Genome Announcements, 2014, 2, .	0.8	13
48	Coxiella burnetii in slaughterhouses in Brazil: A public health concern. PLoS ONE, 2020, 15, e0241246.	2.5	13
49	The effects of expression of an activated rasG mutation on the differentiation of Dictyostelium. Biochemistry and Cell Biology, 1992, 70, 1193-1199.	2.0	12
50	Outbreak of Q fever, Florac, Southern France, Spring 2007. Vector-Borne and Zoonotic Diseases, 2011, 11, 341-347.	1.5	12
51	Experimental infection of the honeybee (Apis mellifera L.) with the chronic bee paralysis virus (CBPV): infectivity of naked CBPV RNAs. Virus Research, 2012, 167, 173-178.	2.2	12
52	Characterisation of Mycoplasma capricolum P60 surface lipoprotein and its evaluation in a recombinant ELISA. Veterinary Microbiology, 2008, 128, 81-89.	1.9	11
53	EXPERIMENTAL INFECTION OF PREGNANT PYRENEAN CHAMOIS (RUPICAPRA PYRENAICA) WITH BORDER DISEASE VIRUS SUBTYPE 4. Journal of Wildlife Diseases, 2013, 49, 55-68.	0.8	11
54	Ras-related genes inDictyostelium discoideum. Genesis, 1991, 12, 147-153.	2.1	10

#	Article	IF	Citations
55	Molecular evolution and phylogeography of infectious hematopoietic necrosis virus with a focus on its presence in France over the last 30 years. Journal of General Virology, 2017, 98, 2438-2446.	2.9	10
56	Development of a semiquantitative PCR assay using internal standard and colorimetric detection on microwell plate for pseudorabies virus. Molecular and Cellular Probes, 1997, 11, 439-448.	2.1	9
57	Validation study for using lab-on-chip technology for Coxiella burnetii multi-locus-VNTR-analysis (MLVA) typing: application for studying genotypic diversity of strains from domestic ruminants in France. Microbes and Infection, 2015, 17, 782-788.	1.9	9
58	A fluorescence-based quantitative PCR method for investigation of pseudorabies virus latency. Journal of Virological Methods, 1996, 61, 79-87.	2.1	8
59	Characterisation of Structural Proteins from Chronic Bee Paralysis Virus (CBPV) Using Mass Spectrometry. Viruses, 2015, 7, 3329-3344.	3.3	8
60	Coxiella burnetii Transcriptional Analysis Reveals Serendipity Clusters of Regulation in Intracellular Bacteria. PLoS ONE, 2010, 5, e15321.	2.5	7
61	Staphylococcus aureus proteins differentially recognized by the ovine immune response in mastitis or nasal carriage. Veterinary Microbiology, 2012, 157, 439-447.	1.9	7
62	RNA 1 and RNA 2 Genomic Segments of Chronic Bee Paralysis Virus Are Infectious and Induce Chronic Bee Paralysis Disease. Journal of Immunology Research, 2015, 2015, 1-8.	2.2	7
63	Whole genome PCR scanning (WGPS) of Coxiella burnetii strains from ruminants. Microbes and Infection, 2015, 17, 772-775.	1.9	7
64	Phorbol 12-myristate 13-acetate modulates the cAMP-induced light-scattering response of aDictyostelium discoÃ¯deumcell population. FEBS Letters, 1988, 241, 149-153.	2.8	6
65	Development of a PCR-based method coupled with a microplate colorimetric assay for the detection of Porcine Parvovirus and application to diagnosis in piglet tissues and human plasma. Molecular and Cellular Probes, 1998, 12, 407-416.	2.1	6
66	Identification of Kashmir bee virus in France using a new RT-PCR method which distinguishes closely related viruses. Journal of Virological Methods, 2014, 198, 82-85.	2.1	6
67	Molecular detection of Coxiella burnetii in aborted bovine fetuses in Brazil. Acta Tropica, 2022, 227, 106258.	2.0	6
68	Staphylococcus aureus proteins differentially produced in ewe gangrenous mastitis or ewe milk. Veterinary Microbiology, 2013, 164, 150-157.	1.9	5
69	Increase of DPH fluorescence polarization during development ofDictyostelium discoideumcells. FEBS Letters, 1987, 223, 381-386.	2.8	3
70	A sporadic case of acute Q fever and identification of the animal source of the infection. Folia Microbiologica, 2020, 65, 797-800.	2.3	3