

Pablo Daniel Ghiringhelli

List of Publications by Year in descending order

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70
papers

1,523
citations

331670

21
h-index

361022

35
g-index

71
all docs

71
docs citations

71
times ranked

1790
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluation of the Nucleopolyhedrovirus of <i>Anticarsia gemmatalis</i> as a Vector for Gene Therapy in Mammals. <i>Current Gene Therapy</i> , 2021, 21, 177-189.	2.0	2
2	Valorization of brewer's spent grain by different strategies of structural destabilization and enzymatic saccharification. <i>Industrial Crops and Products</i> , 2021, 163, 113329.	5.2	14
3	Advances in the Bioinformatics Knowledge of mRNA Polyadenylation in Baculovirus Genes. <i>Viruses</i> , 2020, 12, 1395.	3.3	0
4	Novel insights into cardiac regeneration based on differential fetal and adult ovine heart transcriptomic analysis. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2020, 318, H994-H1007.	3.2	11
5	Impact of hepatitis B virus genotype F on in vitro diagnosis: detection efficiency of HBsAg from Amerindian subgenotypes F1b and F4. <i>Archives of Virology</i> , 2019, 164, 2297-2307.	2.1	5
6	High level production of a recombinant acid stable exoinulinase from <i>Aspergillus kawachii</i> . <i>Protein Expression and Purification</i> , 2018, 147, 29-37.	1.3	6
7	A comprehensive bioinformatic analysis of hepatitis D virus full-length genomes. <i>Journal of Viral Hepatitis</i> , 2018, 25, 860-869.	2.0	8
8	Potential of betabaculoviruses to control the tomato leafminer <i>Tuta absoluta</i> (Meyrick). <i>Journal of Applied Entomology</i> , 2018, 142, 67-77.	1.8	14
9	Relevance of Bacteriophage 933W in the Development of Hemolytic Uremic Syndrome (HUS). <i>Frontiers in Microbiology</i> , 2018, 9, 3104.	3.5	14
10	<i>Bacillus wiedmannii</i> biovar <i>thuringiensis</i> : a specialized mosquitocidal pathogen with plasmids from diverse origins. <i>Genome Biology and Evolution</i> , 2018, 10, 2823-2833.	2.5	28
11	Genomic analysis of an Argentinean isolate of <i>Spodoptera frugiperda</i> granulovirus reveals that various baculoviruses code for Lef-7 proteins with three F-box domains. <i>PLoS ONE</i> , 2018, 13, e0202598.	2.5	10
12	Control biológico de fitopatógenos, insectos y ácaros: Aplicaciones y perspectivas (volumen 2)., 2018, , .		2
13	Variability study of entomopathogenic nematode populations (Heterorhabditidae) from Argentina. <i>Brazilian Journal of Biology</i> , 2017, 77, 569-579.	0.9	12
14	Cationic Antimicrobial Peptides Inactivate Shiga Toxin-Encoding Bacteriophages. <i>Frontiers in Chemistry</i> , 2017, 5, 122.	3.6	5
15	Identification of <i>Diatraea</i> spp. (Lepidoptera: Crambidae) based on cytochrome oxidase II. <i>PLoS ONE</i> , 2017, 12, e0184053.	2.5	15
16	<i>Heterorhabditis bacteriophora</i> pampean-strain VElI (Nematoda): identification and pathogenicity against the strawberry pest <i>Lobiopa insularis</i> (Coleoptera: Nitidulidae). <i>Revista Colombiana De Entomologia</i> , 2017, 43, 223.	0.4	0
17	Telomerase as a Cancer Target. Development of New Molecules. <i>Current Topics in Medicinal Chemistry</i> , 2016, 16, 2432-2440.	2.1	62
18	Comparison of the efficiency of 5 methods for fungal DNA extraction from crop debris and evaluation of its suitability for the amplification of <i>Fusarium graminearum</i> by PCR. <i>Crop Protection</i> , 2016, 82, 7-9.	2.1	1

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19	Nucleotide sequence differentiation of argentine isolates of the mosquito parasitic nematode <i>Strelkovimermis spiculatus</i> (Nematoda: Mermithidae). <i>Journal of Vector Ecology</i> , 2015, 40, 415-418.	1.0	5
20	Evidence of recent interspecies horizontal gene transfer regarding nucleopolyhedrovirus infection of <i>Spodoptera frugiperda</i> . <i>BMC Genomics</i> , 2015, 16, 1008.	2.8	15
21	Novel Insights into the Evolution and Structural Characterization of Dyskerin Using Comprehensive Bioinformatics Analysis. <i>Journal of Proteome Research</i> , 2015, 14, 874-887.	3.7	5
22	Misregulation effect of a novel allelic variant in the Z promoter region found in cis with the CYP21A2 p.P482S mutation: implications for 21-hydroxylase deficiency. <i>Endocrine</i> , 2015, 50, 72-78.	2.3	4
23	The Complete Sequence of the First <i>Spodoptera frugiperda</i> Betabaculovirus Genome: A Natural Multiple Recombinant Virus. <i>Viruses</i> , 2015, 7, 394-421.	3.3	23
24	Potential Conservation of Circadian Clock Proteins in the phylum Nematoda as Revealed by Bioinformatic Searches. <i>PLoS ONE</i> , 2014, 9, e112871.	2.5	13
25	Protein universe containing a PUA RNA-binding domain. <i>FEBS Journal</i> , 2014, 281, 74-87.	4.7	18
26	Gramineous and non-gramineous weed species as alternative hosts of <i>Fusarium graminearum</i> , causal agent of <i>Fusarium</i> head blight of wheat, in Argentina. <i>Crop Protection</i> , 2014, 65, 100-104.	2.1	35
27	Role of bacteriophages in STEC infections: new implications for the design of prophylactic and treatment approaches. <i>F1000Research</i> , 2014, 3, 74.	1.6	11
28	Role of bacteriophages in STEC infections: new implications for the design of prophylactic and treatment approaches. <i>F1000Research</i> , 2014, 3, 74.	1.6	22
29	Promoter Sequence of Shiga Toxin 2 (Stx2) Is Recognized <i>In Vivo</i> , Leading to Production of Biologically Active Stx2. <i>MBio</i> , 2013, 4, e00501-13.	4.1	33
30	Family-Specific Degenerate Primer Design: A Tool to Design Consensus Degenerated Oligonucleotides. <i>Biotechnology Research International</i> , 2013, 2013, 1-9.	1.4	25
31	Identification of a Wee1-Like Kinase Gene Essential for Procyclic <i>Trypanosoma brucei</i> Survival. <i>PLoS ONE</i> , 2013, 8, e79364.	2.5	7
32	Functional Capacity of Shiga-Toxin Promoter Sequences in Eukaryotic Cells. <i>PLoS ONE</i> , 2013, 8, e57128.	2.5	37
33	<i>Aspergillus kawachii</i> produces an inulinase in cultures with yacon (<i>Smallanthus sonchifolius</i>) as substrate. <i>Electronic Journal of Biotechnology</i> , 2013, 16, .	2.2	5
34	First record of a mosquito iridescent virus in <i>Culex pipiens</i> L. (Diptera: Culicidae). <i>Archives of Virology</i> , 2012, 157, 1569-1571.	2.1	10
35	The <i>ac53</i> , <i>ac78</i> , <i>ac101</i> , and <i>ac103</i> Genes Are Newly Discovered Core Genes in the Family Baculoviridae. <i>Journal of Virology</i> , 2012, 86, 12069-12079.	3.4	132
36	Genome of <i>Epipotia aporema</i> granulovirus (EpapGV), a polyorganotropic fast killing betabaculovirus with a novel thymidylate kinase gene. <i>BMC Genomics</i> , 2012, 13, 548.	2.8	33

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37	Telomere structure and telomerase in health and disease. <i>International Journal of Oncology</i> , 2012, 41, 1561-1569.	3.3	126
38	Identification of nucleopolyhedrovirus that infect Nymphalid butterflies <i>Agraulis vanillae</i> and <i>Dione juno</i> . <i>Journal of Invertebrate Pathology</i> , 2011, 106, 255-262.	3.2	4
39	Production of heterologous polygalacturonase I from <i>Aspergillus kawachii</i> in <i>Saccharomyces cerevisiae</i> in batch and fed-batch cultures. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2011, 38, 1437-1447.	3.0	11
40	Baculovirus: Molecular Insights on Their Diversity and Conservation. <i>International Journal of Evolutionary Biology</i> , 2011, 2011, 1-15.	1.0	88
41	Molecular analysis of the virulence attenuation process in Jun \bar{A} n virus vaccine genealogy. <i>Virus Genes</i> , 2010, 40, 320-328.	1.6	24
42	Effects of Fetal Bovine Serum deprivation in cell cultures on the production of <i>Anticarsia gemmatalis</i> Multinucleopolyhedrovirus. <i>BMC Biotechnology</i> , 2010, 10, 68.	3.3	9
43	Expression and Purification of Z Protein from Jun \bar{A} n Virus. <i>Journal of Biomedicine and Biotechnology</i> , 2010, 2010, 1-14.	3.0	4
44	A DNA Vaccine Encoding the Enterohemorrhagic <i>Escherichia coli</i> Shiga-Like Toxin 2 A 2 and B Subunits Confers Protective Immunity to Shiga Toxin Challenge in the Murine Model. <i>Vaccine Journal</i> , 2009, 16, 712-718.	3.1	33
45	Multiplex PCR and quality control of <i>Epipotia aporema</i> granulovirus production. <i>Virus Genes</i> , 2008, 37, 203-211.	1.6	6
46	Argentine hemorrhagic fever diagnostic test based on recombinant Jun \bar{A} n virus N protein. <i>Journal of Medical Virology</i> , 2008, 80, 2127-2133.	5.0	10
47	Two simultaneous hepatitis B virus epidemics among injecting drug users and men who have sex with men in Buenos Aires, Argentina: characterization of the first D/A recombinant from the American continent. <i>Journal of Viral Hepatitis</i> , 2008, 15, 080527190031013-???	2.0	21
48	Functional and structural characterisation of AgMNPV ie1. <i>Virus Genes</i> , 2007, 35, 549-562.	1.6	7
49	Genomic Features of Attenuated Jun \bar{A} n Virus Vaccine Strain Candidate. <i>Virus Genes</i> , 2006, 32, 37-41.	1.6	42
50	Sequencing and Characterisation of p74 Gene in Two Isolates of <i>Anticarsia Gemmatalis</i> MNPV. <i>Virus Genes</i> , 2006, 32, 59-70.	1.6	6
51	Molecular cloning and sequence analysis of the <i>Anticarsia gemmatalis</i> multicapsid nuclear polyhedrosis virus GP64 glycoprotein. <i>Virus Genes</i> , 2003, 26, 57-69.	1.6	7
52	Evaluation of the proacrosin/acrosin system and its mechanism of activation in human sperm extracts. <i>Journal of Reproductive Immunology</i> , 2002, 54, 43-63.	1.9	31
53	Identification and characterization of the ecdysteroid UDP-glycosyltransferase gene of <i>Epipotia aporema</i> granulovirus. <i>Virus Genes</i> , 2002, 24, 119-130.	1.6	8
54	Physical and genetic map of <i>Epipotia aporema</i> granulovirus genome. <i>Virus Genes</i> , 2002, 25, 329-341.	1.6	9

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55	Arenavirus nucleocapsid protein displays a transcriptional antitermination activity in vivo. <i>Virus Research</i> , 2001, 73, 41-55.	2.2	49
56	Generation of a recombinant <i>Anticarsia gemmatalis</i> multicapsid nucleopolyhedrovirus expressing a foreign gene under the control of a very late promoter. <i>Virus Genes</i> , 2001, 22, 363-372.	1.6	8
57	Characterization of a Granulovirus Isolated from <i>Epinotia aporema</i> Wals. (Lepidoptera: Tortricidae) Larvae. <i>Applied and Environmental Microbiology</i> , 2001, 67, 3702-3706.	3.1	28
58	Zinc-binding properties of JunÃn virus nucleocapsid protein. <i>Journal of General Virology</i> , 2001, 82, 121-128.	2.9	22
59	Characterization of human group C rotavirus in Argentina. <i>Journal of Medical Virology</i> , 2000, 62, 199-207.	5.0	22
60	Engineering a compact non-native state of intestinal fatty acid-binding protein. <i>BBA - Proteins and Proteomics</i> , 2000, 1476, 203-218.	2.1	19
61	Arenavirus phylogeny: a new insight. <i>Virus Genes</i> , 1998, 16, 39-46.	1.6	20
62	Molecular characterization of attenuated Junin virus strains.. <i>Journal of General Virology</i> , 1997, 78, 1605-1610.	2.9	39
63	Characterization of arenaviruses using a family-specific primer set for RT-PCR amplification and RFLP analysis. <i>Virus Research</i> , 1997, 49, 79-89.	2.2	53
64	Expression of Properly Folded Human Glutamate Decarboxylase 65 as a Fusion Protein in <i>Escherichia Coli</i> . <i>FEBS Journal</i> , 1997, 246, 350-359.	0.2	25
65	The Glycoprotein Precursor Gene of the Attenuated Junin Virus Vaccine Strain (Candid #1). <i>American Journal of Tropical Medicine and Hygiene</i> , 1997, 56, 216-225.	1.4	13
66	Computational characterisation of potential RNA-binding sites in arenavirus nucleocapsid proteins. <i>Virus Genes</i> , 1996, 13, 247-254.	1.6	11
67	A simple nucleic acid amplification assay for the rapid detection of JunÃn virus in whole blood samples. <i>Virus Research</i> , 1993, 27, 37-53.	2.2	23
68	Molecular organization of Junin virus S RNA: complete nucleotide sequence, relationship with other members of the Arenaviridae and unusual secondary structures. <i>Journal of General Virology</i> , 1991, 72, 2129-2141.	2.9	69
69	Nucleocapsid protein gene of Junin arenavirus (cDNA sequence). <i>Nucleic Acids Research</i> , 1989, 17, 8001-8001.	14.5	18
70	Cloned cDNA Probes for the Detection of Tomato Spotted Wilt Virus. <i>Phytopathology</i> , 1989, 79, 1309.	2.2	15