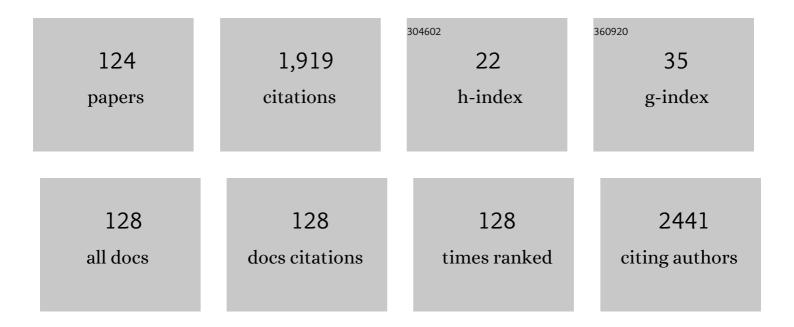
## Bergmann M Ribeiro

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identification of potential new mosquito-associated viruses of adult Aedes aegypti mosquitoes from Tocantins state, Brazil. Brazilian Journal of Microbiology, 2022, 53, 51-62.	0.8	6
2	An easy pipeline for one-step purification of SARS-CoV-2 nucleocapsid protein from insect cell suspension culture. Journal of Virological Methods, 2022, 299, 114341.	1.0	5
3	Genomic Epidemiology of SARS-CoV-2 in Tocantins State and the Diffusion of P.1.7 and AY.99.2 Lineages in Brazil. Viruses, 2022, 14, 659.	1.5	8
4	Characterization and genomic analyses of a novel alphabaculovirus isolated from the black armyworm, Spodoptera cosmioides (Lepidoptera: Noctuidae). Virus Research, 2022, 316, 198797.	1.1	2
5	Mosaic genome evolution and phylogenetics of Chrysodeixis includens nucleopolyhedrovirus (ChinNPV) and virulence of seven new isolates from the Brazilian states of Minas Gerais and Mato Grosso. Archives of Virology, 2021, 166, 125-138.	0.9	1
6	Genomic analyses of Biston suppressaria nucleopolyhedrovirus: a viral isolate obtained from the tea looper caterpillar, Biston suppressaria (Guenée, 1857). Brazilian Journal of Microbiology, 2021, 52, 219-227.	0.8	2
7	Molecular characterization and sequence analysis of four Brazilian rice stripe necrosis virus isolates. Archives of Virology, 2021, 166, 1763-1767.	0.9	2
8	Cassava starch-based essential oil microparticles preparations: Functionalities in mosquito control and selectivity against non-target organisms. Industrial Crops and Products, 2021, 162, 113289.	2.5	11
9	Molecular characterization of a putative new cavemovirus isolated from wild chicory (Cichorium) Tj ETQq1 10.	784314 rgl 0.9	3T /Overlock 1 1
10	Sialovirome of Brazilian tropical anophelines. Virus Research, 2021, 302, 198494.	1.1	6
11	Real-Time Genomic Surveillance during the 2021 Re-Emergence of the Yellow Fever Virus in Rio Grande do Sul State, Brazil. Viruses, 2021, 13, 1976.	1.5	23
12	Easily purified baculovirus/insect-system-expressed recombinant hepatitis B virus surface antigen fused to the N- or C-terminus of polyhedrin. Archives of Virology, 2021, , 1.	0.9	2
13	No Evidence of SARS-CoV-2 Infection in Neotropical Primates Sampled During COVID-19 Pandemic in Minas Gerais and Rio Grande do Sul, Brazil. EcoHealth, 2021, 18, 414-420.	0.9	3
14	Chikungunya virus produced by a persistently infected mosquito cell line comprises a shorter genome and is non-infectious to mammalian cells. Journal of General Virology, 2021, 102, .	1.3	1
15	Identification of <i>Physalis angulata</i> (Solanaceae) as a Natural Alternative Weed Host of Tomato Severe Rugose Virus in Brazil. Plant Disease, 2020, 104, 600-600.	0.7	7
16	Evaluation of theÂanti-apoptotic activity of bovine alphaherpesvirus type 5 US3 protein kinase in insect cells using a recombinant baculovirus. Brazilian Journal of Microbiology, 2020, 51, 827-835.	0.8	1
17	Dengue and Zika virus multi-epitope antigen expression in insect cells. Molecular Biology Reports, 2020, 47, 7333-7340.	1.0	6
18	Identification and genome sequencing of RNA viruses in the eucalyptus snout beetle Gonipterus spp. (Coleoptera: Curculionidae). Archives of Virology, 2020, 165, 2993-2997.	0.9	3

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19	Entomopathogenic Viruses in the Neotropics: Current Status and Recently Discovered Species. Neotropical Entomology, 2020, 49, 315-331.	0.5	18
20	Complete Genome Sequences of Seven New Chrysodeixis includens Nucleopolyhedrovirus Isolates from Minas Gerais and Mato Grosso States in Brazil. Microbiology Resource Announcements, 2020, 9, .	0.3	2
21	Haematological and biochemical parameters of wild capuchin monkeys in BrasÃłia, Federal District—Brazil. Journal of Medical Primatology, 2020, 49, 211-217.	0.3	4
22	Trichoplusia ni and Chrysodeixis includens larvae show different susceptibility to Chrysodeixis includens single nucleopolyhedrovirus per os infection. Journal of Pest Science, 2020, 93, 1019-1029.	1.9	3
23	Characterization of a novel alphabaculovirus isolated from the Southern armyworm, Spodoptera eridania (Cramer, 1782) (Lepidoptera: Noctuidae) and the evolution of odv-e66, a bacterium-acquired baculoviral chondroitinase gene. Genomics, 2020, 112, 3903-3914.	1.3	4
24	Comparative genomic analysis and mosquito larvicidal activity of four Bacillus thuringiensis serovar israelensis strains. Scientific Reports, 2020, 10, 5518.	1.6	6
25	A novel cypovirus found in a betabaculovirus co-infection context contains a poxvirus immune nuclease (poxin)-related gene. Journal of General Virology, 2020, 101, 667-675.	1.3	2
26	Prolonged mosquitocidal activity of Siparuna guianensis essential oil encapsulated in chitosan nanoparticles. PLoS Neglected Tropical Diseases, 2019, 13, e0007624.	1.3	50
27	A Nymphalid-Infecting Group I Alphabaculovirus Isolated from the Major Passion Fruit Caterpillar Pest Dione juno juno (Lepidoptera: Nymphalidae). Viruses, 2019, 11, 602.	1.5	4
28	Novel Viruses in Mosquitoes from Brazilian Pantanal. Viruses, 2019, 11, 957.	1.5	13
29	Faecal Virome Analysis of Wild Animals from Brazil. Viruses, 2019, 11, 803.	1.5	51
30	Inactivated alpha toxin from Clostridium novyi type B in nano-emulsion protect partially protects Swiss mice from lethal alpha toxin challenge. Scientific Reports, 2019, 9, 14082.	1.6	5
31	Characterization of a bacteriophage with broad host range against strains of Pseudomonas aeruginosa isolated from domestic animals. BMC Microbiology, 2019, 19, 134.	1.3	35
32	The complete genome of Rachiplusia nu nucleopolyhedrovirus (RanuNPV) and the identification of a baculoviral CPD-photolyase homolog. Virology, 2019, 534, 64-71.	1.1	8
33	An iflavirus found in stink bugs (Hemiptera: Pentatomidae) of four different species. Virology, 2019, 534, 72-79.	1.1	14
34	A new virus found in garlic virus complex is a member of possible novel genus of the family Betaflexiviridae (order Tymovirales). PeerJ, 2019, 7, e6285.	0.9	20
35	Cell-line-dependent crystal morphology and sublocalization of the Thyrinteina arnobia cypovirus polyhedrin expressed from a recombinant baculovirus. Archives of Virology, 2019, 164, 1677-1682.	0.9	2
36	Assembly of tomato blistering mosaic virus-like particles using a baculovirus expression vector system. Archives of Virology, 2019, 164, 1753-1760.	0.9	3

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37	Complete Sequences of Two Plasmids Found in a Brazilian Bacillus thuringiensis Serovar israelensis Strain. Microbiology Resource Announcements, 2019, 8, .	0.3	0
38	Nanopore sequencing of a novel bipartite New World begomovirus infecting cowpea. Archives of Virology, 2019, 164, 1907-1910.	0.9	22
39	Biological and molecular characterization of two Anticarsia gemmatalis multiple nucleopolyhedrovirus clones exhibiting contrasting virulence. Journal of Invertebrate Pathology, 2019, 164, 23-31.	1.5	3
40	Tailed giant Tupanvirus possesses the most complete translational apparatus of the known virosphere. Nature Communications, 2018, 9, 749.	5.8	247
41	The complete genome sequence of the first hesperiid-infecting alphabaculovirus isolated from the leguminous pest Urbanus proteus (Lepidoptera: Hesperiidae). Virus Research, 2018, 249, 76-84.	1.1	12
42	Genome-wide diversity in temporal and regional populations of the betabaculovirus Erinnyis ello granulovirus (ErelGV). BMC Genomics, 2018, 19, 698.	1.2	11
43	External Quality Assessment for Zika Virus Molecular Diagnostic Testing, Brazil. Emerging Infectious Diseases, 2018, 24, 888-892.	2.0	29
44	Genome Sequence of a New Siphoviridae Phage Found in a Brazilian Bacillus thuringiensis Serovar israelensis Strain. Genome Announcements, 2018, 6, .	0.8	0
45	A Novel Betabaculovirus Isolated from the Monocot Pest Mocis latipes (Lepidoptera: Noctuidae) and the Evolution of Multiple-Copy Genes. Viruses, 2018, 10, 134.	1.5	7
46	Viola phlebovirus is a novel Phlebotomus fever serogroup member identified in Lutzomyia (Lutzomyia) longipalpis from Brazilian Pantanal. Parasites and Vectors, 2018, 11, 405.	1.0	14
47	Essential oil of Siparuna guianensis as an alternative tool for improved lepidopteran control and resistance management practices. Scientific Reports, 2018, 8, 7215.	1.6	33
48	Genomic analysis of a cypovirus isolated from the eucalyptus brown looper, Thyrinteina arnobia (Stoll, 1782) (Lepidoptera: Geometridae). Virus Research, 2018, 253, 62-67.	1.1	5
49	Production of GP64-free virus-like particles from baculovirus-infected insect cells. Journal of General Virology, 2018, 99, 265-274.	1.3	15
50	Discovery of two small circular ssDNA viruses associated with the whitefly Bemisia tabaci. Archives of Virology, 2017, 162, 2835-2838.	0.9	6
51	Expression of Bacillus thuringiensis Toxins in Insect Cells. , 2017, , 99-110.		2
52	The genome sequence of Condylorrhiza vestigialis NPV, a novel baculovirus for the control of the Alamo moth on Populus spp. in Brazil. Journal of Invertebrate Pathology, 2017, 148, 152-161.	1.5	5
53	Real-Time Expression Analysis of Selected Anticarsia gemmatalis multiple nucleopolyhedrovirus Gene Promoters during Infection of Permissive, Semipermissive and Nonpermissive Cell Lines. Viruses, 2017, 9, 132.	1.5	3
54	Novel viruses in salivary glands of mosquitoes from sylvatic Cerrado, Midwestern Brazil. PLoS ONE, 2017, 12, e0187429.	1.1	34

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55	New Sequencing Technologies and Genomic Analysis Applied to Bacillus thuringiensis. , 2017, , 89-97.		0
56	Complete Genome Sequences of Six <i>Chrysodeixis includens</i> Nucleopolyhedrovirus Isolates from Brazil and Guatemala. Genome Announcements, 2016, 4, .	0.8	11
57	Genome sequence of Perigonia lusca single nucleopolyhedrovirus: insights into the evolution of a nucleotide metabolism enzyme in the family Baculoviridae. Scientific Reports, 2016, 6, 24612.	1.6	11
58	The complete genome of a baculovirus isolated from an insect of medical interest: Lonomia obliqua (Lepidoptera: Saturniidae). Scientific Reports, 2016, 6, 23127.	1.6	11
59	A betabaculovirus encoding a gp64 homolog. BMC Genomics, 2016, 17, 94.	1.2	8
60	A Betabaculovirus-Encoded gp64 Homolog Codes for a Functional Envelope Fusion Protein. Journal of Virology, 2016, 90, 1668-1672.	1.5	12
61	Draft genome sequence of Bacillus thuringiensis 147, a Brazilian strain with high insecticidal activity. Memorias Do Instituto Oswaldo Cruz, 2015, 110, 822-823.	0.8	6
62	Insecticidal and Repellent Activity of Siparuna guianensis Aubl. (Negramina) against Aedes aegypti and Culex quinquefasciatus. PLoS ONE, 2015, 10, e0116765.	1.1	49
63	Production of Brazilian human norovirus VLPs and comparison of purification methods. Brazilian Journal of Microbiology, 2015, 46, 1265-1268.	0.8	6
64	Functional characterization of hesp018, a baculovirus-encoded serpin gene. Journal of General Virology, 2015, 96, 1150-1160.	1.3	17
65	Complete genome sequence and structural characterization of a novel iflavirus isolated from Opsiphanes invirae (Lepidoptera: Nymphalidae). Journal of Invertebrate Pathology, 2015, 130, 136-140.	1.5	17
66	The genome sequence of Pseudoplusia includens single nucleopolyhedrovirus and an analysis of p26 gene evolution in the baculoviruses. BMC Genomics, 2015, 16, 127.	1.2	22
67	The silencing suppressor (NSs) protein of the plant virus Tomato spotted wilt virus enhances heterologous protein expression and baculovirus pathogenicity in cells and lepidopteran insects. Archives of Virology, 2015, 160, 2873-2879.	0.9	12
68	CHARACTERIZATION OF HELICOVERPA ZEA SINGLE NUCLEOPOLYHEDROVIRUS ISOLATED IN BRAZIL DURING THE FIRST OLD WORLD BOLLWORM (NOCTUIDAE: HELICOVERPA ARMIGERA) NATIONWIDE OUTBREAK. Virus Reviews & Research: Journal of the Brazilian Society for Virology, 2015, 20, .	0.1	10
69	Complete Genome Sequence of a Tobacco-Infecting, Tomato-Blistering Mosaic Virus. Genome Announcements, 2014, 2, .	0.8	8
70	Genome sequence of Erinnyis ello granulovirus (ErelGV), a natural cassava hornworm pesticide and the first sequenced sphingid-infecting betabaculovirus. BMC Genomics, 2014, 15, 856.	1.2	23
71	Proteomic analyses of baculovirus Anticarsia gemmatalis multiple nucleopolyhedrovirus budded and occluded virus. Journal of General Virology, 2014, 95, 980-989.	1.3	19
72	Complete genome sequence of the first non-Asian isolate of Bombyx mori nucleopolyhedrovirus. Virus Genes, 2014, 49, 477-484.	0.7	18

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73	High genetic stability of peroral infection factors from Anticarsia gemmatalis MNPV over 20years of sampling. Journal of Invertebrate Pathology, 2014, 118, 66-70.	1.5	4
74	Pseudoplusia includens single nucleopolyhedrovirus: Genetic diversity, phylogeny and hypervariability of the pif-2 gene. Journal of Invertebrate Pathology, 2013, 114, 258-267.	1.5	19
75	A baculovirus-mediated strategy for full-length plant virus coat protein expression and purification. Virology Journal, 2013, 10, 262.	1.4	11
76	Immunological effects of Anticarsia gemmatalis multiple nucleopolyhedrovirus (AgMNPV) by stimulation of mice in vivo and in vitro. Virus Research, 2013, 176, 119-127.	1.1	3
77	New Binding Site Conformations of the Dengue Virus NS3 Protease Accessed by Molecular Dynamics Simulation. PLoS ONE, 2013, 8, e72402.	1.1	19
78	A Recombinant Anticarsia gemmatalis MNPV Harboring chiA and v-cath Genes from Choristoneura fumiferana Defective NPV Induce Host Liquefaction and Increased Insecticidal Activity. PLoS ONE, 2013, 8, e74592.	1.1	16
79	A New theraphosid Spider Toxin Causes Early Insect Cell Death by Necrosis When Expressed In Vitro during Recombinant Baculovirus Infection. PLoS ONE, 2013, 8, e84404.	1.1	13
80	Cytotoxicity Analysis of Three Bacillus thuringiensis Subsp. israelensis δ-Endotoxins towards Insect and Mammalian Cells. PLoS ONE, 2012, 7, e46121.	1.1	13
81	A silencing suppressor protein (NSs) of a tospovirus enhances baculovirus replication in permissive and semipermissive insect cell lines. Virus Research, 2011, 155, 259-267.	1.1	25
82	Expression of recombinant Araraquara Hantavirus nucleoprotein in insect cells and its use as an antigen for immunodetection compared to the same antigen expressed in Escherichia coli. Virology Journal, 2011, 8, 218.	1.4	4
83	Identification and sequence analysis of the Condylorrhiza vestigialis MNPV p74 gene. Virus Genes, 2011, 43, 471-475.	0.7	2
84	A-type inclusion bodies: a factor influencing cowpox virus lesion pathogenesis. Archives of Virology, 2011, 156, 617-628.	0.9	7
85	Yellow fever virus envelope protein expressed in insect cells is capable of syncytium formation in lepidopteran cells and could be used for immunodetection of YFV in human sera. Virology Journal, 2011, 8, 261.	1.4	8
86	The major leucyl aminopeptidase of Trypanosoma cruzi (LAPTc) assembles into a homohexamer and belongs to the M17 family of metallopeptidases. BMC Biochemistry, 2011, 12, 46.	4.4	33
87	Comparative analysis of American Dengue virus type 1 full-genome sequences. Virus Genes, 2010, 40, 60-66.	0.7	15
88	Insecticidal activity of two proteases against Spodoptera frugiperda larvae infected with recombinant baculoviruses. Virology Journal, 2010, 7, 143.	1.4	40
89	Midgut GPI-anchored proteins with alkaline phosphatase activity from the cotton boll weevil (Anthonomus grandis) are putative receptors for the Cry1B protein of Bacillus thuringiensis. Insect Biochemistry and Molecular Biology, 2010, 40, 138-145.	1.2	40
90	Molecular analysis of a mutant <i>Anticarsia gemmatalis multiple nucleopolyhedrovirus</i> (AgMNPV) shows an interruption of an inhibitor of apoptosis gene ( <i>iapâ€3</i> ) by a new classâ€I <i>piggyBac</i> â€related insect transposon. Insect Molecular Biology, 2009, 18, 747-757.	1.0	6

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91	Characterization of a new Autographa californica multiple nucleopolyhedrovirus (AcMNPV) polyhedra mutant. Virus Research, 2009, 140, 1-7.	1.1	10
92	Identification of a new nucleopolyhedrovirus from naturally-infected Condylorrhiza vestigialis (Guenée) (Lepidoptera: Crambidae) larvae on poplar plantations in South Brazil. Journal of Invertebrate Pathology, 2009, 102, 149-154.	1.5	13
93	Cry2A toxins from Bacillus thuringiensis expressed in insect cells are toxic to two lepidopteran insects. World Journal of Microbiology and Biotechnology, 2008, 24, 2941-2948.	1.7	15
94	Recombinant Cry11a protein is highly toxic to cotton boll weevil (Anthonomus grandis Boheman) and fall armyworm (Spodoptera frugiperda). Journal of Applied Microbiology, 2008, 104, 1363-1371.	1.4	33
95	Recombinant expression of Garlic virus C (GARV-C) capsid protein in insect cells and its potential for the production of specific antibodies. Microbiological Research, 2008, 163, 354-361.	2.5	8
96	Structural and ultrastructural alterations of Malpighian tubules of Anticarsia gemmatalis (Hübner) (Lepidoptera: Noctuidae) larvae infected with different Anticarsia gemmatalis multiple nucleopolyhedrovirus (AgMNPV) recombinant viruses. Journal of Invertebrate Pathology, 2008, 98, 7-19.	1.5	7
97	An Anticarsia gemmatalis multiple nucleopolyhedrovirus mutant, vApAg, induces hemocytes apoptosis in vivo and displays reduced infectivity in larvae of Anticarsia gemmatalis (Hübner) (Lepidoptera:) Tj ETQq1 1 (	).784314	rgBa /Overlo
98	Characterization of PfDYN2, a dynamin-like protein of Plasmodium falciparum expressed in schizonts. Microbes and Infection, 2007, 9, 797-805.	1.0	18
99	Functional and structural characterisation of AgMNPV ie1. Virus Genes, 2007, 35, 549-562.	0.7	7
100	Introduction of the anti-apoptotic baculovirus p35 gene in passion fruit induces herbicide tolerance, reduced bacterial lesions, but does not inhibits passion fruit woodiness disease progress induced by cowpea aphid-borne mosaic virus (CABMV). Biotechnology Letters, 2007, 29, 79-87.	1.1	23
101	Infectivity of Anticarsia gemmatalis nucleopolyhedrovirus to different insect cell lines: Morphology, viral production, and protein synthesis. Biological Control, 2006, 36, 299-304.	1.4	14
102	A Recombinant Truncated Cry1Ca Protein Is Toxic to Lepidopteran Insects and Forms Large Cuboidal Crystals in Insect Cells. Current Microbiology, 2006, 53, 287-292.	1.0	10
103	Genome of the most widely used viral biopesticide: Anticarsia gemmatalis multiple nucleopolyhedrovirus. Journal of General Virology, 2006, 87, 3233-3250.	1.3	76
104	The inhibitor of apoptosis gene (iap-3) of Anticarsia gemmatalis multicapsid nucleopolyhedrovirus (AgMNPV) encodes a functional IAP. Archives of Virology, 2005, 150, 1549-1562.	0.9	26
105	Cell Culture Derived AgMNPV Bioinsecticide: Biological Constraints and Bioprocess Issues. Cytotechnology, 2005, 48, 27-39.	0.7	21
106	Pathology of Anticarsia gemmatalis larvae infected by two recombinant A. gemmatalis multicapsid nucleopolyhedroviruses. Research in Microbiology, 2005, 156, 263-269.	1.0	14
107	In vivo apoptosis induction and reduction of infectivity by an Autographa californica multiple nucleopolyhedrovirus p35â^' recombinant in hemocytes from the velvet bean caterpillar Anticarsia gemmatalis (Hübner) (Lepidoptera: Noctuidae). Research in Microbiology, 2005, 156, 1014-1025.	1.0	9
108	Identification, Expression and Phylogenetic Analysis of the Anticarsia gemmatalis multicapsid nucleopolyhedrovirus (AgMNPV) Helicase. Virus Genes, 2004, 29, 345-352.	0.7	5

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109	Morphological characterization of Anticarsia gemmatalis M nucleopolyhedrovirus infection in haemocytes from its natural larval host, the velvet bean caterpillar Anticarsia gemmatalis (Hübner) (Lepidoptera: Noctuidae). Tissue and Cell, 2004, 36, 171-180.	1.0	6
110	The gp64 locus of Anticarsia gemmatalis multicapsid nucleopolyhedrovirus contains a 3′ repair exonuclease homologue and lacks v-cath and ChiA genes. Journal of General Virology, 2004, 85, 211-219.	1.3	27
111	Transcriptome characterization of the dimorphic and pathogenic fungusParacoccidioides brasiliensisby EST analysis. Yeast, 2003, 20, 263-271.	0.8	74
112	Inactivation of the ecdysteroid UDP-glucosyltransferase (egt) gene of Anticarsia gemmatalis nucleopolyhedrovirus (AgMNPV) improves its virulence towards its insect host. Biological Control, 2003, 27, 336-344.	1.4	30
113	Characterization of the p10 gene region of Anticarsia gemmatalis nucleopolyhedrovirus. Virus Genes, 2002, 24, 243-247.	0.7	10
114	Construction of a recombinant Anticarsia gemmatalis nucleopolyhedrovirus (AgMNPV-2D) harbouring the Î <sup>2</sup> -galactosidase gene. Archives of Virology, 2001, 146, 1355-1367.	0.9	13
115	Characterization of the ecdysteroid UDP-glucosyltransferase (egt) gene of Anticarsia gemmatalis nucleopolyhedrovirus. Virus Genes, 2001, 22, 103-112.	0.7	24
116	Production of viral progeny in insect cells undergoing apoptosis induced by a mutant Anticarsia gemmatalis nucleopolyhedrovirus. Microbiological Research, 2001, 156, 369-376.	2.5	6
117	Screening and characterization of Bacillus thuringiensis isolates from Brazil for the presence of coleoptera-specific cry genes. Microbiological Research, 2000, 154, 355-362.	2.5	6
118	Molecular biology of baculovirus and its use in biological control in Brazil. Pesquisa Agropecuaria Brasileira, 1999, 34, 1733-1761.	0.9	8
119	Structural and ultrastructural studies of Anticarsia gemmatalis midgut cells infected with the baculovirus A. gemmatalis nucleopolyhedrovirus. Arthropod Structure and Development, 1999, 28, 195-201.	0.4	14
120	Characterization of <i>Bacillus thuringiensis</i> subsp. <i>kurstaki</i> strain S93 effective against the	0.8	8
121	Characterization of <i>Bacillus thuringiensis</i> subsp. <i>kurstaki</i> strain S93 effective against	0.8	1
122	Structural and Ultrastructural Changes during the Infection of UFL-AG-286 Cells with the Baculovirus AgMNPV. Journal of Invertebrate Pathology, 1998, 72, 239-245.	1.5	16
123	Expression of Full-Length and Truncated Forms of Crystal Protein Genes from Bacillus thuringiensis subsp. kurstaki in a Baculovirus and Pathogenicity of the Recombinant Viruses. Journal of Invertebrate Pathology, 1993, 62, 121-130.	1.5	34
124	Selection and characterization of Bacillus thuringiensis isolates with a high insecticidal activity against Spodoptera frugiperda (Lepidoptera: Noctuidae). Bioscience Journal, 0, , 1522-1536.	0.4	3