

# Vânia Aparecida Vicente

## List of Publications by Year in descending order

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123  
papers

3,641  
citations

126708

33  
h-index

155451

55  
g-index

125  
all docs

125  
docs citations

125  
times ranked

3569  
citing authors

#	ARTICLE	IF	CITATIONS
1	Chromoblastomycosis. <i>Clinical Microbiology Reviews</i> , 2017, 30, 233-276.	5.7	234
2	Waterborne <i>Exophiala</i> species causing disease in cold-blooded animals. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2011, 27, 46-72.	1.6	191
3	Proposed nomenclature for <i>Pseudallescheria</i> , <i>Scedosporium</i> and related genera. <i>Fungal Diversity</i> , 2014, 67, 1-10.	4.7	152
4	Exploring the genomic diversity of black yeasts and relatives ( <i>Chaetothyriales</i> , <i>Ascomycota</i> ). <i>Studies in Mycology</i> , 2017, 86, 1-28.	4.5	144
5	Species Diversity and Polymorphism in the <i>Exophiala spinifera</i> Clade Containing Opportunistic Black Yeast-Like Fungi. <i>Journal of Clinical Microbiology</i> , 2003, 41, 4767-4778.	1.8	141
6	Fungal infections in animals: a patchwork of different situations. <i>Medical Mycology</i> , 2018, 56, S165-S187.	0.3	141
7	Environmental isolation of black yeast-like fungi involved in human infection. <i>Studies in Mycology</i> , 2008, 61, 137-144.	4.5	136
8	Molecular ecology and pathogenic potential of <i>Fonsecaea</i> species. <i>Medical Mycology</i> , 2004, 42, 405-416.	0.3	126
9	The capability of endophytic fungi for production of hemicellulases and related enzymes. <i>BMC Biotechnology</i> , 2013, 13, 94.	1.7	89
10	<i>Fonsecaea nubica</i> sp. nov, a new agent of human chromoblastomycosis revealed using molecular data. <i>Medical Mycology</i> , 2010, 48, 800-806.	0.3	87
11	Shifts in taxonomic and functional microbial diversity with agriculture: How fragile is the Brazilian Cerrado?. <i>BMC Microbiology</i> , 2016, 16, 42.	1.3	78
12	Metagenomic analysis reveals microbial functional redundancies and specificities in a soil under different tillage and crop-management regimes. <i>Applied Soil Ecology</i> , 2015, 86, 106-112.	2.1	76
13	Black yeast-like fungi associated with Lethargic Crab Disease (LCD) in the mangrove-land crab, <i>Ucides cordatus</i> (Ocypodidae). <i>Veterinary Microbiology</i> , 2012, 158, 109-122.	0.8	71
14	Molecular Epidemiology of <i>Fonsecaea</i> Species. <i>Emerging Infectious Diseases</i> , 2011, 17, 464-469.	2.0	68
15	Molecular Epidemiology of Agents of Human Chromoblastomycosis in Brazil with the Description of Two Novel Species. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0005102.	1.3	66
16	Selective factors involved in oil flotation isolation of black yeasts from the environment. <i>Studies in Mycology</i> , 2008, 61, 157-163.	4.5	62
17	Cyphellophora and its relatives in Phialophora: biodiversity and possible role in human infection. <i>Fungal Diversity</i> , 2014, 65, 17-45.	4.7	62
18	<i>Fonsecaea pugnacius</i> , a Novel Agent of Disseminated Chromoblastomycosis. <i>Journal of Clinical Microbiology</i> , 2015, 53, 2674-2685.	1.8	62

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19	<i>Cladophialophora saturnica</i> sp. nov., a new opportunistic species of Chaetothyriales revealed using molecular data. <i>Medical Mycology</i> , 2009, 47, 51-62.	0.3	59
20	Propolis Extract for Onychomycosis Topical Treatment: From Bench to Clinic. <i>Frontiers in Microbiology</i> , 2018, 9, 779.	1.5	57
21	Environmental siblings of black agents of human chromoblastomycosis. <i>Fungal Diversity</i> , 2014, 65, 47-63.	4.7	56
22	Rapid detection of pathogenic fungi using loop-mediated isothermal amplification, exemplified by <i>Fonsecaea</i> agents of chromoblastomycosis. <i>Journal of Microbiological Methods</i> , 2010, 80, 19-24.	0.7	55
23	Antiadherent activity of <i>Schinus terebinthifolius</i> and <i>Croton urucurana</i> extracts on in vitro biofilm formation of <i>Candida albicans</i> and <i>Streptococcus mutans</i> . <i>Archives of Oral Biology</i> , 2014, 59, 887-896.	0.8	53
24	Analysis of the in vitro adherence of <i>Streptococcus mutans</i> and <i>Candida albicans</i> . <i>Brazilian Journal of Microbiology</i> , 2007, 38, 624-631.	0.8	50
25	A re-evaluation of the Chaetothyriales using criteria of comparative biology. <i>Fungal Diversity</i> , 2020, 103, 47-85.	4.7	43
26	Isolation of <i>Fonsecaea pedrosoi</i> from the Shell of the Babassu Coconut ( <i>Orbignya phalerata</i> Martius) in the Amazon Region of Maranhao Brazil. <i>Medical Mycology Journal</i> , 2006, 47, 305-311.	0.9	42
27	Rapid identification of fungal pathogens by rolling circle amplification using <i>Fonsecaea</i> as a model. <i>Mycoses</i> , 2011, 54, e577-82.	1.8	41
28	Histopathology of the mangrove land crab <i>Ucides cordatus</i> (Ocypodidae) affected by lethargic crab disease. <i>Diseases of Aquatic Organisms</i> , 2007, 78, 73-81.	0.5	41
29	The global burden of chromoblastomycosis. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009611.	1.3	40
30	<i>Fonsecaea multimorphosa</i> sp. nov, a new species of Chaetothyriales isolated from a feline cerebral abscess. <i>Fungal Biology</i> , 2011, 115, 1066-1076.	1.1	39
31	Genomic Understanding of an Infectious Brain Disease from the Desert. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 909-922.	0.8	39
32	Molecular identification of <i>Penicillium marneffeii</i> using rolling circle amplification. <i>Mycoses</i> , 2011, 54, e751-e759.	1.8	36
33	Molecular Epidemiology of <i>Fonsecaea</i> Species. <i>Emerging Infectious Diseases</i> , 2011, 17, 464-9.	2.0	35
34	The role of melanin pathways in extremotolerance and virulence of <i>Fonsecaea</i> revealed by <i>de novo</i> assembly transcriptomics using illumina paired-end sequencing. <i>Studies in Mycology</i> , 2016, 83, 1-18.	4.5	35
35	Comparative Genomics of Sibling Species of <i>Fonsecaea</i> Associated with Human Chromoblastomycosis. <i>Frontiers in Microbiology</i> , 2017, 8, 1924.	1.5	31
36	Susceptibility and molecular characterization of <i>Candida</i> species from patients with vulvovaginitis. <i>Brazilian Journal of Microbiology</i> , 2016, 47, 373-380.	0.8	30

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37	Molecular Characterization of Pathogenic Members of the Genus <i>Fonsecaea</i> Using Multilocus Analysis. <i>PLoS ONE</i> , 2012, 7, e41512.	1.1	28
38	Phylogenomic analyses reveal the diversity of laccase-coding genes in <i>Fonsecaea</i> genomes. <i>PLoS ONE</i> , 2017, 12, e0171291.	1.1	28
39	Black yeasts in the omics era: Achievements and challenges. <i>Medical Mycology</i> , 2018, 56, S32-S41.	0.3	28
40	Black Yeasts-Like Fungi Isolated from Dialysis Water in Hemodialysis Units. <i>Mycopathologia</i> , 2013, 175, 413-420.	1.3	27
41	Diversity of opportunistic black fungi on babassu coconut shells, a rich source of esters and hydrocarbons. <i>Fungal Biology</i> , 2017, 121, 488-500.	1.1	27
42	Molecular characterisation and antifungal susceptibility of clinical <i>Cryptococcus deuterogattii</i> (AFLP6/VGII) isolates from Southern Brazil. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2016, 35, 1803-1810.	1.3	24
43	Isolation of herpotrichiellaceous fungi from the environment. <i>Brazilian Journal of Microbiology</i> , 2001, 32, 47-51.	0.8	24
44	Molecular and morphological markers for rapid distinction between 2 <i>Colletotrichum</i> species. <i>Canadian Journal of Microbiology</i> , 2009, 55, 1076-1088.	0.8	22
45	Isolation and characterization of the nematophagous fungus <i>Arthrobotrys conoides</i> . <i>Parasitology Research</i> , 2013, 112, 177-185.	0.6	22
46	<i>Fusarium oxysporum</i> is an onychomycosis etiopathogenic agent. <i>Future Microbiology</i> , 2018, 13, 1745-1756.	1.0	22
47	Comparative Genomic Analysis of Capsule-Producing Black Yeasts <i>Exophiala dermatitidis</i> and <i>Exophiala spinifera</i> , Potential Agents of Disseminated Mycoses. <i>Frontiers in Microbiology</i> , 2020, 11, 586.	1.5	22
48	<i>Cladophialophora abundans</i> , a novel species of Chaetothyriales isolated from the natural environment. <i>Mycological Progress</i> , 2014, 13, 381-391.	0.5	21
49	A Model for Trans-Kingdom Pathogenicity in <i>Fonsecaea</i> Agents of Human Chromoblastomycosis. <i>Frontiers in Microbiology</i> , 2018, 9, 2211.	1.5	20
50	Fulfilling Koch's postulates confirms the mycotic origin of Lethargic Crab Disease. <i>Antonie Van Leeuwenhoek</i> , 2011, 99, 601-608.	0.7	19
51	Black Yeast Biota in the Mangrove, in Search of the Origin of the Lethargic Crab Disease (LCD). <i>Mycopathologia</i> , 2013, 175, 421-430.	1.3	19
52	Influence of Culturing Conditions on Bioprospecting and the Antimicrobial Potential of Endophytic Fungi from <i>Schinus terebinthifolius</i> . <i>Current Microbiology</i> , 2016, 72, 173-183.	1.0	18
53	<i>Arthrocladium</i> , an unexpected human opportunist in Trichomeriaceae (Chaetothyriales). <i>Fungal Biology</i> , 2016, 120, 207-218.	1.1	17
54	Genomic analysis of ant domatia-associated melanized fungi (Chaetothyriales, Ascomycota). <i>Mycological Progress</i> , 2019, 18, 541-552.	0.5	17

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55	Biological activity of <i>Diaporthe terebinthifolii</i> extracts against <i>Phyllosticta citricarpa</i> . FEMS Microbiology Letters, 2017, 364, .	0.7	16
56	Shed Light in the DaRk LineagES of the Fungal Tree of Lifeâ€™STRES. Life, 2020, 10, 362.	1.1	16
57	Bioprospecting highly diverse endophytic <i>Pestalotiopsis</i> spp. with antibacterial properties from <i>Maytenus ilicifolia</i> , a medicinal plant from Brazil. Canadian Journal of Microbiology, 2007, 53, 1123-1132.	0.8	15
58	Molecular characterization and antifungal susceptibility testing of <i>Cryptococcus neoformans sensu stricto</i> from southern Brazil. Journal of Medical Microbiology, 2018, 67, 560-569.	0.7	15
59	Molecular and Phenotypic Characterization of <i>Nannizzia</i> (Arthrodermataceae). Mycopathologia, 2020, 185, 9-35.	1.3	14
60	Control of pathogens in fresh pork sausage by inclusion of <i>Lactobacillus sakei</i> BASO117. Canadian Journal of Microbiology, 2019, 65, 831-841.	0.8	13
61	Molecular Identification and Antimicrobial Activity of Foliar Endophytic Fungi on the Brazilian Pepper Tree ( <i>Schinus terebinthifolius</i> ) Reveal New Species of <i>Diaporthe</i> . Current Microbiology, 2021, 78, 3218-3229.	1.0	13
62	Methodological variations in the isolation of genomic DNA from <i>Streptococcus</i> bacteria. Brazilian Archives of Biology and Technology, 2010, 53, 845-849.	0.5	12
63	<i>In Vitro</i> Activities of Eight Antifungal Drugs against 106 Waterborne and Cutaneous Exophiala Species. Antimicrobial Agents and Chemotherapy, 2013, 57, 6395-6398.	1.4	12
64	Molecular identification of <i>Histoplasma capsulatum</i> using rolling circle amplification. Mycoses, 2016, 59, 12-19.	1.8	12
65	In vitro susceptibility and molecular characterization of <i>Candida</i> spp. from candidemic patients. Revista Iberoamericana De Micologia, 2015, 32, 221-228.	0.4	11
66	Genetic manipulation of <i>Fonsecaea pedrosoi</i> using particles bombardment and <i>Agrobacterium</i> mediated transformation. Microbiological Research, 2018, 207, 269-279.	2.5	11
67	New Molecular Markers Distinguishing <i>Fonsecaea</i> Agents of Chromoblastomycosis. Mycopathologia, 2019, 184, 493-504.	1.3	11
68	Chromoblastomycosis in an Endemic Area of Brazil: A Clinical-Epidemiological Analysis and a Worldwide Haplotype Network. Journal of Fungi (Basel, Switzerland), 2020, 6, 204.	1.5	11
69	Environmental Detection of SARS-CoV-2 Virus RNA in Health Facilities in Brazil and a Systematic Review on Contamination Sources. International Journal of Environmental Research and Public Health, 2021, 18, 3824.	1.2	11
70	Black Fungi and Hydrocarbons: An Environmental Survey for Alkylbenzene Assimilation. Microorganisms, 2021, 9, 1008.	1.6	11
71	Microbiological and virulence aspects of. EXCLI Journal, 2020, 19, 687-704.	0.5	10
72	Onychomycosis by <i>Fusarium oxysporum</i> probably acquired in utero. Medical Mycology Case Reports, 2014, 6, 58-61.	0.7	9

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73	Draft Genome Sequence of the Ant-Associated Fungus <i>Phialophora attae</i> (CBS 131958). <i>Genome Announcements</i> , 2015, 3, .	0.8	9
74	Rapid Identification of Seven Waterborne <i>Exophiala</i> Species by RCA DNA Padlock Probes. <i>Mycopathologia</i> , 2018, 183, 669-677.	1.3	9
75	A case of disseminated sporotrichosis caused by <i>Sporothrix brasiliensis</i> . <i>Medical Mycology Case Reports</i> , 2018, 21, 34-36.	0.7	9
76	Environmental prospecting of black yeast-like agents of human disease using culture-independent methodology. <i>Scientific Reports</i> , 2020, 10, 14229.	1.6	9
77	Selective isolation of agents of chromoblastomycosis from insect-associated environmental sources. <i>Fungal Biology</i> , 2020, 124, 194-204.	1.1	9
78	Chromoblastomycosis Caused by <i>Phialophora</i> âProven Cases from Mexico. <i>Journal of Fungi (Basel)</i> , Tj ETQq0 0 0 rgBT /Overlock 10 Tf	1.5	9
79	Hypericin-P123-photodynamic therapy in an ex vivo model as an alternative treatment approach for onychomycosis caused by <i>Fusarium</i> spp.. <i>Photodiagnosis and Photodynamic Therapy</i> , 2021, 35, 102414.	1.3	9
80	Sporotrichosis in Children: Case series and Narrative Review. <i>Current Fungal Infection Reports</i> , 2022, 16, 33-46.	0.9	9
81	Some biomolecules and a partially O-acetylated exo-galactomannan containing Î²-Galf units from pathogenic <i>Exophiala jeanselmei</i> , having a pronounced immunogenic response. <i>International Journal of Biological Macromolecules</i> , 2011, 48, 177-182.	3.6	8
82	The bright future of darknessâthe rising power of black fungi: black yeasts, microcolonial fungi, and their relatives. <i>Mycopathologia</i> , 2013, 175, 365-368.	1.3	8
83	Resistance to Extended-Spectrum Î²-Lactamases in <i>Salmonella</i> from a Broiler Supply Chain. <i>International Journal of Environmental Research and Public Health</i> , 2014, 11, 11718-11726.	1.2	8
84	Specific primers for the detection of the black-yeast fungus associated with lethargic crab disease (LCD). <i>Diseases of Aquatic Organisms</i> , 2011, 94, 73-75.	0.5	8
85	Draft Genome Sequence of <i>Fonsecaea monophora</i> Strain CBS 269.37, an Agent of Human Chromoblastomycosis. <i>Genome Announcements</i> , 2016, 4, .	0.8	7
86	Comparative genomics of opportunistic <i>Phialophora</i> species involved in divergent disease types. <i>Mycoses</i> , 2021, 64, 555-568.	1.8	7
87	Draft Genome Sequence of <i>Fonsecaea nubica</i> Strain CBS 269.64, Causative Agent of Human Chromoblastomycosis. <i>Genome Announcements</i> , 2016, 4, .	0.8	6
88	Peritonitis by <i>Exophiala dermatitidis</i> in a pediatric patient. <i>Medical Mycology Case Reports</i> , 2019, 24, 18-22.	0.7	6
89	Comparative Analysis of Clinical and Environmental Strains of <i>Exophiala spinifera</i> by Long-Reads Sequencing and RNAseq Reveal Adaptive Strategies. <i>Frontiers in Microbiology</i> , 2020, 11, 1880.	1.5	6
90	Black fungi and ants: a genomic comparison of species inhabiting carton nests versus domatia. <i>IMA Fungus</i> , 2022, 13, 4.	1.7	6

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91	New method for early detection of two random amplified polymorphic DNA (RAPD) groups of <i>Staphylococcus aureus</i> causing bovine mastitis infection in Paran State, Brazil. <i>Brazilian Archives of Biology and Technology</i> , 2010, 53, 353-360.	0.5	5
92	Occurrence of sulphate reducing bacteria (SRB) associated with biocorrosion on metallic surfaces in a hydroelectric power station in Ibirama (SC) - Brazil. <i>Brazilian Archives of Biology and Technology</i> , 2013, 56, 801-809.	0.5	5
93	Is Marine Dispersion of the Lethargic Crab Disease Possible? Assessing the Tolerance of <i>Exophiala cancerae</i> to a Broad Combination of Salinities, Temperatures, and Exposure Times. <i>Mycopathologia</i> , 2017, 182, 997-1004.	1.3	5
94	Mixed secondary bacterial infection is associated with severe lesions of chromoblastomycosis in a neglected population from Brazil. <i>Diagnostic Microbiology and Infectious Disease</i> , 2019, 95, 201-207.	0.8	5
95	FATAL cryptococcal meningitis in a child with hyper-immunoglobulin M syndrome, with an emphasis on the agent. <i>Journal De Mycologie Medicale</i> , 2019, 29, 273-277.	0.7	5
96	Genomics and Virulence of <i>Fonsecaea pugnacius</i> , Agent of Disseminated Chromoblastomycosis. <i>Frontiers in Genetics</i> , 2020, 11, 822.	1.1	5
97	<i>Paecilomyces niveus</i> Stolk & Samson, 1971 (Ascomycota: Thermoascaceae) as a pathogen of <i>Nasonovia ribisnigri</i> (Mosley, 1841) (Hemiptera, Aphididae) in Brazil. <i>Brazilian Journal of Biology</i> , 2015, 75, 158-162.	0.4	5
98	A Case of Subcutaneous Phaeohyphomycosis Associated with Leprosy. <i>Infectious Disorders - Drug Targets</i> , 2017, 17, 223-226.	0.4	5
99	Vacuuming method as a successful strategy in the diagnosis of active infestation by <i>Pediculus humanus capitis</i> . <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2020, 62, e7.	0.5	5
100	Genetic variability of <i>Streptococcus mutans</i> isolated from low-income families, as shown by RAPD markers. <i>Brazilian Journal of Microbiology</i> , 2007, 38, 729-735.	0.8	4
101	Technological Potential of Antimicrobial Peptides: a Systematic Review. , 2019, 81, .		4
102	Pathogenicity and Growth Conditions Modulate <i>Fonsecaea</i> Extracellular Vesiclesâ€™ Ability to Interact With Macrophages. <i>Frontiers in Cellular and Infection Microbiology</i> , 0, 12, .	1.8	4
103	Glycan analysis of <i>Fonsecaea monophora</i> from clinical and environmental origins reveals different structural profile and human antigenic response. <i>Frontiers in Cellular and Infection Microbiology</i> , 2014, 4, 153.	1.8	3
104	Detection of <i>Streptococcus mutans</i> using padlock probe based on Rolling Circle Amplification (RCA). <i>Brazilian Archives of Biology and Technology</i> , 2015, 58, 54-60.	0.5	3
105	In vitro establishment of shoot meristems of <i>Ilex paraguariensis</i> and identification of endophytic bacteria. <i>Journal of Forestry Research</i> , 2019, 30, 1765-1777.	1.7	3
106	Genome Sequence of the Human Opportunistic Fungus <i>Arthrocladium fulminans</i> (CBS 136243). G3: Genes, Genomes, Genetics, 2020, 10, 1817-1821.	0.8	3
107	Environmental Screening of <i>Fonsecaea</i> Agents of Chromoblastomycosis Using Rolling Circle Amplification. <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 290.	1.5	3
108	In vitro activities of 8 antifungal drugs against 126 clinical and environmental <i>Exophiala</i> isolates. <i>Mycoses</i> , 2021, 64, 1328-1333.	1.8	3

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109	Genome Sequence of Type Strain <i>Fonsecaea multimorphosa</i> CBS 980.96 <sup>T</sup> , a Causal Agent of Feline Cerebral Phaeohyphomycosis. <i>Genome Announcements</i> , 2017, 5, .	0.8	2
110	Scalp microbiota alterations in children with pediculosis. <i>Infection, Genetics and Evolution</i> , 2019, 73, 322-331.	1.0	2
111	<i>Agrobacterium tumefaciens</i> -Mediated Transformation of <i>Fonsecaea monophora</i> and <i>Fonsecaea erecta</i> for Host-Environment Interaction Studies. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020, 6, 325.	1.5	2
112	Primary Central Nervous System Infection by <i>Histoplasma</i> in an Immunocompetent Adult. <i>Mycopathologia</i> , 2020, 185, 331-338.	1.3	2
113	An Atypical Etiology of Fungal Keratitis Caused by <i>Rousoella neopustulans</i> . <i>Journal of Fungi (Basel)</i> 11 0.784314 1.5 rgBT /Overlo	1.5	2
114	Shared Physiological Traits of <i>Exophiala</i> Species in Cold-Blooded Vertebrates, as Opportunistic Black Yeasts. <i>Mycopathologia</i> , 2016, 181, 353-362.	1.3	1
115	Lethargic Crab Disease: Now You See, Now You Donâ€™t. , 2018, , 233-247.		1
116	A Review on COVID-19 Diagnosis Tests Approved for Use in Brazil and the Impact on Pandemic Control. <i>Brazilian Archives of Biology and Technology</i> , 2021, 64, .	0.5	1
117	<i>In vitro</i> activity of eight antifungal drugs against <i>Chaetomiaceae</i> . <i>Medical Mycology</i> , 2021, 60, .	0.3	1
118	Chromoblastomycosis-Leprosy Co-Infection in Central West Brazil. Presentation of Three Cases and Literature Review. <i>Mycopathologia</i> , 2022, 187, 363-374.	1.3	1
119	Using molecular markers to assess <i>Streptococcus mutans</i> variability and the biological risk for caries. <i>Brazilian Journal of Oral Sciences</i> , 2014, 13, 235-241.	0.1	0
120	New perspectives on active pediculosis detection in schoolchildren from Southern Brazil. <i>Research, Society and Development</i> , 2021, 10, e58210615793.	0.0	0
121	Molecular characterization of <i>Streptococcus mutans</i> <i>gtfB</i> gene isolated from families. <i>Revista Odonto Ciencia</i> , 2018, 33, 40.	0.0	0
122	Unveiling Xylanolytic Enzymes Production of <i>Talaromyces wortmannii</i> DR49 on Industrial Agro Wastes. <i>Brazilian Archives of Biology and Technology</i> , 0, 64, .	0.5	0
123	New Insights on Environmental Occurrence of Pathogenic Fungi Based on Metagenomic Data from Brazilian Cerrado Biome. <i>Brazilian Archives of Biology and Technology</i> , 0, 65, .	0.5	0