## Vânia Aparecida Vicente

List of Publications by Year in descending order

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123 papers 3,641 citations

126708 33 h-index 55 g-index

125 all docs

125 docs citations

125 times ranked 3569 citing authors

#	Article	IF	Citations
1	Sporotrichosis in Children: Case series and Narrative Review. Current Fungal Infection Reports, 2022, 16, 33-46.	0.9	9
2	Black fungi and ants: a genomic comparison of species inhabiting carton nests versus domatia. IMA Fungus, 2022, 13, 4.	1.7	6
3	An Atypical Etiology of Fungal Keratitis Caused by Roussoella neopustulans. Journal of Fungi (Basel,) Tj ETQq1 1 (	0.784314 1.5	rgBT /Overloc
4	Chromoblastomycosis-Leprosy Co-Infection in Central West Brazil. Presentation of Three Cases and Literature Review. Mycopathologia, 2022, 187, 363-374.	1.3	1
5	Chromoblastomycosis Caused by Phialophora—Proven Cases from Mexico. Journal of Fungi (Basel,) Tj ETQq1 1	0.784314	4 rgBT /Over <mark>lo</mark>
6	A Review on COVID-19 Diagnosis Tests Approved for Use in Brazil and the Impact on Pandemic Control. Brazilian Archives of Biology and Technology, 2021, 64, .	0.5	1
7	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
8	Black Fungi and Hydrocarbons: An Environmental Survey for Alkylbenzene Assimilation. Microorganisms, 2021, 9, 1008.	1.6	11
9	New perspectives on active pediculosis detection in schoolchildren from Southern Brazil. Research, Society and Development, 2021, 10, e58210615793.	0.0	O
10	Molecular Identification and Antimicrobial Activity of Foliar Endophytic Fungi on the Brazilian Pepper Tree (Schinus terebinthifolius) Reveal New Species of Diaporthe. Current Microbiology, 2021, 78, 3218-3229.	1.0	13
11	In vitro activities of 8 antifungal drugs against 126 clinical and environmental <i>Exophiala</i> isolates. Mycoses, 2021, 64, 1328-1333.	1.8	3
12	The global burden of chromoblastomycosis. PLoS Neglected Tropical Diseases, 2021, 15, e0009611.	1.3	40
13	Hypericin-P123-photodynamic therapy in an ex vivo model as an alternative treatment approach for onychomycosis caused by Fusarium spp Photodiagnosis and Photodynamic Therapy, 2021, 35, 102414.	1.3	9
14	Comparative genomics of opportunistic <i>Phialophora</i> species involved in divergent disease types. Mycoses, 2021, 64, 555-568.	1.8	7
15	<i>In vitro</i> activity of eight antifungal drugs against <i>Chaetomiaceae</i> . Medical Mycology, 2021, 60, .	0.3	1
16	Molecular and Phenotypic Characterization of Nannizzia (Arthrodermataceae). Mycopathologia, 2020, 185, 9-35.	1.3	14
17	Genome Sequence of the Human Opportunistic Fungus Arthrocladium fulminans (CBS 136243). G3: Genes, Genomes, Genetics, 2020, 10, 1817-1821.	0.8	3
18	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

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19	Environmental Screening of Fonsecaea Agents of Chromoblastomycosis Using Rolling Circle Amplification. Journal of Fungi (Basel, Switzerland), 2020, 6, 290.	1.5	3
20	Agrobacterium tumefaciens-Mediated Transformation of Fonsecaea monophora and Fonsecaea erecta for Host-Environment Interaction Studies. Journal of Fungi (Basel, Switzerland), 2020, 6, 325.	1.5	2
21	Genomics and Virulence of Fonsecaea pugnacius, Agent of Disseminated Chromoblastomycosis. Frontiers in Genetics, 2020, 11, 822.	1.1	5
22	A re-evaluation of the Chaetothyriales using criteria of comparative biology. Fungal Diversity, 2020, 103, 47-85.	4.7	43
23	Environmental prospecting of black yeast-like agents of human disease using culture-independent methodology. Scientific Reports, 2020, 10, 14229.	1.6	9
24	Comparative Analysis of Clinical and Environmental Strains of Exophiala spinifera by Long-Reads Sequencing and RNAseq Reveal Adaptive Strategies. Frontiers in Microbiology, 2020, 11, 1880.	1.5	6
25	Shed Light in the DaRk LineagES of the Fungal Tree of Lifeâ€"STRES. Life, 2020, 10, 362.	1.1	16
26	Selective isolation of agents of chromoblastomycosis from insect-associated environmental sources. Fungal Biology, 2020, 124, 194-204.	1.1	9
27	Primary Central Nervous System Infection by Histoplasma in an Immunocompetent Adult. Mycopathologia, 2020, 185, 331-338.	1.3	2
28	Comparative Genomic Analysis of Capsule-Producing Black Yeasts Exophiala dermatitidis and Exophiala spinifera, Potential Agents of Disseminated Mycoses. Frontiers in Microbiology, 2020, 11, 586.	1.5	22
29	Vaccuuming method as a successful strategy in the diagnosis of active infestation by Pediculus humanus capitis. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2020, 62, e7.	0.5	5
30	Microbiological and virulence aspects of. EXCLI Journal, 2020, 19, 687-704.	0.5	10
31	In vitro establishment of shoot meristems of llex paraguariensis and identification of endophytic bacteria. Journal of Forestry Research, 2019, 30, 1765-1777.	1.7	3
32	Control of pathogens in fresh pork sausage by inclusion of <i>Lactobacillus sakei</i> BAS0117. Canadian Journal of Microbiology, 2019, 65, 831-841.	0.8	13
33	New Molecular Markers Distinguishing Fonsecaea Agents of Chromoblastomycosis. Mycopathologia, 2019, 184, 493-504.	1.3	11
34	Mixed secondary bacterial infection is associated with severe lesions of chromoblastomycosis in a neglected population from Brazil. Diagnostic Microbiology and Infectious Disease, 2019, 95, 201-207.	0.8	5
35	FATAL cryptococcal meningitis in a child with hyper-immunoglobulin M syndrome, with an emphasis on the agent. Journal De Mycologie Medicale, 2019, 29, 273-277.	0.7	5
36	Scalp microbiota alterations in children with pediculosis. Infection, Genetics and Evolution, 2019, 73, 322-331.	1.0	2

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37	Genomic analysis of ant domatia-associated melanized fungi (Chaetothyriales, Ascomycota). Mycological Progress, 2019, 18, 541-552.	0.5	17
38	Peritonitis by Exophiala dermatitidis in a pediatric patient. Medical Mycology Case Reports, 2019, 24, 18-22.	0.7	6
39	Technological Potential of Antimicrobial Peptides: a Systematic Review. , 2019, 81, .		4
40	Rapid Identification of Seven Waterborne Exophiala Species by RCA DNA Padlock Probes. Mycopathologia, 2018, 183, 669-677.	1.3	9
41	Black yeasts in the omics era: Achievements and challenges. Medical Mycology, 2018, 56, S32-S41.	0.3	28
42	A case of disseminated sporotrichosis caused by Sporothrix brasiliensis. Medical Mycology Case Reports, 2018, 21, 34-36.	0.7	9
43	Genetic manipulation of Fonsecaea pedrosoi using particles bombardment and Agrobacterium mediated transformation. Microbiological Research, 2018, 207, 269-279.	2.5	11
44	Genomic Understanding of an Infectious Brain Disease from the Desert. G3: Genes, Genomes, Genetics, 2018, 8, 909-922.	0.8	39
45	Fungal infections in animals: a patchwork of different situations. Medical Mycology, 2018, 56, S165-S187.	0.3	141
46	<i>Fusarium oxysporum</i> is an onychomycosis etiopathogenic agent. Future Microbiology, 2018, 13, 1745-1756.	1.0	22
47	A Model for Trans-Kingdom Pathogenicity in Fonsecaea Agents of Human Chromoblastomycosis. Frontiers in Microbiology, 2018, 9, 2211.	1.5	20
48	Propolis Extract for Onychomycosis Topical Treatment: From Bench to Clinic. Frontiers in Microbiology, 2018, 9, 779.	1.5	57
49	Lethargic Crab Disease: Now You See, Now You Don't. , 2018, , 233-247.		1
50	Molecular characterization and antifungal susceptibility testing of Cryptococcus neoformans sensu stricto from southern Brazil. Journal of Medical Microbiology, 2018, 67, 560-569.	0.7	15
51	Molecular characterization of Streptococcus mutans gtfB gene isolated from families. Revista Odonto Ciencia, 2018, 33, 40.	0.0	O
52	Exploring the genomic diversity of black yeasts and relatives ( <i>Chaetothyriales</i> , <i>Ascomycota</i> ). Studies in Mycology, 2017, 86, 1-28.	<b>4.</b> 5	144
53	Genome Sequence of Type Strain <i>Fonsecaea multimorphosa</i> CBS 980.96 <sup>T</sup> , a Causal Agent of Feline Cerebral Phaeohyphomycosis. Genome Announcements, 2017, 5, .	0.8	2
54	Diversity of opportunistic black fungi on babassu coconut shells, a rich source of esters and hydrocarbons. Fungal Biology, 2017, 121, 488-500.	1,1	27

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55	Biological activity of Diaporthe terebinthifolii extracts against Phyllosticta citricarpa. FEMS Microbiology Letters, 2017, 364, .	0.7	16
56	Is Marine Dispersion of the Lethargic Crab Disease Possible? Assessing the Tolerance of Exophiala cancerae to a Broad Combination of Salinities, Temperatures, and Exposure Times. Mycopathologia, 2017, 182, 997-1004.	1.3	5
57	Chromoblastomycosis. Clinical Microbiology Reviews, 2017, 30, 233-276.	5.7	234
58	Comparative Genomics of Sibling Species of Fonsecaea Associated with Human Chromoblastomycosis. Frontiers in Microbiology, 2017, 8, 1924.	1.5	31
59	Phylogenomic analyses reveal the diversity of laccase-coding genes in Fonsecaea genomes. PLoS ONE, 2017, 12, e0171291.	1.1	28
60	A Case of Subcutaneous Phaeohyphomycosis Associated with Leprosy. Infectious Disorders - Drug Targets, 2017, 17, 223-226.	0.4	5
61	Molecular Epidemiology of Agents of Human Chromoblastomycosis in Brazil with the Description of Two Novel Species. PLoS Neglected Tropical Diseases, 2016, 10, e0005102.	1.3	66
62	Shared Physiological Traits of Exophiala Species in Cold-Blooded Vertebrates, as Opportunistic Black Yeasts. Mycopathologia, 2016, 181, 353-362.	1.3	1
63	Arthrocladium, an unexpected human opportunist in Trichomeriaceae (Chaetothyriales). Fungal Biology, 2016, 120, 207-218.	1.1	17
64	Draft Genome Sequence of Fonsecaea monophora Strain CBS 269.37, an Agent of Human Chromoblastomycosis. Genome Announcements, 2016, 4, .	0.8	7
65	Molecular characterisation and antifungal susceptibility of clinical Cryptococcus deuterogattii (AFLP6/VGII) isolates from Southern Brazil. European Journal of Clinical Microbiology and Infectious Diseases, 2016, 35, 1803-1810.	1.3	24
66	Molecular identification of <i><scp>H</scp>istoplasma capsulatum</i> using rolling circle amplification. Mycoses, 2016, 59, 12-19.	1.8	12
67	Draft Genome Sequence of <i>Fonsecaea nubica</i> Strain CBS 269.64, Causative Agent of Human Chromoblastomycosis. Genome Announcements, 2016, 4, .	0.8	6
68	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. Studies in Mycology, 2016, 83, 1-18.	4.5	35
69	Shifts in taxonomic and functional microbial diversity with agriculture: How fragile is the Brazilian Cerrado?. BMC Microbiology, 2016, 16, 42.	1.3	78
70	Influence of Culturing Conditions on Bioprospecting and the Antimicrobial Potential of Endophytic Fungi from Schinus terebinthifolius. Current Microbiology, 2016, 72, 173-183.	1.0	18
71	Susceptibility and molecular characterization of Candida species from patients with vulvovaginitis. Brazilian Journal of Microbiology, 2016, 47, 373-380.	0.8	30
72	Detection of Streptococcus mutans using padlock probe based on Rolling Circle Amplification (RCA). Brazilian Archives of Biology and Technology, 2015, 58, 54-60.	0.5	3

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73	In vitro susceptibility and molecular characterization of Candida spp. from candidemic patients. Revista Iberoamericana De Micologia, 2015, 32, 221-228.	0.4	11
74	Fonsecaea pugnacius, a Novel Agent of Disseminated Chromoblastomycosis. Journal of Clinical Microbiology, 2015, 53, 2674-2685.	1.8	62
75	Draft Genome Sequence of the Ant-Associated Fungus Phialophora attae (CBS 131958). Genome Announcements, 2015, 3, .	0.8	9
76	Metagenomic analysis reveals microbial functional redundancies and specificities in a soil under different tillage and crop-management regimes. Applied Soil Ecology, 2015, 86, 106-112.	2.1	76
77	Paecilomyces niveus Stolk & Damson, 1971 (Ascomycota: Thermoascaceae) as a pathogen of Nasonovia ribisnigri (Mosley, 1841) (Hemiptera, Aphididae) in Brazil. Brazilian Journal of Biology, 2015, 75, 158-162.	0.4	5
78	Glycan analysis of Fonsecaea monophora from clinical and environmental origins reveals different structural profile and human antigenic response. Frontiers in Cellular and Infection Microbiology, 2014, 4, 153.	1.8	3
79	Resistance to Extended-Spectrum $\hat{l}^2$ -Lactamases in Salmonella from a Broiler Supply Chain. International Journal of Environmental Research and Public Health, 2014, 11, 11718-11726.	1.2	8
80	Using molecular markers to assess Streptococcus mutans variability and the biological risk for caries. Brazilian Journal of Oral Sciences, 2014, 13, 235-241.	0.1	0
81	Onychomycosis by Fusarium oxysporum probably acquired in utero. Medical Mycology Case Reports, 2014, 6, 58-61.	0.7	9
82	Antiadherent activity of Schinus terebinthifolius and Croton urucurana extracts on in vitro biofilm formation of Candida albicans and Streptococcus mutans. Archives of Oral Biology, 2014, 59, 887-896.	0.8	53
83	Environmental siblings of black agents of human chromoblastomycosis. Fungal Diversity, 2014, 65, 47-63.	4.7	56
84	Cyphellophora and its relatives in Phialophora: biodiversity and possible role in human infection. Fungal Diversity, 2014, 65, 17-45.	4.7	62
85	Cladophialophora abundans, a novel species of Chaetothyriales isolated from the natural environment. Mycological Progress, 2014, 13, 381-391.	0.5	21
86	Proposed nomenclature for Pseudallescheria, Scedosporium and related genera. Fungal Diversity, 2014, 67, 1-10.	4.7	152
87	The bright future of darkness—the rising power of black fungi: black yeasts, microcolonial fungi, and their relatives. Mycopathologia, 2013, 175, 365-368.	1.3	8
88	Isolation and characterization of the nematophagous fungus Arthrobotrys conoides. Parasitology Research, 2013, 112, 177-185.	0.6	22
89	Black Yeasts-Like Fungi Isolated from Dialysis Water in Hemodialysis Units. Mycopathologia, 2013, 175, 413-420.	1.3	27
90	Black Yeast Biota in the Mangrove, in Search of the Origin of the Lethargic Crab Disease (LCD). Mycopathologia, 2013, 175, 421-430.	1.3	19

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91	<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
92	The capability of endophytic fungi for production of hemicellulases and related enzymes. BMC Biotechnology, 2013, 13, 94.	1.7	89
93	Occurrence of sulphate reducing bacteria (SRB) associated with biocorrosion on metallic surfaces in a hydroelectric power station in Ibirama (SC) - Brazil. Brazilian Archives of Biology and Technology, 2013, 56, 801-809.	0.5	5
94	Molecular Characterization of Pathogenic Members of the Genus Fonsecaea Using Multilocus Analysis. PLoS ONE, 2012, 7, e41512.	1.1	28
95	Black yeast-like fungi associated with Lethargic Crab Disease (LCD) in the mangrove-land crab, Ucides cordatus (Ocypodidae). Veterinary Microbiology, 2012, 158, 109-122.	0.8	71
96	Fonsecaea multimorphosa sp. nov, a new species of Chaetothyriales isolated from a feline cerebral abscess. Fungal Biology, 2011, 115, 1066-1076.	1.1	39
97	Some biomolecules and a partially O-acetylated exo-galactomannan containing $\hat{l}^2$ -Galf units from pathogenic Exophiala jeanselmei, having a pronounced immunogenic response. International Journal of Biological Macromolecules, 2011, 48, 177-182.	3 <b>.</b> 6	8
98	Molecular Epidemiology of <i>Fonsecaea </i> Species. Emerging Infectious Diseases, 2011, 17, 464-469.	2.0	68
99	Waterborne <i>Exophiala</i> species causing disease in cold-blooded animals. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2011, 27, 46-72.	1.6	191
100	Rapid identification of fungal pathogens by rolling circle amplification using <i>Fonsecaea</i> as a model. Mycoses, 2011, 54, e577-82.	1.8	41
101	Molecular identification of Penicillium marneffei using rolling circle amplification. Mycoses, 2011, 54, e751-e759.	1.8	36
102	Fulfilling Koch's postulates confirms the mycotic origin of Lethargic Crab Disease. Antonie Van Leeuwenhoek, 2011, 99, 601-608.	0.7	19
103	Molecular Epidemiology of Fonsecaea Species. Emerging Infectious Diseases, 2011, 17, 464-9.	2.0	35
104	Specific primers for the detection of the black-yeast fungus associated with lethargic crab disease (LCD). Diseases of Aquatic Organisms, 2011, 94, 73-75.	0.5	8
105	New method for early detection of two random amplified polymorphic DNA (RAPD) groups of Staphylococcus aureus causing bovine mastitis infection in ParanA <sub>i</sub> State, Brazil. Brazilian Archives of Biology and Technology, 2010, 53, 353-360.	0.5	5
106	Methodological variations in the isolation of genomic DNA from Streptococcus bacteria. Brazilian Archives of Biology and Technology, 2010, 53, 845-849.	0.5	12
107	<i>Fonsecaea nubica</i> sp. nov, a new agent of human chromoblastomycosis revealed using molecular data. Medical Mycology, 2010, 48, 800-806.	0.3	87
108	Rapid detection of pathogenic fungi using loop-mediated isothermal amplification, exemplified by Fonsecaea agents of chromoblastomycosis. Journal of Microbiological Methods, 2010, 80, 19-24.	0.7	55

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109	Molecular and morphological markers for rapid distinction between 2 Colletotrichum species. Canadian Journal of Microbiology, 2009, 55, 1076-1088.	0.8	22
110	<i>Cladophialophorasaturnica</i> sp. nov., a new opportunistic species of <i>Chaetothyriales</i> revealed using molecular data. Medical Mycology, 2009, 47, 51-62.	0.3	59
111	Selective factors involved in oil flotation isolation of black yeasts from the environment. Studies in Mycology, 2008, 61, 157-163.	4.5	62
112	Environmental isolation of black yeast-like fungi involved in human infection. Studies in Mycology, 2008, 61, 137-144.	4.5	136
113	Bioprospecting highly diverse endophytic Pestalotiopsis spp. with antibacterial properties from Maytenus ilicifolia, a medicinal plant from Brazil. Canadian Journal of Microbiology, 2007, 53, 1123-1132.	0.8	15
114	Analysis of the in vitro adherence of Streptococcus mutans and Candida albicans. Brazilian Journal of Microbiology, 2007, 38, 624-631.	0.8	50
115	Genetic variability of Streptococcus mutans isolated from low-income families, as shown by RAPD markers. Brazilian Journal of Microbiology, 2007, 38, 729-735.	0.8	4
116	Histopathology of the mangrove land crab Ucides cordatus (Ocypodidae) affected by lethargic crab disease. Diseases of Aquatic Organisms, 2007, 78, 73-81.	0.5	41
117	Isolation of Fonsecaea pedrosoi from the Shell of the Babassu Coconut (Orbignya phalerata Martius) in the Amazon Region of Maranhao Brazil. Medical Mycology Journal, 2006, 47, 305-311.	0.9	42
118	Molecular ecology and pathogenic potential ofFonsecaeaspecies. Medical Mycology, 2004, 42, 405-416.	0.3	126
119	Species Diversity and Polymorphism in the Exophiala spinifera Clade Containing Opportunistic Black Yeast-Like Fungi. Journal of Clinical Microbiology, 2003, 41, 4767-4778.	1.8	141
120	Isolation of herpotrichiellacious fungi from the environment. Brazilian Journal of Microbiology, 2001, 32, 47-51.	0.8	24
121	Unveiling Xylanolytic Enzymes Production of Talaromyces wortmannii DR49 on Industrial Agro Wastes. Brazilian Archives of Biology and Technology, 0, 64, .	0.5	0
122	New Insights on Environmental Occurrence of Pathogenic Fungi Based on Metagenomic Data from Brazilian Cerrado Biome. Brazilian Archives of Biology and Technology, 0, 65, .	0.5	0
123	Pathogenicity and Growth Conditions Modulate Fonsecaea Extracellular Vesicles' Ability to Interact With Macrophages. Frontiers in Cellular and Infection Microbiology, 0, 12, .	1.8	4