Eliana Gertrudes de Macedo Lemos

List of Publications by Year
in descending order

139 papers	4,859 citations	236833 25 h-index	106281 65 g-index
140 all docs	140 docs citations	140 times ranked	5244 citing authors

#	Article	IF	Citations
1	Comparison of the genomes of two Xanthomonas pathogens with differing host specificities. Nature, 2002, 417, 459-463.	13.7	1,074
2	The genome sequence of the plant pathogen Xylella fastidiosa. Nature, 2000, 406, 151-157.	13.7	827
3	Comparative Genomics of Two Leptospira interrogans Serovars Reveals Novel Insights into Physiology and Pathogenesis. Journal of Bacteriology, 2004, 186, 2164-2172.	1.0	406
4	Comparative Analyses of the Complete Genome Sequences of Pierce's Disease and Citrus Variegated Chlorosis Strains of Xylella fastidiosa. Journal of Bacteriology, 2003, 185, 1018-1026.	1.0	307
5	Analysis and Functional Annotation of an Expressed Sequence Tag Collection for Tropical Crop Sugarcane. Genome Research, 2003, 13, 2725-2735.	2.4	254
6	The Genome Sequence of the Gram-Positive Sugarcane Pathogen Leifsonia xyli subsp. xyli. Molecular Plant-Microbe Interactions, 2004, 17, 827-836.	1.4	119
7	Brazilian coffee genome project: an EST-based genomic resource. Brazilian Journal of Plant Physiology, 2006, 18, 95-108.	0.5	112
8	Biotechnology of polyketides: new breath of life for the novel antibiotic genetic pathways discovery through metagenomics. Brazilian Journal of Microbiology, 2013, 44, 1007-1034.	0.8	67
9	Characterization of novel Acidobacteria exopolysaccharides with potential industrial and ecological applications. Scientific Reports, 2017, 7, 41193.	1.6	61
10	Genomics-based design of defined growth media for the plant pathogenXylella fastidiosa. FEMS Microbiology Letters, 2003, 219, 39-45.	0.7	59
11	Evaluation of the biotechnological potential of Rhizobium tropici strains for exopolysaccharide production. Carbohydrate Polymers, 2014, 111, 191-197.	5.1	54
12	Production of exopolysaccharide from rhizobia with potential biotechnological and bioremediation applications. International Journal of Biological Macromolecules, 2015, 74, 515-522.	3.6	53
13	Transcriptional Profiles of Roots of Different Soybean Genotypes Subjected to Drought Stress. Plant Molecular Biology Reporter, 2011, 29, 19-34.	1.0	52
14	An Evaluation of the Genetic Diversity of Xylella fastidiosa Isolated from Diseased Citrus and Coffee in SÃ£o Paulo, Brazil. Phytopathology, 2001, 91, 599-605.	1.1	48
15	The Family Rhizobiaceae. , 2014, , 419-437.		39
16	The Family Bradyrhizobiaceae. , 2014, , 135-154.		37
17	Characterization of new exopolysaccharide production by Rhizobium tropici during growth on hydrocarbon substrate. International Journal of Biological Macromolecules, 2017, 96, 361-369.	3.6	37
18	Cryopreservation of Dendrobium hybrid seeds and protocorms as affected by phloroglucinol and Supercool X1000. Scientia Horticulturae, 2012, 148, 154-160.	1.7	35

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19	Nitrogen assimilation in Citrus based on CitEST data mining. Genetics and Molecular Biology, 2007, 30, 810-818.	0.6	33
20	Molecular characterization of bacterial populations of different soils. Brazilian Journal of Microbiology, 2006, 37, 439-447.	0.8	32
21	Characterization of Exopolysaccharides Produced by Rhizobia Species. Revista Brasileira De Ciencia Do Solo, 2015, 39, 1566-1575.	0.5	31
22	From a metagenomic source to a high-resolution structure of a novel alkaline esterase. Applied Microbiology and Biotechnology, 2017, 101, 4935-4949.	1.7	31
23	Bacterial communities in mining soils and surrounding areas under regeneration process in a former ore mine. Brazilian Journal of Microbiology, 2018, 49, 489-502.	0.8	30
24	Diversidade genÃ©tica em maracujazeiro-amarelo utilizando marcadores moleculares fAFLP. Revista Brasileira De Fruticultura, 2004, 26, 494-498.	0.2	28
25	Identification of sex in Carica papaya L. using RAPD markers. Euphytica, 2002, 127, 179-184.	0.6	27
26	BactÃ©rias produtoras de auxinas isoladas de raÃzes de Cattleya walkeriana, orquÃdea Brasileira ameaÃ§ada de extinÃ§Ã£o, e sua funÃ§Ã£o na aclimatizaÃ§Ã£o. Revista Brasileira De Ciencia Do Solo, 2011, 35, 729-737.	0.5	27
27	Identification of reference genes for expression analysis by real-time quantitative PCR in drought-stressed soybean. Pesquisa Agropecuaria Brasileira, 2011, 46, 58-65.	0.9	27
28	Liming in the sugarcane burnt system and the green harvest practice affect soil bacterial community in northeastern SÃ£o Paulo, Brazil. Antonie Van Leeuwenhoek, 2016, 109, 1643-1654.	0.7	26
29	Est16, a New Esterase Isolated from a Metagenomic Library of a Microbial Consortium Specializing in Diesel Oil Degradation. PLoS ONE, 2015, 10, e0133723.	1.1	26
30	Bacterial diversity of soil under eucalyptus assessed by 16S rDNA sequencing analysis. Pesquisa Agropecuaria Brasileira, 2006, 41, 1507-1516.	0.9	24
31	Intense Exercise and Aerobic Conditioning Associated with Chromium or L-Carnitine Supplementation Modified the Fecal Microbiota of Fillies. PLoS ONE, 2016, 11, e0167108.	1.1	24
32	Chemical and rheological properties of exopolysaccharides produced by four isolates of rhizobia. International Journal of Biological Macromolecules, 2015, 81, 291-298.	3.6	23
33	Cryopreservation, early seedling development, and genetic stability of Oncidium flexuosum Sims. Plant Cell, Tissue and Organ Culture, 2013, 114, 139-148.	1.2	22
34	IdentificaÃ§Ã£o e avaliaÃ§Ã£o de rizobactÃ©rias isoladas de raÃzes de milho. Bragantia, 2010, 69, 905-911.	1.3	21
35	Identification and enzymatic characterization of acid phosphatase from Burkholderia gladioli. BMC Research Notes, 2014, 7, 221.	0.6	21
36	Optimized medium culture forAcidobacteriasubdivision 1 strains. FEMS Microbiology Letters, 2016, 363, fnw245.	0.7	21

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37	A nested-PCR assay for detection of Xylella fastidiosa in citrus plants and sharpshooter leafhoppers. Journal of Applied Microbiology, 2004, 96, 546-551.	1.4	20
38	Whole-Genome Expression Profiling ofXylella fastidiosain Response to Growth on Glucose. OMICS A Journal of Integrative Biology, 2005, 9, 77-90.	1.0	20
39	Draft Genome Sequence of the Nitrogen-Fixing Symbiotic Bacterium Bradyrhizobium elkanii 587. Journal of Bacteriology, 2012, 194, 3547-3548.	1.0	20
40	Validation of absolute quantitative real-time PCR for the diagnosis of Streptococcus agalactiae in fish. Journal of Microbiological Methods, 2015, 119, 168-175.	0.7	20
41	Genetic variations among passion fruit species using rapd markers. Revista Brasileira De Fruticultura, 2002, 24, 738-740.	0.2	19
42	Lack of host specificity of Colletotrichum spp. isolates associated with anthracnose symptoms on mango in Brazil. Plant Pathology, 2013, 62, 1038-1047.	1.2	19
43	ComposiÃ§Ã£o de exopolissacarÃdeos produzidos por estirpes de rizÃ³bios cultivados em diferentes fontes de carbono. Pesquisa Agropecuaria Brasileira, 2007, 42, 1503-1506.	0.9	18
44	Crescimento in vitro e aclimatizaÃ§Ã£o de Cattleya loddigesii Lindley (Orchidaceae) com carvÃ£o ativado sob dois espectros luminosos. Ciencia Rural, 2012, 42, 801-807.	0.3	18
45	Characterization of EST3: a metagenome-derived esterase with suitable properties for biotechnological applications. Applied Microbiology and Biotechnology, 2016, 100, 5815-5827.	1.7	18
46	Growth Optimization Procedures for the Phytopathogen Xylella fastidiosa. Current Microbiology, 2003, 46, 99-102.	1.0	17
47	A microsatellite library for Carica papaya L. cv. Sunrise solo. Revista Brasileira De Fruticultura, 2003, 25, 263-267.	0.2	17
48	Seedling Development and Evaluation of Genetic Stability of Cryopreserved Dendrobium Hybrid Mature Seeds. Applied Biochemistry and Biotechnology, 2014, 172, 2521-2529.	1.4	17
49	Bg10: A Novel Metagenomics Alcohol-Tolerant and Glucose-Stimulated GH1 ÃŸ-Glucosidase Suitable for Lactose-Free Milk Preparation. PLoS ONE, 2016, 11, e0167932.	1.1	17
50	Evaluation of the Genetic Diversity of Xylella fastidiosa Strains from Citrus and Coffee Hosts by Single-Nucleotide Polymorphism Markers. Phytopathology, 2007, 97, 1543-1549.	1.1	15
51	Uso de filtros de carvÃ£o ativado granular associado a microrganismos para remoÃ§Ã£o de fÃ¡rmacos no tratamento de Ã¡gua de abastecimento. Engenharia Sanitaria E Ambiental, 2016, 21, 709-720.	0.1	15
52	Molecular analysis of the bacterial diversity in a specialized consortium for diesel oil degradation. Revista Brasileira De Ciencia Do Solo, 2010, 34, 773-781.	0.5	14
53	Polyhydroxybutyrate in Rhizobium and Bradyrhizobium: quantification and phbC gene expression. World Journal of Microbiology and Biotechnology, 2011, 27, 773-778.	1.7	14
54	Exopolysaccharides produced by Pandoraea shows emulsifying and anti-biofilm activities. Journal of Polymer Research, 2019, 26, 1.	1.2	14

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55	CaracterizaÃ§Ã£o morfolÃ³gica e quÃmica de frutos de rambutan. Revista Brasileira De Fruticultura, 2008, 30, 958-963.	0.2	14
56	Î±-Hydroxynitrile lyase protein from Xylella fastidiosa: Cloning, expression, and characterization. Microbial Pathogenesis, 2009, 47, 118-127.	1.3	13
57	Influence of Vinasse Application in the Structure and Composition of the Bacterial Community of the Soil under Sugarcane Cultivation. International Journal of Microbiology, 2016, 2016, 1-11.	0.9	13
58	Draft Genome Sequence of a Chitinophaga Strain Isolated from a Lignocellulose Biomass-Degrading Consortium. Genome Announcements, 2017, 5, .	0.8	13
59	Desenvolvimento inicial e crescimento in vitro de Cattleya violacea (Kunth) Rolfe em diferentes concentraÃ§Ãµes de sacarose. Acta Amazonica, 2013, 43, 127-134.	0.3	13
60	CaracterizaÃ§Ã£o quÃmica dos gÃ©is produzidos pelas bactÃ©rias diazotrÃ³ficas Rhizobium tropici e Mesorhizobium sp.. Quimica Nova, 2012, 35, 705-708.	0.3	13
61	Mating system of a population of Myracrodruon urundeuva F.F. & M.F. AllemÃ£o using the fAFLP molecular marker. Genetics and Molecular Biology, 2004, 27, 425-431.	0.6	12
62	VariaÃ§Ã£o genÃ©tica em progÃªnies de Myracrodruon urundeuva F.F. & M.F. AllemÃ£o em trÃªs sistemas de cultivo. Revista Arvore, 2006, 30, 319-329.	0.5	12
63	ExpressÃ£o protÃ©ica diferencial entre plÃ¢ntulas apomÃticas e zigÃ³ticas de citros. Revista Brasileira De Fruticultura, 2004, 26, 1-4.	0.2	11
64	Propriedades reolÃ³gicas e efeito da adiÃ§Ã£o de sal na viscosidade de exopolissacarÃdeos produzidos por bactÃ©rias do gÃªnero Rhizobium. Quimica Nova, 2010, 33, 895-899.	0.3	11
65	Molecular and phylogenetic characterization based on the complete genome of a virulent pathotype of Newcastle disease virus isolated in the 1970s in Brazil. Infection, Genetics and Evolution, 2014, 26, 160-167.	1.0	11
66	Genetic Diversity and Population Differentiation of the Causal Agent of Citrus Black Spot in Brazil. Scientific World Journal, The, 2012, 2012, 1-14.	0.8	10
67	A kinetic model for Xylella fastidiosa adhesion, biofilm formation, and virulence. FEMS Microbiology Letters, 2004, 236, 313-318.	0.7	10
68	RelaÃ§Ãµes filogenÃ©ticas e diversidade de isolados de Guignardia spp oriundos de diferentes hospedeiros nas regiÃµes ITS1-5,8S-ITS2. Revista Brasileira De Fruticultura, 2009, 31, 360-380.	0.2	10
69	Molecular Identification of Fungal Communities in a Soil Cultivated with Vegetables and Soil Suppressiveness toRhizoctonia solani. Applied and Environmental Soil Science, 2013, 2013, 1-7.	0.8	9
70	Investigation, Expression, and Molecular Modeling of ORF2, a Metagenomic Lipolytic Enzyme. Applied Biochemistry and Biotechnology, 2015, 175, 3875-3887.	1.4	8
71	Identification of Microsatellite DNA Markers for the Giant Anteater Myrmecophaga tridactyla. Journal of Heredity, 2005, 96, 600-602.	1.0	7
72	Strain variability in the DNA immigration control region (ICR) of Xylella fastidiosa. Research in Microbiology, 2006, 157, 254-262.	1.0	7

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73	IdentificaÃ§Ã£o de hÃbridos de citros resistentes Ã mancha-marrom-de-alternÃ¡ria por meio de fAFLP e testes de patogenicidade. Pesquisa Agropecuaria Brasileira, 2007, 42, 975-983.	0.9	7
74	Impact of sewage sludge on the soil bacterial communities by DNA microarray analysis. World Journal of Microbiology and Biotechnology, 2011, 27, 1997-2003.	1.7	7
75	<b>Bacterial diversity in bovine rumen by metagenomic 16S rDNA sequencing and scanning electron microscopy. Acta Scientiarum - Animal Sciences, 2015, 37, 251.	0.3	7
76	Metagenomic Assembly and Draft Genome Sequence of an Uncharacterized Prevotella sp. from Nelore Rumen. Genome Announcements, 2015, 3, .	0.8	7
77	Properties of Polyhydroxyalkanoate Granules and Bioemulsifiers from Pseudomonas sp. and Burkholderia sp. Isolates Growing on Glucose. Applied Biochemistry and Biotechnology, 2016, 178, 990-1001.	1.4	7
78	Halotolerant aminopeptidase M29 from Mesorhizobium SEMIA 3007 with biotechnological potential and its impact on biofilm synthesis. Scientific Reports, 2017, 7, 10684.	1.6	7
79	Metagenome-assembled genome of a Chitinophaga sp. and its potential in plant biomass degradation, as well of affiliated Pandoraea and Labrys species. World Journal of Microbiology and Biotechnology, 2021, 37, 162.	1.7	7
80	Use of lipopolysaccharides to characterize Bradyrhizobium spp.. Soil Biology and Biochemistry, 1996, 28, 1227-1234.	4.2	6
81	Cloning, expression, purification and characterization of recombinant glutathione-S-transferase from Xylella fastidiosa. Protein Expression and Purification, 2008, 59, 153-160.	0.6	6
82	Kinetic characterization of a novel acid ectophosphatase from Enterobacter asburiae. Journal of Microbiology, 2016, 54, 106-113.	1.3	6
83	Seed cryopreservation, in vitro propagation and ex vitro growth of Cattleya walkeriana Gardner, a vulnerable ornamental orchid. Australian Journal of Crop Science, 2017, 11, 485-490.	0.1	6
84	Gene expression during the germination of coffee seed. Journal of Seed Science, 2019, 41, 168-179.	0.7	6
85	The potential of extracellular biopolymer production by Mesorhizobium sp. from monosaccharide constituents of lignocellulosic biomass. Biotechnology Letters, 2021, 43, 1385-1394.	1.1	6
86	UtilizaÃ§Ã£o de marcador molecular SCAR na identificaÃ§Ã£o de Fusarium subglutinans, agente causal da malformaÃ§Ã£o da mangueira. Revista Brasileira De Fruticultura, 2007, 29, 563-570.	0.2	6
87	Evaluation of Xylella fastidiosa genetic diversity by fAFPL markers. Revista Brasileira De Fruticultura, 2008, 30, 202-208.	0.2	6
88	Carbohydrate metabolism of Xylella fastidiosa: Detection of glycolytic and pentose phosphate pathway enzymes and cloning and expression of the enolase gene. Genetics and Molecular Biology, 2003, 26, 203-211.	0.6	5
89	Genetic characterization and nitrogen fixation capacity of Rhizobium strains on common bean. Pesquisa Agropecuaria Brasileira, 2008, 43, 1177-1184.	0.9	5
90	Genetic Diversity and Population Differentiation ofGuignardia mangiferaefrom â€œTahitiâ€•Acid Lime. Scientific World Journal, The, 2012, 2012, 1-11.	0.8	5

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91	Reclassification of the taxonomic status of SEMIA3007 isolated in Mexico B-11A Mex as Rhizobium leguminosarum bv. viceae by bioinformatic tools. BMC Microbiology, 2016, 16, 260.	1.3	5
92	Rhizobium spp exopolysaccharides production and xanthan lyase use on its structural modification. International Journal of Biological Macromolecules, 2019, 136, 424-435.	3.6	5
93	The Influence of Cooper and Chromium Ions on the Production of Exopolysaccharide and Polyhydroxybutyrate by Rhizobium tropici LBMP-C01. Journal of Polymers and the Environment, 2019, 27, 445-455.	2.4	5
94	Reconstruction and in silico analysis of new Marinobacter adhaerens t76_800 with potential for long-chain hydrocarbon bioremediation associated with marine environmental lipases. Marine Genomics, 2020, 49, 100685.	0.4	5
95	Rhizobial Exopolysaccharides and Type VI Secretion Systems: A Promising Way to Improve Nitrogen Acquisition by Legumes. Frontiers in Agronomy, 2021, 3, .	1.5	5
96	BiodegradaÃ§Ã£o da hepatotoxina(D-Leu1)-microcistina-LR por bactÃ©rias presentes em filtros biolÃ³gicos de carvÃ£o. Engenharia Sanitaria E Ambiental, 2013, 18, 205-214.	0.1	5
97	ConcentraÃ§Ãµes de sacarose no desenvolvimento in vitro e na aclimatizaÃ§Ã£o de Cattleya loddigesii Lindley. Semina:Ciencias Agrarias, 2013, 34, 583-592.	0.1	5
98	Bioremediation of heavy metalâ€“polluted environments by non-living cells from rhizobial isolates. Environmental Science and Pollution Research, 2022, 29, 46953-46967.	2.7	5
99	Marcadores fAFLP na caracterizaÃ§Ã£o de trÃªs genÃ³tipos de umezeiro selecionados como porta-enxertos para pessegueiro. Pesquisa Agropecuaria Brasileira, 2007, 42, 1741-1746.	0.9	4
100	Xylella fastidiosa gene expression analysis by DNA microarrays. Genetics and Molecular Biology, 2009, 32, 340-353.	0.6	4
101	CaracterizaÃ§Ã£o morfolÃ³gica de plantas de rambutan. Acta Scientiarum - Agronomy, 2009, 31, .	0.6	4
102	Xylella fastidiosa: An in vivo system to study possible survival strategies within citrus xylem vessels based on global gene expression analysis. Electronic Journal of Biotechnology, 2012, 15, .	1.2	4
103	Characterization of ruminal bacteria in grazing Nellore steers. Revista Colombiana De Ciencias Pecuarias, 2019, 32, 248-260.	0.4	4
104	DetecÃ§Ã£o de polimorfismo em porta-enxertos para citros. Revista Brasileira De Fruticultura, 2007, 29, 345-349.	0.2	4
105	SeleÃ§Ã£o de agentes alternativos ao Ã¡gar para propagaÃ§Ã£o de plÃ¢ntulas de Cattleya loddigesii Lindley (Orchidaceae). Revista Brasileirade Ciencias Agrarias, 2012, 7, 756-760.	0.3	4
106	Broad thermal spectrum metagenomic laccase with action for dye decolorization and fentin hydroxide treatment. AMB Express, 2022, 12, 38.	1.4	4
107	Resazurin reducing time as an indicator of Bradyrhizobium viable cell count. World Journal of Microbiology and Biotechnology, 1997, 14, 139-141.	1.7	3
108	Comportamento de cultivares de manga (Mangifera indica L.) em relaÃ§Ã£o Ã malformaÃ§Ã£o. Revista Brasileira De Fruticultura, 2007, 29, 115-119.	0.2	3

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109	CaracterizaÃ§Ã£o de rizÃ³bios indicados para produÃ§Ã£o de inoculantes por meio de sequenciamento parcial do 16S rRNA. Pesquisa Agropecuaria Brasileira, 2009, 44, 384-391.	0.9	3
110	Nitrogen Metabolism in Citrus Based on Expressed Tag Analysis. , 2012, , 245-255.		3
111	Erratum to â€œGenetic Diversity and Population Differentiation of Guignardia mangiferae from â€œTahitiâ€•Acid Limeâ€• Scientific World Journal, The, 2012, 2012, 1-1.	0.8	3
112	Differential Expression of Monocarboxylate Transporter 1 and Ancillary Protein CD147 in Red Blood Cells of Show Jumping Horses. Journal of Equine Veterinary Science, 2019, 81, 102791.	0.4	3
113	Production of DNA microarray and expression analysis of genes from Xylella fastidiosa in different culture media. Brazilian Archives of Biology and Technology, 2009, 52, 555-566.	0.5	3
114	Title is missing!. World Journal of Microbiology and Biotechnology, 1999, 15, 185-192.	1.7	2
115	AVALIAÃ‡Ãƒ O DE PLANTAS MATRIZES DE ABACAXIZEIRO CULTIVAR SMOOTH CAYENNE UTILIZANDO MARCADORES RAPD E PADRÃ•ES ISOENZIMÃTICOS. Revista Brasileira De Fruticultura, 2001, 23, 463-467.	0.2	2
116	Transposon Tn1721 distribution among strains ofXylella fastidiosa. FEMS Microbiology Letters, 2002, 208, 163-168.	0.7	2
117	AvaliaÃ§Ã£o de populaÃ§Ãµes de possÃveis rizobactÃ©rias em solos sob espÃ©cies florestais. Revista Brasileira De Ciencia Do Solo, 2008, 32, 1921-1927.	0.5	2
118	Discovery of genes related to antibiotic biosynthesis into a metagenomic library. New Biotechnology, 2009, 25, S105.	2.4	2
119	A putative twin-arginine translocation system in the phytopathogenic bacterium Xylella fastidiosa. Canadian Journal of Microbiology, 2011, 57, 149-154.	0.8	2
120	Exopolysaccharide Produced from Rhizobium spp. - An Interesting Product for Industry and Environment. Current Applied Polymer Science, 2020, 3, 157-166.	0.2	2
121	Bagasse minority pathway expression: Real time study of GH2 Î²-mannosidases from bacteroidetes. PLoS ONE, 2021, 16, e0247822.	1.1	2
122	ReaÃ§Ã£o enzimÃ¡tica ao estresse salino durante a germinaÃ§Ã£o de estilosantes. Pesquisa Agropecuaria Brasileira, 2004, 39, 93-95.	0.9	2
123	Molecular and Pathogenic Study of <i>Guignardia</i> spp. Isolates Associated to Different Hosts. Advances in Microbiology, 2014, 04, 116-125.	0.3	2
124	Electrophoretic patterns of lipopolysaccharides and antigenic analysis of soybean bradyrhizobia. World Journal of Microbiology and Biotechnology, 1999, 15, 205-215.	1.7	1
125	CaracterizaÃ§Ã£o molecular atravÃ©s da tÃ©cnica fAFLP de isolados de Diaporthe citri. Summa Phytopathologica, 2006, 32, 147-150.	0.3	1
126	Bacterial community in two subtropical fishponds in So Paulo, Brazil. African Journal of Microbiology Research, 2015, 9, 404-413.	0.4	1

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127	Carbohydrates on the growth of Cattleya schilleriana Reichb.f. seedlings (Orchidaceae). Ciencia Rural, 2018, 48, .	0.3	1
128	RemoÃ§Ã£o de herbicida atrazina por meio de filtros de carvÃ£o ativado granular associados com microrganismos no tratamento de Ã¡gua para abastecimento. Engenharia Sanitaria E Ambiental, 2021, 26, 263-272.	0.1	1
129	Transposon Tn1721 distribution among strains of Xylella fastidiosa. FEMS Microbiology Letters, 2002, 208, 163-168.	0.7	1
130	Algebraic approach to optimal clone selection applied in metagenomic projects. , 2010, , .		1
131	CaracterizaÃ§Ã£o de trÃªs genÃ³tipos de umezeiro (Prunus mume Sieb. et Zucc.) por marcadores RAPD. Revista Brasileira De Fruticultura, 2008, 30, 1045-1050.	0.2	1
132	CaracterizaÃ§Ã£o de rizÃ³bios (Bradyrhizobium japonicum) e produtividade da soja. Scientia Agricola, 1998, 55, 360-366.	0.6	1
133	Transcriptional Profile of Bradyrhizobium elkanii Semia 587 in Symbiosis with Soybean (Glycine max L.) Tj ETQq1 1 0.784314 rgBT /Ov 299-300.	0.0	1
134	Title is missing!. FEMS Microbiology Letters, 2004, 231, 299.	0.7	0
135	Microenxertia em cultivares de manga. Revista Brasileira De Fruticultura, 2006, 28, 533-535.	0.2	0
136	Diversidade genÃ©tica entre progÃªnies e matrizes de rambutan. Revista Brasileira De Fruticultura, 2012, 34, 630-634.	0.2	0
137	Differential Expression of Monocarboxylate Transporter 1 and Ancillary Protein CD147 in Red Blood Cells of Show Jumping Horses. Journal of Equine Veterinary Science, 2017, 50, 139-144.	0.4	0
138	Draft Genome Sequence of Bradyrhizobium elkanii Tn phoA 33, a Producer of Polyhydroxyalkanoates. Genome Announcements, 2017, 5, .	0.8	0
139	A Comprehensive Bioinformatics Analysis of the Lipoxygenases Superfamily in Shewanella Woodyi Strain (Strain ATCC 51908/MS32). Journal of Materials Science and Engineering B, 2014, 4, .	0.2	0