Aline Maria da Silva

List of Publications by Year in descending order

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82 papers

4,512 citations

32 h-index 63 g-index

86 all docs 86 docs citations

86 times ranked 5715 citing authors

#	Article	IF	CITATIONS
1	Characterization and (i) in vitro (/i) testing of newly isolated lytic bacteriophages for the Âbiocontrol of (i) Pseudomonas aeruginosa (/i). Future Microbiology, 2022, 17, 111-141.	2.0	7
2	Isolation and characterization of vB_XciM_LucasX, a new jumbo phage that infects Xanthomonas citri and Xanthomonas fuscans. PLoS ONE, 2022, 17, e0266891.	2.5	5
3	Comparative Genomics of Xylella fastidiosa Explores Candidate Host-Specificity Determinants and Expands the Known Repertoire of Mobile Genetic Elements and Immunity Systems. Microorganisms, 2022, 10, 914.	3.6	8
4	The XadA Trimeric Autotransporter Adhesins in <i>Xylella fastidiosa</i> Differentially Contribute to Cell Aggregation, Biofilm Formation, Insect Transmission and Virulence to Plants. Molecular Plant-Microbe Interactions, 2022, 35, 857-866.	2.6	3
5	Genomic Characterization of <i>Bacillus safensis</i> Isolated from Mine Tailings in Peru and Evaluation of Its Cyanide-Degrading Enzyme CynD. Applied and Environmental Microbiology, 2022, 88, .	3.1	3
6	Isolation and Molecular Characterization of a Novel Lytic Bacteriophage That Inactivates MDR Klebsiella pneumoniae Strains. Pharmaceutics, 2022, 14, 1421.	4. 5	13
7	A genomic catalog of Earth's microbiomes. Nature Biotechnology, 2021, 39, 499-509.	17.5	457
8	High-Quality Draft Genome Sequence of Pantanalinema sp. GBBB05, a Cyanobacterium From Cerrado Biome. Frontiers in Ecology and Evolution, 2021, 9, .	2.2	1
9	A comparative genomic analysis of Xanthomonas arboricola pv. juglandis strains reveal hallmarks of mobile genetic elements in the adaptation and accelerated evolution of virulence. Genomics, 2021, 113, 2513-2525.	2.9	9
10	Microbiomes of Field-Grown Maize and Soybean in Southeastern and Central Brazil Inferred by High-Throughput 16S and Internal Transcribed Spacer Amplicon Sequencing. Microbiology Resource Announcements, 2021, 10, e0052821.	0.6	1
11	Genome-resolved metagenome and metatranscriptome analyses of thermophilic composting reveal key bacterial players and their metabolic interactions. BMC Genomics, 2021, 22, 652.	2.8	16
12	Xylella fastidiosa subsp. pauca Strains Fb7 and 9a5c from Citrus Display Differential Behavior, Secretome, and Plant Virulence. International Journal of Molecular Sciences, 2020, 21, 6769.	4.1	6
13	High-Quality Draft Genome Sequence Resources of Eight Xylella fastidiosa Strains Isolated from Citrus, Coffee, Plum, and Hibiscus in South America. Phytopathology, 2020, 110, 1751-1755.	2.2	3
14	Genetic Diversity of Xylella fastidiosa Plasmids Assessed by Comparative Genomics. Tropical Plant Pathology, 2020, 45, 342-360.	1.5	8
15	Complete genome sequence and analysis of Alcaligenes faecalis strain Mc250, a new potential plant bioinoculant. PLoS ONE, 2020, 15, e0241546.	2.5	9
16	A Tropical Composting Operation Unit at São Paulo Zoo as a Source of Bacterial Proteolytic Enzymes. Applied Biochemistry and Biotechnology, 2019, 187, 282-297.	2.9	10
17	First report of cis-1,4-polyisoprene degradation by Gordonia paraffinivorans. Brazilian Journal of Microbiology, 2019, 50, 1051-1062.	2.0	9
18	A ligand motif enables differential vascular targeting of endothelial junctions between brain and retina. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 2300-2305.	7.1	14

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19	Characterization of novel hydrocarbon-degrading Gordonia paraffinivorans and Gordonia sihwensis strains isolated from composting. PLoS ONE, 2019, 14, e0215396.	2.5	19
20	Proteomic and Metabolomic Analyses of <i>Xylella fastidiosa</i> OMV-Enriched Fractions Reveal Association with Virulence Factors and Signaling Molecules of the DSF Family. Phytopathology, 2019, 109, 1344-1353.	2.2	51
21	Where do we aspire to publish? A position paper on scientific communication in biochemistry and molecular biology. Brazilian Journal of Medical and Biological Research, 2019, 52, e8935.	1.5	1
22	<i>Xanthomonas citri</i> T6SS mediates resistance to <i>Dictyostelium</i> predation and is regulated by an ECF \parallel f factor and cognate Ser/Thr kinase. Environmental Microbiology, 2018, 20, 1562-1575.	3.8	47
23	Comparative Metagenomics. Methods in Molecular Biology, 2018, 1704, 243-260.	0.9	2
24	MARVEL, a Tool for Prediction of Bacteriophage Sequences in Metagenomic Bins. Frontiers in Genetics, 2018, 9, 304.	2.3	133
25	Bacterial Diversification in the Light of the Interactions with Phages: The Genetic Symbionts and Their Role in Ecological Speciation. Frontiers in Ecology and Evolution, 2018, 6, .	2.2	32
26	Three novel Pseudomonas phages isolated from composting provide insights into the evolution and diversity of tailed phages. BMC Genomics, 2017, 18, 346.	2.8	32
27	Genome-Centric Analysis of a Thermophilic and Cellulolytic Bacterial Consortium Derived from Composting. Frontiers in Microbiology, 2017, 8, 644.	3.5	61
28	Plasmid-mediated mcr-1 in carbapenem-susceptible Escherichia coli ST156 causing a blood infection: an unnoticeable spread of colistin resistance in Brazil?. Clinics, 2017, 72, 642-644.	1.5	35
29	Characterization of mycobacteria and mycobacteriophages isolated from compost at the São Paulo Zoo Park Foundation in Brazil and creation of the new mycobacteriophage Cluster U. BMC Microbiology, 2016, 16, 111.	3.3	12
30	Microbial community structure and dynamics in thermophilic composting viewed through metagenomics and metatranscriptomics. Scientific Reports, 2016, 6, 38915.	3.3	183
31	<i>Xylella fastidiosa</i> outer membrane vesicles modulate plant colonization by blocking attachment to surfaces. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E3910-8.	7.1	143
32	Diffusible Signal Factor (DSF) Synthase RpfF of Xylella fastidiosa Is a Multifunction Protein Also Required for Response to DSF. Journal of Bacteriology, 2013, 195, 5273-5284.	2.2	41
33	Phenotype Overlap in Xylella fastidiosa Is Controlled by the Cyclic Di-GMP Phosphodiesterase Eal in Response to Antibiotic Exposure and Diffusible Signal Factor-Mediated Cell-Cell Signaling. Applied and Environmental Microbiology, 2013, 79, 3444-3454.	3.1	22
34	Metagenomic Analysis of a Tropical Composting Operation at the São Paulo Zoo Park Reveals Diversity of Biomass Degradation Functions and Organisms. PLoS ONE, 2013, 8, e61928.	2.5	91
35	Melatonin triggers PKA activation in the rodent malaria parasite Plasmodium chabaudi. Journal of Pineal Research, 2011, 50, 64-70.	7.4	35
36	Novel insights into the genomic basis of citrus canker based on the genome sequences of two strains of Xanthomonas fuscans subsp. aurantifolii. BMC Genomics, 2010, 11, 238.	2.8	102

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37	Development and validation of a Xanthomonas axonopodis pv. citri DNA microarray platform (XACarray) generated from the shotgun libraries previously used in the sequencing of this bacterial genome. BMC Research Notes, 2010, 3, 150.	1.4	3
38	Effects of the antimicrobial peptide gomesin on the global gene expression profile, virulence and biofilm formation of Xylella fastidiosa. FEMS Microbiology Letters, 2010, 306, 152-159.	1.8	25
39	Heterologous expression in Escherichia coli of Neurospora crassa neutral trehalase as an active enzyme. Protein Expression and Purification, 2009, 65, 185-189.	1.3	14
40	The P450 oxidoreductase, RedA, controls development beyond the mound stage in Dictyostelium discoideum. BMC Developmental Biology, 2008, 8, 8.	2.1	4
41	The Iron Stimulon of <i>Xylella fastidiosa</i> Includes Genes for Type IV Pilus and Colicin V-Like Bacteriocins. Journal of Bacteriology, 2008, 190, 2368-2378.	2.2	44
42	Identification and domain mapping of Dictyostelium discoideum type-1 protein phosphatase inhibitor-2â [†] t. Biochimie, 2007, 89, 692-701.	2.6	2
43	Genome mapping and expression analyses of human intronic noncoding RNAs reveal tissue-specific patterns and enrichment in genes related to regulation of transcription. Genome Biology, 2007, 8, R43.	9.6	209
44	Staurosporine induces tyrosine phosphorylation inDictyostelium discoideum proteins. Cell Biochemistry and Function, 2007, 25, 555-561.	2.9	0
45	Androgen responsive intronic non-coding RNAs. BMC Biology, 2007, 5, 4.	3.8	73
46	Transcription Profiling of Signal Transduction-Related Genes in Sugarcane Tissues. DNA Research, 2005, 12, 27-38.	3.4	77
47	Cyclic AMP and calcium interplay as second messengers in melatonin-dependent regulation of Plasmodium falciparum cell cycle. Journal of Cell Biology, 2005, 170, 551-557.	5.2	119
48	Large-scale Transcriptome Analyses Reveal New Genetic Marker Candidates of Head, Neck, and Thyroid Cancer. Cancer Research, 2005, 65, 1693-1699.	0.9	55
49	Whole-Genome Expression Profiling of Xylella fastidiosain Response to Growth on Glucose. OMICS A Journal of Integrative Biology, 2005, 9, 77-90.	2.0	20
50	DNA Microarray-Based Genome Comparison of a Pathogenic and a Nonpathogenic Strain of Xylella fastidiosa Delineates Genes Important for Bacterial Virulence. Journal of Bacteriology, 2004, 186, 5442-5449.	2.2	74
51	Polyductin, the PKHD1 gene product, comprises isoforms expressed in plasma membrane, primary cilium, and cytoplasm. Kidney International, 2004, 66, 1345-1355.	5.2	138
52	Antisense intronic non-coding RNA levels correlate to the degree of tumor differentiation in prostate cancer. Oncogene, 2004, 23, 6684-6692.	5.9	150
53	RASL11A, member of a novel small monomeric GTPase gene family, is down-regulated in prostate tumors. Biochemical and Biophysical Research Communications, 2004, 316, 618-627.	2.1	29
54	Analysis and Functional Annotation of an Expressed Sequence Tag Collection for Tropical Crop Sugarcane. Genome Research, 2003, 13, 2725-2735.	5.5	254

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55	Zerg: a very fast BLAST parser library. Bioinformatics, 2003, 19, 1035-1036.	4.1	17
56	The generation and utilization of a cancer-oriented representation of the human transcriptome by using expressed sequence tags. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 13418-13423.	7.1	105
57	ESTWeb: bioinformatics services for EST sequencing projects. Bioinformatics, 2003, 19, 1587-1588.	4.1	36
58	Dictyostelium discoideum protein phosphatase-1 catalytic subunit exhibits distinct biochemical properties. Biochemical Journal, 2003, 373, 703-711.	3.7	6
59	Genetic Organization of Plasmid pXF51 from the Plant Pathogen Xylella fastidiosa. Plasmid, 2001, 45, 184-199.	1.4	45
60	The contribution of 700,000 ORF sequence tags to the definition of the human transcriptome. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 12103-12108.	7.1	123
61	The sugarcane signal transduction (SUCAST) catalogue: prospecting signal transduction in sugarcane. Genetics and Molecular Biology, 2001, 24, 25-34.	1.3	26
62	The genome sequence of the plant pathogen Xylella fastidiosa. Nature, 2000, 406, 151-157.	27.8	827
63	Neutral trehalases catalyse intracellular trehalose breakdown in the filamentous fungi Aspergillus nidulans and Neurospora crassa. Molecular Microbiology, 1999, 32, 471-483.	2.5	101
64	Nitric oxide stimulates tyrosine phosphorylation in murine fibroblasts in the absence and presence of epidermal growth factor. Biochemical Journal, 1995, 305, 613-619.	3.7	55
65	Calcium uptake and gp80 messenger RNA destabilization follows cAMP receptor down regulation in Dictyostelium discoideum. Cellular Signalling, 1994, 6, 883-895.	3.6	2
66	BIOCHEMICAL AND FUNCTIONAL CHARACTERIZATION OF A GLYCOLIPID ANCHORED CELL ADHESION MOLECULE IN DICTYOSTELIUM DISCOIDEUM. , 1992 , , $211-228$.		0
67	Biochemical and functional characterization of a glycolipid anchored cell adhesion molecule in. Cell Biology International Reports, 1991, 15, 1065-1082.	0.6	2
68	A rapid posttranslational myristylation of a 68-kD protein in D. discoideum Journal of Cell Biology, 1990, 111, 401-407.	5.2	34
69	Cell adhesion in transformed D. discoideum cells: Expression of gp80 and its biochemical characterization. Developmental Biology, 1990, 140, 139-148.	2.0	22
70	Characterization of a glycosyl-phosphatidylinositol degrading activity in membranes. Experimental Cell Research, 1989, 185, 464-472.	2.6	14
71	Biosynthesis of 117 antigen: A cell cohesion molecule inDictyostelium discoideum. Genesis, 1988, 9, 561-567.	2.1	4
72	Regulation of tubulin and actin synthesis and accumulation during Blastocladiella emersonii development. Cell Differentiation, 1988, 24, 45-54.	0.4	4

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73	Evidence that a glycolipid tail anchors antigen 117 to the plasma membrane of Dictyostelium discoideum cells Proceedings of the National Academy of Sciences of the United States of America, 1988, 85, 5512-5515.	7.1	34
74	Acquisition of thermotolerance during development of Blastocladieila emersonii. Biochemical and Biophysical Research Communications, 1987, 144, 491-498.	2.1	4
75	Changes in the pattern of protein synthesis during zoospore germination in Blastocladiella emersonii. Journal of Bacteriology, 1987, 169, 2069-2078.	2.2	16
76	Effect of heat shock on S6 phosphorylation during the development of Blastocladiella emersonii. Molecular and Cellular Biochemistry, 1987, 78, 27-35.	3.1	1
77	Differential expression of heat-shock proteins and spontaneous synthesis of HSP70 during the life cycle of Blastocladiella emersonii. FEBS Journal, 1987, 163, 211-220.	0.2	27
78	Heat shock protein synthesis during development in Caulobacter crescentus. Journal of Bacteriology, 1986, 168, 923-930.	2.2	38
79	Developmental changes in translatable RNA species and protein synthesis during sporulation in the aquatic fungus Blastocladiella emersonii. Cell Differentiation, 1986, 18, 263-274.	0.4	16
80	Phosphorylation of ribosomal protein S6 in the aquatic fungus Blastocladiella emersonii. FEBS Journal, 1984, 144, 597-606.	0.2	16
81	CHAPTER 14: Iron as a Regulator of Virulence in Plant-Pathogenic Bacteria. , 0, , 263-283.		0
82	Diversity assessment of photosynthesizers: comparative analysis of pre-cultivated and natural microbiome of sediments from Cerrado biome in MaranhÃ \pm o, Brazil. Environmental Science and Pollution Research, 0, , .	5.3	0