John A Fuerst

List of Publications by Year in descending order

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0.0	0.200	76326	48315
98	8,309 citations	40	88
papers	citations	h-index	g-index
106	106	106	5790
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Missing lithotroph identified as new planctomycete. Nature, 1999, 400, 446-449.	27.8	1,382
2	Candidatus "Scalindua brodaeâ€, sp. nov., Candidatus "Scalindua wagneriâ€, sp. nov., Two New Species of Anaerobic Ammonium Oxidizing Bacteria. Systematic and Applied Microbiology, 2003, 26, 529-538.	2.8	535
3	Microbiology and application of the anaerobic ammonium oxidation (â€~anammox') process. Current Opinion in Biotechnology, 2001, 12, 283-288.	6.6	534
4	Candidatus "Anammoxoglobus propionicus―a new propionate oxidizing species of anaerobic ammonium oxidizing bacteria. Systematic and Applied Microbiology, 2007, 30, 39-49.	2.8	511
5	Beyond the bacterium: planctomycetes challenge our concepts of microbial structure and function. Nature Reviews Microbiology, 2011, 9, 403-413.	28.6	410
6	New concepts of microbial treatment processes for the nitrogen removal in wastewater. FEMS Microbiology Reviews, 2003, 27, 481-492.	8.6	407
7	Cell compartmentalisation in planctomycetes: novel types of structural organisation for the bacterial cell. Archives of Microbiology, 2001, 175, 413-429.	2.2	334
8	INTRACELLULAR COMPARTMENTATION IN PLANCTOMYCETES. Annual Review of Microbiology, 2005, 59, 299-328.	7.3	256
9	The anammoxosome: an intracytoplasmic compartment in anammox bacteria. FEMS Microbiology Letters, 2004, 233, 7-13.	1.8	243
10	Endocytosis-like protein uptake in the bacterium <i>Gemmata obscuriglobus</i> . Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12883-12888.	7.1	210
11	The occurrence of hopanoids in planctomycetes: implications for the sedimentary biomarker record. Organic Geochemistry, 2004, 35, 561-566.	1.8	179
12	Membrane-bounded nucleoid in the eubacterium Gemmatata obscuriglobus Proceedings of the National Academy of Sciences of the United States of America, 1991, 88, 8184-8188.	7.1	168
13	Linking Ultrastructure and Function in Four Genera of Anaerobic Ammonium-Oxidizing Bacteria: Cell Plan, Glycogen Storage, and Localization of Cytochrome <i>c</i> Proteins. Journal of Bacteriology, 2008, 190, 708-717.	2.2	163
14	Anaerobic ammonium oxidation by marine and freshwater planctomycete-like bacteria. Applied Microbiology and Biotechnology, 2003, 63, 107-114.	3.6	156
15	Discovery of a New Source of Rifamycin Antibiotics in Marine Sponge Actinobacteria by Phylogenetic Prediction. Applied and Environmental Microbiology, 2006, 72, 2118-2125.	3.1	128
16	Marine actinomycetes related to the 'Salinospora' group from the Great Barrier Reef sponge Pseudoceratina clavata. Environmental Microbiology, 2005, 7, 509-518.	3.8	123
17	Phylum Verrucomicrobia representatives share a compartmentalized cell plan with members of bacterial phylum Planctomycetes. BMC Microbiology, 2009, 9, 5.	3.3	120
18	Pirellulosomes: a new type of membrane-bounded cell compartment in planctomycete bacteria of the genus Pirellula. Microbiology (United Kingdom), 1997, 143, 739-748.	1.8	112

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19	Isolation of Gemmata -Like and Isosphaera -Like Planctomycete Bacteria from Soil and Freshwater. Applied and Environmental Microbiology, 2002, 68, 417-422.	3.1	110
20	A comparison of five methods for assaying bacterial hydrophobicity. Journal of Microbiological Methods, 1986, 6, 13-19.	1.6	95
21	Demonstration of lipopolysaccharide on sheathed flagella of Vibrio cholerae O:1 by protein A-gold immunoelectron microscopy. Journal of Bacteriology, 1988, 170, 1488-1494.	2.2	90
22	The anammox case-a new experimental manifesto for microbiological eco-physiology. Antonie Van Leeuwenhoek, 2002, 81, 693-702.	1.7	89
23	Improved nitrogen removal by application of new nitrogen-cycle bacteria. Reviews in Environmental Science and Biotechnology, 2002, 1, 51-63.	8.1	88
24	Isolation and molecular identification of planctomycete bacteria from postlarvae of the giant tiger prawn, Penaeus monodon. Applied and Environmental Microbiology, 1997, 63, 254-262.	3.1	84
25	Diversity of polyketide synthase genes from bacteria associated with the marine sponge <i>Pseudoceratina clavata</i> : cultureâ€dependent and cultureâ€independent approaches. Environmental Microbiology, 2006, 8, 1460-1470.	3.8	78
26	Intracellular localization of membraneâ€bound ATPases in the compartmentalized anammox bacterium â€~ <i>Candidatus</i> Kuenenia stuttgartiensis'. Molecular Microbiology, 2010, 77, 701-715.	2.5	71
27	Phenotypic conversion of Pseudomonas aeruginosa in cystic fibrosis. Journal of Clinical Microbiology, 1990, 28, 1143-1146.	3.9	68
28	Heterotrophic bacteria in an air-handling system. Applied and Environmental Microbiology, 1992, 58, 3914-3920.	3.1	67
29	The Order Planctomycetales, Including the Genera Planctomyces, Pirellula, Gemmata and Isosphaera and the Candidatus Genera Brocadia, Kuenenia and Scalindua. , 2006, , 757-793.		63
30	Phylogenetic Analysis of <i>Bradyrhizobium japonicum</i> and Photosynthetic Stem-Nodulating Bacteria from <i>Aeschynomene</i> Species Grown in Separated Geographical Regions. Applied and Environmental Microbiology, 1994, 60, 940-946.	3.1	63
31	The cell cycle of the planctomycete Gemmata obscuriglobus with respect to cell compartmentalization. BMC Cell Biology, 2009, 10, 4.	3.0	61
32	Widespread Distribution of Poribacteria in Demospongiae. Applied and Environmental Microbiology, 2009, 75, 5695-5699.	3.1	60
33	Culturable Bacterial Symbionts Isolated from Two Distinct Sponge Species (Pseudoceratina clavata) Tj ETQq1 Microbial Ecology, 2005, 50, 213-220.	1 0.784314 2.8	rgBT /Overlo 56
34	Cell division ring, a new cell division protein and vertical inheritance of a bacterial organelle in anammox planctomycetes. Molecular Microbiology, 2009, 73, 1009-1019.	2.5	53
35	Reclassification of the polyphyletic genus Prosthecomicrobium to form two novel genera, Vasilyevaea gen. nov. and Bauldia gen. nov. with four new combinations: Vasilyevaea enhydra comb. nov., Vasilyevaea mishustinii comb. nov., Bauldia consociata comb. nov. and Bauldia litoralis comb. nov International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 2960-2966.	1.7	52
36	Inhibition of growth ofLegionella species by heterotrophic plate count bacteria isolated from chlorinated drinking water. Current Microbiology, 1990, 21, 139-143.	2.2	48

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37	LC-MS-Based Metabolomics Study of Marine Bacterial Secondary Metabolite and Antibiotic Production in Salinispora arenicola. Marine Drugs, 2015, 13, 249-266.	4.6	45
38	The PVC superphylum: exceptions to the bacterial definition?. Antonie Van Leeuwenhoek, 2013, 104, 451-466.	1.7	44
39	Keys to eukaryality: Planctomycetes and ancestral evolution of cellular complexity. Frontiers in Microbiology, 2012, 3, 167.	3.5	43
40	Structural Studies of Planctomycete Gemmata obscuriglobus Support Cell Compartmentalisation in a Bacterium. PLoS ONE, 2014, 9, e91344.	2.5	42
41	Diversity and biotechnological potential of microorganisms associated with marine sponges. Applied Microbiology and Biotechnology, 2014, 98, 7331-7347.	3.6	41
42	Isolation of a bacterium resembling Pirellula species from primary tissue culture of the giant tiger prawn (Penaeus monodon). Applied and Environmental Microbiology, 1991, 57, 3127-3134.	3.1	36
43	Phylogenetic Analysis of Evolutionary Relationships of the Planctomycete Division of the Domain Bacteria Based on Amino Acid Sequences of Elongation Factor Tu. Journal of Molecular Evolution, 2001, 52, 405-418.	1.8	35
44	Isolation and diversity of planctomycetes from the sponge Niphates sp., seawater, and sediment of Moreton Bay, Australia. Antonie Van Leeuwenhoek, 2013, 104, 533-546.	1.7	35
45	Heterogeneity, persistence, and distribution of Pseudomonas aeruginosa genotypes in cystic fibrosis patients. Journal of Clinical Microbiology, 1991, 29, 2151-2157.	3.9	35
46	Protein uptake by bacteria. Communicative and Integrative Biology, 2010, 3, 572-575.	1.4	33
47	Discovering the Recondite Secondary Metabolome Spectrum of Salinispora Species: A Study of Inter-Species Diversity. PLoS ONE, 2014, 9, e91488.	2.5	33
48	Gene discovery within the planctomycete division of the domain Bacteria using sequence tags from genomic DNA libraries. Genome Biology, 2002, 3, research0031.1.	9.6	28
49	Effects of fixative and buffer on morphology and ultrastructure of a freshwater planctomycete, Gemmata obscuriglobus. Journal of Microbiological Methods, 1995, 21, 45-54.	1.6	27
50	Membrane-bounded nucleoids in microbial symbionts of marine sponges. FEMS Microbiology Letters, 1998, 166, 29-34.	1.8	27
51	A Phylogenetic Analysis of the Genus Blastobacter with a View to its Future Reclassification. Systematic and Applied Microbiology, 1994, 17, 51-57.	2.8	25
52	Novel protein domains and motifs in the marine planctomycete Rhodopirellula baltica. FEMS Microbiology Letters, 2004, 236, 333-340.	1.8	25
53	Tuwongella immobilis gen. nov., sp. nov., a novel non-motile bacterium within the phylum Planctomycetes. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 4923-4929.	1.7	25
54	Nuclear Pore-Like Structures in a Compartmentalized Bacterium. PLoS ONE, 2017, 12, e0169432.	2.5	24

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55	Diversity and distribution of the bioactive actinobacterial genus Salinispora from sponges along the Great Barrier Reef. Antonie Van Leeuwenhoek, 2012, 101, 603-618.	1.7	21
56	Production of <i>N</i> -acyl homoserine lactones by the sponge-associated marine actinobacteria <i>Salinispora arenicola</i> and <i>Salinispora pacifica</i> FEMS Microbiology Letters, 2017, 364, fnx002.	1.8	21
57	Diversity of Mycobacterium species from marine sponges and their sensitivity to antagonism by sponge-derived rifamycin-synthesizing actinobacterium in the genus Salinispora. FEMS Microbiology Letters, 2010, 313, 33-40.	1.8	20
58	Nested Bacterial Boxes: Nuclear and Other Intracellular Compartments in Planctomycetes. Journal of Molecular Microbiology and Biotechnology, 2013, 23, 95-103.	1.0	20
59	Effects of salinity on antibiotic production in sponge-derived <i>Salinispora</i> actinobacteria. Journal of Applied Microbiology, 2014, 117, 109-125.	3.1	19
60	A canonical FtsZ protein in Verrucomicrobium spinosum, a member of the Bacterial phylum Verrucomicrobia that also includes tubulin-producing Prosthecobacter species. BMC Evolutionary Biology, 2007, 7, 37.	3.2	18
61	Reanalysis of 5S rRNA sequence data for the Vibrionaceae with the clustan program suite. Current Microbiology, 1987, 15, 329-335.	2.2	17
62	Bacterial sheathed flagella and the rotary motor model for the mechanism of bacterial motility. Journal of Theoretical Biology, 1980, 84, 761-774.	1.7	16
63	Planctomycetesâ€"New Models for Microbial Cells and Activities. , 2017, , 1-27.		16
64	The Role of Reductionism in the Development of Molecular Biology: Peripheral or Central?. Social Studies of Science, 1982, 12, 241-278.	2.5	15
65	Electron tomography of the nucleoid of Gemmata obscuriglobus reveals complex liquid crystalline cholesteric structure. Frontiers in Microbiology, 2012, 3, 326.	3.5	15
66	Novel protein domains and motifs in the marine planctomycete Rhodopirellula baltica. FEMS Microbiology Letters, 2004, 236, 333-340.	1.8	15
67	Comparative analysis of ribonuclease P RNA of the planctomycetes. International Journal of Systematic and Evolutionary Microbiology, 2004, 54, 1333-1344.	1.7	14
68	Screening of rifamycin producing marine sponge bacteria by LC–MS–MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2007, 852, 362-366.	2.3	14
69	Bioinformatic analyses of integral membrane transport proteins encoded within the genome of the planctomycetes species, Rhodopirellula baltica. Biochimica Et Biophysica Acta - Biomembranes, 2014, 1838, 193-215.	2.6	14
70	The Definition of Molecular Biology and the Definition of Policy: The Role of the Rockefeller Foundation's Policy for Molecular Biology. Social Studies of Science, 1984, 14, 225-237.	2.5	13
71	Developmental cycle and pharmaceutically relevant compounds of Salinispora actinobacteria isolated from Great Barrier Reef marine sponges. Applied Microbiology and Biotechnology, 2013, 97, 3097-3108.	3.6	13
72	Molecular and ultrastructural confirmation of classification of ATCC 35122 as a strain of Pirellula staleyi International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 1663-1667.	1.7	13

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73	Anammoxosomes of Anaerobic Ammonium-oxidizing Planctomycetes. Microbiology Monographs, 2006, , 259-283.	0.6	10
74	Two Peptides, Cycloaspeptide A and Nazumamide A from a Sponge Associated Marine Actinobacterium <i>Salinispora</i> sp. Natural Product Communications, 2014, 9, 1934578X1400900.	0.5	10
75	Microorganisms Should Be High on DNA Preservation List. , 2000, 290, 1503b-1503.		10
76	Immersing undergraduate students in the research experience. Biochemistry and Molecular Biology Education, 2012, 40, 37-45.	1.2	9
77	Towards understanding the molecular mechanism of the endocytosis-like process in the bacterium Gemmata obscuriglobus. Biochimica Et Biophysica Acta - Molecular Cell Research, 2014, 1843, 1732-1738.	4.1	9
78	Ancient, highly conserved proteins from a LUCA with complex cell biology provide evidence in support of the nuclear compartment commonality (NuCom) hypothesis. Research in Microbiology, 2017, 168, 395-412.	2.1	9
79	Making heads or tails of the HU proteins in the planctomycete Gemmata obscuriglobus. Microbiology (United Kingdom), 2011, 157, 2012-2021.	1.8	8
80	Bacterial production of the fungusâ€derived cholesterolâ€lowering agent mevinolin. Biomedical Chromatography, 2014, 28, 1163-1166.	1.7	8
81	Buds from the tree of life: linking compartmentalized prokaryotes and eukaryotes by a non-hyperthermophile common ancestor and implications for understanding Archaean microbial communities. International Journal of Astrobiology, 2004, 3, 183-187.	1.6	7
82	Microbiological material exchanges among scientists. Research in Microbiology, 2010, 161, 446-452.	2.1	7
83	Two peptides, cycloaspeptide A and nazumamide A from a sponge associated marine actinobacterium Salinispora sp. Natural Product Communications, 2014, 9, 545-6.	0.5	7
84	Paralogization and New Protein Architectures in Planctomycetes Bacteria with Complex Cell Structures. Molecular Biology and Evolution, 2020, 37, 1020-1040.	8.9	6
85	Negative staining of freshwater bacterioneuston sampled directly with electron microscope specimen support grids. Microbial Ecology, 1987, 13, 219-228.	2.8	5
86	Close relationship of RNase P RNA in Gemmata and anammox planctomycete bacteria. FEMS Microbiology Letters, 2007, 268, 244-253.	1.8	5
87	Phylum Verrucomicrobia., 2019,, 551-551.		3
88	Membrane-bounded nucleoids in microbial symbionts of marine sponges. FEMS Microbiology Letters, 1998, 166, 29-34.	1.8	3
89	Membrane-bounded Nucleoids and Pirellulosomes of Planctomycetes. Microbiology Monographs, 2006, , 229-257.	0.6	2
90	Microorganisms—A Journal and a Unifying Concept for the Science of Microbiology. Microorganisms, 2014, 2, 140-146.	3.6	2

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91	Microorganismsâ€"A Forum for Understanding Microbial Life in All Its Forms. Microorganisms, 2013, 1, 1-2.	3.6	1
92	Novel Compartmentalistaion in Planctomycete Bacteria. Microscopy and Microanalysis, 2004, 10, 1528-1529.	0.4	0
93	Microbial Evolution: Chlamydial Creatures fromÂthe Deep. Current Biology, 2020, 30, R267-R269.	3.9	O
94	Editorial: Structure, Function and Evolution of Complex Cellular Organization in Bacteria and Archaea. Frontiers in Microbiology, 2021, 12, 751416.	3.5	0
95	Microbial diversity beyond E. coli: new microbial worlds, new concepts in biology. Microbiology Australia, 2011, 32, 73.	0.4	O
96	Planctomycetes: Their Evolutionary Implications for Models for Origins of Eukaryotes and the Eukaryote Nucleus and Endomembranes., 2013,, 243-270.		0
97	A Final Word: The Future of Planctomycetology and Related Studies. , 2013, , 271-273.		O
98	Cell Compartmentalization and Endocytosis in Planctomycetes: Structure and Function in Complex Bacteria., 2013,, 39-75.		0