

# Aurelien L Carlier

## List of Publications by Year in descending order

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44  
papers

1,968  
citations

218677

26  
h-index

265206

42  
g-index

50  
all docs

50  
docs citations

50  
times ranked

1946  
citing authors

#	ARTICLE	IF	CITATIONS
1	Induction of antibiotic specialized metabolism by co-culturing in a collection of phyllosphere bacteria. <i>Environmental Microbiology</i> , 2021, 23, 2132-2151.	3.8	12
2	<i>Paenibacillus foliorum</i> sp. nov., <i>Paenibacillus phytohabitans</i> sp. nov., <i>Paenibacillus plantarum</i> sp. nov., <i>Paenibacillus planticolens</i> sp. nov., <i>Paenibacillus phytorum</i> sp. nov. and <i>Paenibacillus germinis</i> sp. nov., isolated from the <i>Arabidopsis thaliana</i> phyllosphere. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	31
3	Patterns of transmission and horizontal gene transfer in the <i>Dioscorea sansibarensis</i> leaf symbiosis revealed by whole-genome sequencing. <i>Current Biology</i> , 2021, 31, 2666-2673.e4.	3.9	6
4	PaSiT: a novel approach based on short-oligonucleotide frequencies for efficient bacterial identification and typing. <i>Bioinformatics</i> , 2020, 36, 2337-2344.	4.1	5
5	<i>Orrella amnicola</i> sp. nov., isolated from a freshwater river, reclassification of <i>Algicoccus marinus</i> as <i>Orrella marina</i> comb. nov., and emended description of the genus <i>Orrella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 6381-6389.	1.7	13
6	Introducing SPeDE: High-Throughput Dereplication and Accurate Determination of Microbial Diversity from Matrix-Assisted Laser Desorption/Ionization Time of Flight Mass Spectrometry Data. <i>MSystems</i> , 2019, 4, .	3.8	53
7	Adaptations and evolution of a heritable leaf nodule symbiosis between <i>Dioscorea sansibarensis</i> and <i>Orrella dioscoreae</i> . <i>ISME Journal</i> , 2019, 13, 1831-1844.	9.8	17
8	Synthesis and Biological Evaluation of the Novel Growth Inhibitor Streptol Glucoside, Isolated from an Obligate Plant Symbiont. <i>Chemistry - A European Journal</i> , 2019, 25, 1722-1726.	3.3	13
9	<i>Pedobacter gandavensis</i> sp. nov., <i>Pedobacter foliorum</i> sp. nov. and <i>Pedobacter planticolens</i> sp. nov., isolated from leaves of <i>Arabidopsis thaliana</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 71, .	1.7	16
10	Leaf nodule symbiosis: function and transmission of obligate bacterial endophytes. <i>Current Opinion in Plant Biology</i> , 2018, 44, 23-31.	7.1	46
11	<i>Abditibacterium utsteinense</i> sp. nov., the first cultivated member of candidate phylum FBP, isolated from ice-free Antarctic soil samples. <i>Systematic and Applied Microbiology</i> , 2018, 41, 279-290.	2.8	58
12	Heterologous Expression, Biosynthetic Studies, and Ecological Function of the Selective Gq-Signaling Inhibitor FR900359. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 836-840.	13.8	57
13	Heterologe Expression, Biosynthese und Ökologische Funktion des selektiven Gq-Signaltransduktionsinhibitors FR900359. <i>Angewandte Chemie</i> , 2018, 130, 844-849.	2.0	5
14	Patterns of Nucleotide Deletion and Insertion Inferred from Bacterial Pseudogenes. <i>Genome Biology and Evolution</i> , 2018, 10, 1792-1802.	2.5	12
15	Draft genome and description of <i>Orrella dioscoreae</i> gen. nov. sp. nov., a new species of <i>Alcaligenaceae</i> isolated from leaf acumens of <i>Dioscorea sansibarensis</i> . <i>Systematic and Applied Microbiology</i> , 2017, 40, 11-21.	2.8	42
16	The Essential Genome of <i>Burkholderia cenocepacia</i> H111. <i>Journal of Bacteriology</i> , 2017, 199, .	2.2	24
17	Comparative Genomics of <i>Burkholderia singularis</i> sp. nov., a Low G+C Content, Free-Living Bacterium That Defies Taxonomic Dissection of the Genus <i>Burkholderia</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 1679.	3.5	36
18	Characterization of the emerging zoonotic pathogen <i>Arcobacter thereius</i> by whole genome sequencing and comparative genomics. <i>PLoS ONE</i> , 2017, 12, e0180493.	2.5	21

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19	Comparative genomics of <i>Burkholderia multivorans</i> , a ubiquitous pathogen with a highly conserved genomic structure. <i>PLoS ONE</i> , 2017, 12, e0176191.	2.5	17
20	The genome analysis of <i>Candidatus Burkholderia crenata</i> reveals that secondary metabolism may be a key function of the <i>Acridia crenata</i> leaf nodule symbiosis. <i>Environmental Microbiology</i> , 2016, 18, 2507-2522.	3.8	64
21	The role of siderophores in metal homeostasis of members of the genus <i>Burkholderia</i> . <i>Environmental Microbiology Reports</i> , 2016, 8, 103-109.	2.4	17
22	Molecular mechanisms underlying the close association between soil <i>Burkholderia</i> and fungi. <i>ISME Journal</i> , 2016, 10, 253-264.	9.8	118
23	Evidence of horizontal gene transfer between obligate leaf nodule symbionts. <i>ISME Journal</i> , 2016, 10, 2092-2105.	9.8	63
24	Isolation and Total Synthesis of Kirkamide, an Aminocyclitol from an Obligate Leaf Nodule Symbiont. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 7968-7970.	13.8	44
25	Isolation and Total Synthesis of Kirkamide, an Aminocyclitol from an Obligate Leaf Nodule Symbiont. <i>Angewandte Chemie</i> , 2015, 127, 8079-8081.	2.0	10
26	Microbial Biofilms and Quorum Sensing. , 2015, , 45-52.		4
27	Genome Sequence of <i>Burkholderia cenocepacia</i> H111, a Cystic Fibrosis Airway Isolate. <i>Genome Announcements</i> , 2014, 2, .	0.8	39
28	A novel siderophore-independent strategy of iron uptake in the genus <i>Burkholderia</i> . <i>Molecular Microbiology</i> , 2014, 91, 805-820.	2.5	46
29	The Third Replicon of Members of the <i>Burkholderia cepacia</i> Complex, Plasmid pC3, Plays a Role in Stress Tolerance. <i>Applied and Environmental Microbiology</i> , 2014, 80, 1340-1348.	3.1	33
30	Proteomics Analysis of <i>Psychotria</i> Leaf Nodule Symbiosis: Improved Genome Annotation and Metabolic Predictions. <i>Molecular Plant-Microbe Interactions</i> , 2013, 26, 1325-1333.	2.6	27
31	Identification of <i>Burkholderia cenocepacia</i> Strain H111 Virulence Factors Using Nonmammalian Infection Hosts. <i>Infection and Immunity</i> , 2013, 81, 143-153.	2.2	40
32	The AHL- and BDSF-Dependent Quorum Sensing Systems Control Specific and Overlapping Sets of Genes in <i>Burkholderia cenocepacia</i> H111. <i>PLoS ONE</i> , 2012, 7, e49966.	2.5	70
33	Exposing the third chromosome of <i>Burkholderia cepacia</i> complex strains as a virulence plasmid. <i>Molecular Microbiology</i> , 2012, 83, 362-378.	2.5	90
34	The eroded genome of a <i>Psychotria</i> leaf symbiont: hypotheses about lifestyle and interactions with its plant host. <i>Environmental Microbiology</i> , 2012, 14, 2757-2769.	3.8	60
35	Structure/Function Analysis of the <i>Pantoea stewartii</i> Quorum-Sensing Regulator EsaR as an Activator of Transcription. <i>Journal of Bacteriology</i> , 2009, 191, 7402-7409.	2.2	31
36	Identification and characterization of three novel EsaI/EsaR quorum-sensing controlled stewartan exopolysaccharide biosynthetic genes in <i>Pantoea stewartii</i> ssp. <i>stewartii</i> . <i>Molecular Microbiology</i> , 2009, 74, 903-913.	2.5	28

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37	The rcsA Promoter of <i>Pantoea stewartii</i> subsp. <i>stewartii</i> Features a Low-Level Constitutive Promoter and an EsaR Quorum-Sensing-Regulated Promoter. <i>Journal of Bacteriology</i> , 2006, 188, 4581-4584.	2.2	20
38	N-hexanoyl-L-homoserine lactone, a mediator of bacterial quorum-sensing regulation, exhibits plant-dependent stability and may be inactivated by germinating <i>Lotus corniculatus</i> seedlings. <i>FEMS Microbiology Ecology</i> , 2005, 52, 13-20.	2.7	107
39	The cell density-dependent expression of stewartan exopolysaccharide in <i>Pantoea stewartii</i> ssp. <i>stewartii</i> is a function of EsaR-mediated repression of the rcsA gene. <i>Molecular Microbiology</i> , 2005, 56, 189-203.	2.5	64
40	Bacterial populations in the rhizosphere of tobacco plants producing the quorum-sensing signals hexanoyl-homoserine lactone and 3-oxo-hexanoyl-homoserine lactone. <i>FEMS Microbiology Ecology</i> , 2004, 51, 19-29.	2.7	34
41	The Assimilation of $\hat{1}^3$ -Butyrolactone in <i>Agrobacterium tumefaciens</i> C58 Interferes with the Accumulation of the N-Acyl-Homoserine Lactone Signal. <i>Molecular Plant-Microbe Interactions</i> , 2004, 17, 951-957.	2.6	69
42	The Ti Plasmid of <i>Agrobacterium tumefaciens</i> Harbors an attM-Paralogous Gene, <i>aiiB</i> , Also Encoding N-Acyl Homoserine Lactonase Activity. <i>Applied and Environmental Microbiology</i> , 2003, 69, 4989-4993.	3.1	189
43	Novel bacteria degrading N-acylhomoserine lactones and their use as quenchers of quorum-sensing-regulated functions of plant-pathogenic bacteria. <i>Microbiology (United Kingdom)</i> , 2003, 149, 1981-1989.	1.8	213
44	Role of Quorum-Sensing Regulation in Pathogenesis of <i>Pantoea stewartii</i> subsp. <i>stewartii</i> . , 0, , 201-212.		0