Fabien Aujoulat

List of Publications by Year in descending order

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448610 511568 1,115 31 19 30 citations h-index g-index papers 31 31 31 1823 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A new perspective on opportunistic pathogens of the genus Bordetella in cystic fibrosis. Journal of Cystic Fibrosis, 2022, 21, 344-347.	0.3	2
2	From Copper Tolerance to Resistance in Pseudomonas aeruginosa towards Patho-Adaptation and Hospital Success. Genes, 2022, 13, 301.	1.0	18
3	Environmental Antimicrobial Resistance in a Small Urban Mediterranean River: A Focus on Endemic Beta-Lactamases in Clinically Relevant Bacteria. Water (Switzerland), 2021, 13, 2010.	1.2	3
4	The population structure of Ochrobactrum isolated from entomopathogenic nematodes indicates interactions with the symbiotic system. Infection, Genetics and Evolution, 2019, 70, 131-139.	1.0	4
5	Whole-Genome Sequences of a Cluster of 14 Unidentified Related Veillonella sp. Strains from Human Clinical Samples and Type Strains of 3 Veillonella Validated Species. Microbiology Resource Announcements, 2019, 8, .	0.3	O
6	A sequence database analysis of 5-nitroimidazole reductase and related proteins to expand knowledge on enzymes responsible for metronidazole inactivation. Anaerobe, 2019, 55, 29-34.	1.0	4
7	Survey of Staphylococcus aureus in a general pediatric population and focus on isolates with three clinically relevant toxin-encoding genes. World Journal of Pediatrics, 2018, 14, 35-43.	0.8	5
8	Inefficient immune response is associated with microbial permissiveness in juvenile oysters affected by mass mortalities on field. Fish and Shellfish Immunology, 2018, 77, 156-163.	1.6	32
9	First molecular characterization of related cases of healthcare-associated infections involving multidrug-resistant Enterococcus faecium vanA in Algeria. Infection and Drug Resistance, 2018, Volume 11, 1483-1490.	1.1	10
10	Chronic Airway Colonization by Achromobacter xylosoxidans in Cystic Fibrosis Patients Is Not Sustained by Their Domestic Environment. Applied and Environmental Microbiology, 2018, 84, .	1.4	7
11	A new methodology to assess antimicrobial resistance of bacteria in coastal waters; pilot study in a Mediterranean hydrosystem. Comptes Rendus - Geoscience, 2017, 349, 310-318.	0.4	3
12	Highly Diversified Pandoraea pulmonicola Population during Chronic Colonization in Cystic Fibrosis. Frontiers in Microbiology, 2017, 8, 1892.	1.5	19
13	Highly diverse recombining populations of Vibrio cholerae and Vibrio parahaemolyticus in French Mediterranean coastal lagoons. Frontiers in Microbiology, 2015, 6, 708.	1.5	21
14	Bacterial Diversity Associated with Wild Caught Anopheles Mosquitoes from Dak Nong Province, Vietnam Using Culture and DNA Fingerprint. PLoS ONE, 2015, 10, e0118634.	1.1	24
15	Pseudomonas aeruginosa and Achromobacter sp. Clonal Selection Leads to Successive Waves of Contamination of Water in Dental Care Units. Applied and Environmental Microbiology, 2015, 81, 7509-7524.	1.4	30
16	Rarimicrobium hominis gen. nov., sp. nov., representing the fifth genus in the phylum Synergistetes that includes human clinical isolates. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 3965-3970.	0.8	12
17	Temporal dynamics of the very premature infant gut dominant microbiota. BMC Microbiology, 2014, 14, 325.	1.3	41
18	Veillonella seminalis sp. nov., a novel anaerobic Gram-stain-negative coccus from human clinical samples, and emended description of the genus Veillonella. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 3526-3531.	0.8	39

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19	Niches, Population Structure and Genome Reduction in Ochrobactrum intermedium: Clues to Technology-Driven Emergence of Pathogens. PLoS ONE, 2014, 9, e83376.	1.1	45
20	From Environment to Man: Genome Evolution and Adaptation of Human Opportunistic Bacterial Pathogens. Genes, 2012, 3, 191-232.	1.0	51
21	Dynamics and Clinical Evolution of Bacterial Gut Microflora in Extremely Premature Patients. Journal of Pediatrics, 2011, 158, 390-396.	0.9	188
22	Comparative diffusion assay to assess efficacy of topical antimicrobial agents against Pseudomonas aeruginosa in burns care. Annals of Clinical Microbiology and Antimicrobials, 2011, 10, 27.	1.7	8
23	Multilocus Sequence-Based Analysis Delineates a Clonal Population of Agrobacterium (Rhizobium) radiobacter (Agrobacterium tumefaciens) of Human Origin. Journal of Bacteriology, 2011, 193, 2608-2618.	1.0	29
24	Intragenomic and intraspecific heterogeneity in rrs may surpass interspecific variability in a natural population of Veillonella. Microbiology (United Kingdom), 2010, 156, 2080-2091.	0.7	33
25	Multilocus sequence typing supports the hypothesis that Ochrobactrum anthropi displays a human-associated subpopulation. BMC Microbiology, 2009, 9, 267.	1.3	48
26	Optimized PCR-Temporal Temperature Gel Electrophoresis compared to cultivation to assess diversity of gut microbiota in neonates. Journal of Microbiological Methods, 2009, 79, 156-165.	0.7	33
27	Aquaporin molecular characterization in the sea-bass (Dicentrarchus labrax): The effect of salinity on AQP1 and AQP3 expression. Comparative Biochemistry and Physiology Part A, Molecular & Camp; Integrative Physiology, 2007, 148, 430-444.	0.8	90
28	Digestive tract ontogeny of Dicentrarchus labrax: Implication in osmoregulation. Development Growth and Differentiation, 2006, 48, 139-151.	0.6	48
29	Differential freshwater adaptation in juvenile sea-bass Dicentrarchus labrax: involvement of gills and urinary system. Journal of Experimental Biology, 2005, 208, 3859-3871.	0.8	80
30	Ontogeny of osmoregulatory structures and functions in the green crab Carcinus maenas (Crustacea, Decapoda). Journal of Experimental Biology, 2004, 207, 325-336.	0.8	83
31	Putative involvement of crustacean hyperglycemic hormone isoforms in the neuroendocrine mediation of osmoregulation in the crayfishAstacus leptodactylus. Journal of Experimental Biology, 2003, 206, 979-988.	0.8	105