## **Bertrand Aigle**

List of Publications by Year in descending order

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REDTRAND AICLE

#	Article	IF	CITATIONS
1	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. Nature Biotechnology, 2016, 34, 828-837.	9.4	2,802
2	Minimum Information about a Biosynthetic Gene cluster. Nature Chemical Biology, 2015, 11, 625-631.	3.9	715
3	Towards the sustainable discovery and development of new antibiotics. Nature Reviews Chemistry, 2021, 5, 726-749.	13.8	439
4	Identification of a bioactive 51-membered macrolide complex by activation of a silent polyketide synthase in <i>Streptomyces ambofaciens</i> . Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 6258-6263.	3.3	275
5	Evolution of the Terminal Regions of the Streptomyces Linear Chromosome. Molecular Biology and Evolution, 2006, 23, 2361-2369.	3.5	96
6	Characterization and Manipulation of the Pathway-Specific Late Regulator AlpW Reveals <i>Streptomyces ambofaciens</i> as a New Producer of Kinamycins. Journal of Bacteriology, 2011, 193, 1142-1153.	1.0	96
7	Genome mining of <i>Streptomyces ambofaciens</i> . Journal of Industrial Microbiology and Biotechnology, 2014, 41, 251-263.	1.4	85
8	Functional Angucycline-Like Antibiotic Gene Cluster in the Terminal Inverted Repeats of the Streptomyces ambofaciens Linear Chromosome. Antimicrobial Agents and Chemotherapy, 2004, 48, 575-588.	1.4	65
9	Pseudomonas fluorescens Pirates both Ferrioxamine and Ferricoelichelin Siderophores from Streptomyces ambofaciens. Applied and Environmental Microbiology, 2015, 81, 3132-3141.	1.4	62
10	Comparative Genomics among Closely Related Streptomyces Strains Revealed Specialized Metabolite Biosynthetic Gene Cluster Diversity. Antibiotics, 2018, 7, 86.	1.5	53
11	Involvement of AlpV, a New Member of the Streptomyces Antibiotic Regulatory Protein Family, in Regulation of the Duplicated Type II Polyketide Synthase alp Gene Cluster in Streptomyces ambofaciens. Journal of Bacteriology, 2005, 187, 2491-2500.	1.0	40
12	Kinamycin biosynthesis employs a conserved pair of oxidases for B-ring contraction. Chemical Communications, 2015, 51, 8845-8848.	2.2	39
13	Role of secondary metabolites in the interaction between <i>Pseudomonas fluorescens</i> and soil microorganisms under iron-limited conditions. FEMS Microbiology Ecology, 2016, 92, fiw107.	1.3	39
14	Regulation of the Synthesis of the Angucyclinone Antibiotic Alpomycin in <i>Streptomyces ambofaciens</i> by the Autoregulator Receptor AlpZ and Its Specific Ligand. Journal of Bacteriology, 2008, 190, 3293-3305.	1.0	38
15	Waking up Streptomyces Secondary Metabolism by Constitutive Expression of Activators or Genetic Disruption of Repressors. Methods in Enzymology, 2012, 517, 343-366.	0.4	33
16	Identification of Alp1U and Lom6 as epoxy hydrolases and implications for kinamycin and lomaiviticin biosynthesis. Nature Communications, 2015, 6, 7674.	5.8	33
17	Intraspecific Variability of the Terminal Inverted Repeats of the Linear Chromosome of Streptomyces ambofaciens. Journal of Bacteriology, 2006, 188, 6599-6610.	1.0	32
18	A Single Sfp-Type Phosphopantetheinyl Transferase Plays a Major Role in the Biosynthesis of PKS and NRPS Derived Metabolites in Streptomyces ambofaciens ATCC23877. PLoS ONE, 2014, 9, e87607.	1.1	32

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19	An Unprecedented 1,2‣hift in the Biosynthesis of the 3â€Aminosalicylate Moiety of Antimycins. ChemBioChem, 2012, 13, 769-773.	1.3	31
20	End-to-end fusion of linear deleted chromosomes initiates a cycle of genome instability in Streptomyces ambofaciens. Molecular Microbiology, 2003, 50, 411-425.	1.2	30
21	Dynamics of the compartmentalized Streptomyces chromosome during metabolic differentiation. Nature Communications, 2021, 12, 5221.	5.8	30
22	Volatile Lactones from Streptomycetes Arise via the Antimycin Biosynthetic Pathway. ChemBioChem, 2012, 13, 1635-1644.	1.3	29
23	Complete genome sequence of Streptomyces ambofaciens ATCC 23877, the spiramycin producer. Journal of Biotechnology, 2015, 214, 117-118.	1.9	29
24	Cytochrome P450-mediated hydroxylation is required for polyketide macrolactonization in stambomycin biosynthesis. Journal of Antibiotics, 2014, 67, 71-76.	1.0	22
25	N-acylation of L-amino acids in aqueous media: Evaluation of the catalytic performances of Streptomyces ambofaciens aminoacylases. Enzyme and Microbial Technology, 2020, 137, 109536.	1.6	22
26	Differential and Cross-Transcriptional Control of Duplicated Genes Encoding Alternative Sigma Factors in Streptomyces ambofaciens. Journal of Bacteriology, 2004, 186, 5355-5365.	1.0	13
27	An aminoacylase activity from <i>Streptomyces ambofaciens</i> catalyzes the acylation of lysine on αâ€position and peptides on Nâ€terminal position. Engineering in Life Sciences, 2018, 18, 589-599.	2.0	12
28	Characterization of two Streptomyces ambofaciens recA mutants: identification of the RecA protein by immunoblotting. FEMS Microbiology Letters, 2006, 149, 181-187.	0.7	10
29	Gluconic acid-producing Pseudomonas sp. prevent γ-actinorhodin biosynthesis by Streptomyces coelicolor A3(2). Archives of Microbiology, 2014, 196, 619-627.	1.0	10
30	Diversity and antimicrobial activities of Streptomyces isolates from Fetzara Lake, north eastern Algeria. Annales De Biologie Clinique, 2018, 76, 81-95.	0.2	9
31	Engineering the stambomycin modular polyketide synthase yields 37-membered mini-stambomycins. Nature Communications, 2022, 13, 515.	5.8	8
32	Isolation and characterization of a mutator strain of <i>Streptomyces ambofaciens</i> ATCC23877 exhibiting an increased level of genetic instability. Canadian Journal of Microbiology, 1996, 42, 562-570.	0.8	5
33	Inhibitions Dominate but Stimulations and Growth Rescues Are Not Rare Among Bacterial Isolates from Grains of Forest Soil. Microbial Ecology, 2020, 80, 872-884.	1.4	2
34	Molecular Dynamics to Elucidate the DNA-Binding Activity of AlpZ, a Member of the Gamma-Butyrolactone Receptor Family in Streptomyces ambofaciens. Frontiers in Microbiology, 2020, 11, 1255.	1.5	2
35	Draft Whole-Genome Shotgun Sequence of Streptomyces sp. Strain ETH9427. Microbiology Resource Announcements, 2018, 7, .	0.3	1