## Sylvie Lautru

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6193034/publications.pdf

Version: 2024-02-01

24 2,372 16 25
papers citations h-index g-index

27 27 27 3275
all docs docs citations times ranked citing authors

#	Article	lF	CITATIONS
1	Minimum Information about a Biosynthetic Gene cluster. Nature Chemical Biology, 2015, 11, 625-631.	8.0	715
2	Discovery of a new peptide natural product by Streptomyces coelicolor genome mining. Nature Chemical Biology, 2005, 1, 265-269.	8.0	331
3	Cyclodipeptide synthases are a family of tRNA-dependent peptide bond–forming enzymes. Nature Chemical Biology, 2009, 5, 414-420.	8.0	215
4	The nonribosomal synthesis of diketopiperazines in tRNA-dependent cyclodipeptide synthase pathways. Natural Product Reports, 2012, 29, 961.	10.3	140
5	A new approach to bioconjugates for proteins and peptides ("pegylationâ€) utilising living radical polymerisation. Chemical Communications, 2004, , 2026-2027.	4.1	138
6	The Albonoursin Gene Cluster of S. noursei. Chemistry and Biology, 2002, 9, 1355-1364.	6.0	133
7	Multiple biosynthetic and uptake systems mediate siderophore-dependent iron acquisition in Streptomyces coelicolor A3(2) and Streptomyces ambofaciens ATCC 23877. Microbiology (United) Tj ETQq1 1 (	<b>).78.4</b> 314	rgBīī2¦Overlo <mark>c</mark> i
8	Substrate recognition by nonribosomal peptide synthetase multi-enzymes. Microbiology (United) Tj ETQq0 0 0 r	gBT_lOver	ock 10 Tf 50 4
9	MbtH-like protein-mediated cross-talk between non-ribosomal peptide antibiotic and siderophore biosynthetic pathways in Streptomyces coelicolor M145. Microbiology (United Kingdom), 2007, 153, 1405-1412.	1.8	93
10	Genome mining of <i>Streptomyces ambofaciens</i> Journal of Industrial Microbiology and Biotechnology, 2014, 41, 251-263.	3.0	85
11	An Iterative Nonribosomal Peptide Synthetase Assembles the Pyrrole-Amide Antibiotic Congocidine in Streptomyces ambofaciens. Chemistry and Biology, 2009, 16, 421-431.	6.0	54
12	Glycosylation Steps during Spiramycin Biosynthesis in <i>Streptomyces ambofaciens</i> : Involvement of Three Glycosyltransferases and Their Interplay with Two Auxiliary Proteins. Antimicrobial Agents and Chemotherapy, 2010, 54, 2830-2839.	3.2	36
13	Cyclic dipeptide oxidase from Streptomyces noursei. FEBS Journal, 2001, 268, 1712-1721.	0.2	34
14	Natural Combinatorial Biosynthesis Involving Two Clusters for the Synthesis of Three Pyrrolamides in <i>Streptomyces netropsis</i> . ACS Chemical Biology, 2015, 10, 601-610.	3.4	30
15	Dynamics of the compartmentalized Streptomyces chromosome during metabolic differentiation. Nature Communications, 2021, 12, 5221.	12.8	30
16	Complete genome sequence of Streptomyces ambofaciens ATCC 23877, the spiramycin producer. Journal of Biotechnology, 2015, 214, 117-118.	3.8	29
17	A Sweet Origin for the Key Congocidine Precursor 4â€Acetamidopyrroleâ€2â€carboxylate. Angewandte Chemie - International Edition, 2012, 51, 7454-7458.	13.8	17
18	Cyclic dipeptide oxidase from Streptomyces nourseilsolation, purification and partial characterization of a novel, amino acyl alpha, beta-dehydrogenase. FEBS Journal, 2001, 268, 1712-1721.	0.2	17

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#	Article	IF	CITATION
19	Modular and Integrative Vectors for Synthetic Biology Applications in <i>Streptomyces</i> spp. Applied and Environmental Microbiology, 2019, 85, .	3.1	14
20	Post-PKS Tailoring Steps of the Spiramycin Macrolactone Ring in Streptomyces ambofaciens. Antimicrobial Agents and Chemotherapy, 2013, 57, 3836-3842.	3.2	13
21	Study of bicyclomycin biosynthesis in Streptomyces cinnamoneus by genetic and biochemical approaches. Scientific Reports, 2019, 9, 20226.	3.3	12
22	Revised Structure of Anthelvencin A and Characterization of the Anthelvencin Biosynthetic Gene Cluster. ACS Chemical Biology, 2020, 15, 945-951.	3.4	9
23	Draft Genome Sequence of <i>Streptomyces</i> sp. M1013, a Close Relative of Streptomyces ambofaciens and Streptomyces coelicolor. Genome Announcements, 2017, 5, .	0.8	3
24	Transcriptional regulation of congocidine (netropsin) biosynthesis and resistance Applied and Environmental Microbiology, 2021, 87, e0138021.	3.1	0