

Jean-Luc Pernodet

List of Publications by Year in descending order

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69
papers

4,800
citations

126858

33
h-index

98753

67
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73
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73
docs citations

73
times ranked

4542
citing authors

#	ARTICLE	IF	CITATIONS
1	Marker-Free Genome Engineering in <i>Amycolatopsis</i> Using the pSAM2 Site-Specific Recombination System. <i>Microorganisms</i> , 2022, 10, 828.	1.6	3
2	Towards the sustainable discovery and development of new antibiotics. <i>Nature Reviews Chemistry</i> , 2021, 5, 726-749.	13.8	439
3	Transcriptional regulation of congocidine (netropsin) biosynthesis and resistance.. <i>Applied and Environmental Microbiology</i> , 2021, 87, e0138021.	1.4	0
4	Dynamics of the compartmentalized <i>Streptomyces</i> chromosome during metabolic differentiation. <i>Nature Communications</i> , 2021, 12, 5221.	5.8	30
5	Revised Structure of Anthelvencin A and Characterization of the Anthelvencin Biosynthetic Gene Cluster. <i>ACS Chemical Biology</i> , 2020, 15, 945-951.	1.6	9
6	Modular and Integrative Vectors for Synthetic Biology Applications in <i>Streptomyces</i> spp. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	14
7	Design of a generic CRISPR-Cas9 approach using the same sgRNA to perform gene editing at distinct loci. <i>BMC Biotechnology</i> , 2019, 19, 18.	1.7	11
8	Study of bicyclomycin biosynthesis in <i>Streptomyces cinnamoneus</i> by genetic and biochemical approaches. <i>Scientific Reports</i> , 2019, 9, 20226.	1.6	12
9	A Comprehensive Overview of the Cyclodipeptide Synthase Family Enriched with the Characterization of 32 New Enzymes. <i>Frontiers in Microbiology</i> , 2018, 9, 46.	1.5	52
10	Complete Genome Sequence of <i>Streptomyces</i> sp. TN58, a Producer of Acyl Alpha-1-Rhamnopyranosides. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
11	Draft Genome Sequence of <i>Streptomyces</i> sp. M1013, a Close Relative of <i>Streptomyces ambofaciens</i> and <i>Streptomyces coelicolor</i> . <i>Genome Announcements</i> , 2017, 5, .	0.8	3
12	Analysis of 51 cyclodipeptide synthases reveals the basis for substrate specificity. <i>Nature Chemical Biology</i> , 2015, 11, 721-727.	3.9	70
13	Complete genome sequence of <i>Streptomyces ambofaciens</i> ATCC 23877, the spiramycin producer. <i>Journal of Biotechnology</i> , 2015, 214, 117-118.	1.9	29
14	The Absence of Pupylation (Prokaryotic Ubiquitin-Like Protein Modification) Affects Morphological and Physiological Differentiation in <i>Streptomyces coelicolor</i> . <i>Journal of Bacteriology</i> , 2015, 197, 3388-3399.	1.0	35
15	Minimum Information about a Biosynthetic Gene cluster. <i>Nature Chemical Biology</i> , 2015, 11, 625-631.	3.9	715
16	Characterization of Sviceucin from <i>Streptomyces</i> Provides Insight into Enzyme Exchangeability and Disulfide Bond Formation in Lasso Peptides. <i>ACS Chemical Biology</i> , 2015, 10, 2641-2649.	1.6	73
17	Natural Combinatorial Biosynthesis Involving Two Clusters for the Synthesis of Three Pyrrolamides in <i>Streptomyces netropsis</i> . <i>ACS Chemical Biology</i> , 2015, 10, 601-610.	1.6	30
18	Genome mining of <i>Streptomyces ambofaciens</i> . <i>Journal of Industrial Microbiology and Biotechnology</i> , 2014, 41, 251-263.	1.4	85

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19	The Genome Sequence of <i>Streptomyces lividans</i> 66 Reveals a Novel tRNA-Dependent Peptide Biosynthetic System within a Metal-Related Genomic Island. <i>Genome Biology and Evolution</i> , 2013, 5, 1165-1175.	1.1	99
20	Post-PKS Tailoring Steps of the Spiramycin Macrolactone Ring in <i>Streptomyces ambofaciens</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 3836-3842.	1.4	13
21	The nonribosomal synthesis of diketopiperazines in tRNA-dependent cyclodipeptide synthase pathways. <i>Natural Product Reports</i> , 2012, 29, 961.	5.2	140
22	A Sweet Origin for the Key Congocidine Precursor 4- <i>N</i> -Acetamidopyrrole-2-carboxylate. <i>Angewandte Chemie - International Edition</i> , 2012, 51, 7454-7458.	7.2	17
23	Nonribosomal Peptide Synthesis in Animals: The Cyclodipeptide Synthase of <i>Nematostella</i> . <i>Chemistry and Biology</i> , 2011, 18, 1362-1368.	6.2	50
24	Cyclodipeptide synthases, a family of class-I aminoacyl-tRNA synthetase-like enzymes involved in non-ribosomal peptide synthesis. <i>Nucleic Acids Research</i> , 2011, 39, 4475-4489.	6.5	83
25	Regulation of the Biosynthesis of the Macrolide Antibiotic Spiramycin in <i>Streptomyces ambofaciens</i> . <i>Journal of Bacteriology</i> , 2010, 192, 5813-5821.	1.0	31
26	Glycosylation Steps during Spiramycin Biosynthesis in <i>Streptomyces ambofaciens</i> : Involvement of Three Glycosyltransferases and Their Interplay with Two Auxiliary Proteins. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 2830-2839.	1.4	36
27	Transcriptional regulation of the novobiocin biosynthetic gene cluster. <i>Microbiology (United Kingdom)</i> Tj ETQq1 1 0.784314 rgBT / Overlock 10	0.7	15
28	Cyclodipeptide synthases are a family of tRNA-dependent peptide bond-forming enzymes. <i>Nature Chemical Biology</i> , 2009, 5, 414-420.	3.9	215
29	An Iterative Nonribosomal Peptide Synthetase Assembles the Pyrrole-Amide Antibiotic Congocidine in <i>Streptomyces ambofaciens</i> . <i>Chemistry and Biology</i> , 2009, 16, 421-431.	6.2	54
30	Identification and structural basis of the reaction catalyzed by CYP121, an essential cytochrome P450 in <i>Mycobacterium tuberculosis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 7426-7431.	3.3	192
31	The integrative element pSAM2 from <i>Streptomyces</i> : kinetics and mode of conjugal transfer. <i>Molecular Microbiology</i> , 2008, 42, 159-166.	1.2	86
32	MbtH-like protein-mediated cross-talk between non-ribosomal peptide antibiotic and siderophore biosynthetic pathways in <i>Streptomyces coelicolor</i> M145. <i>Microbiology (United Kingdom)</i> , 2007, 153, 1405-1412.	0.7	93
33	Organization of the biosynthetic gene cluster for the macrolide antibiotic spiramycin in <i>Streptomyces ambofaciens</i> . <i>Microbiology (United Kingdom)</i> , 2007, 153, 4111-4122.	0.7	54
34	Intraspecific Variability of the Terminal Inverted Repeats of the Linear Chromosome of <i>Streptomyces ambofaciens</i> . <i>Journal of Bacteriology</i> , 2006, 188, 6599-6610.	1.0	32
35	Evolution of the Terminal Regions of the <i>Streptomyces</i> Linear Chromosome. <i>Molecular Biology and Evolution</i> , 2006, 23, 2361-2369.	3.5	96
36	Construction and testing of a bacterial luciferase reporter gene system for in vivo measurement of nonsense suppression in <i>Streptomyces</i> . <i>Folia Microbiologica</i> , 2006, 51, 62-4.	1.1	2

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37	Excisable Cassettes: New Tools for Functional Analysis of Streptomyces Genomes. Applied and Environmental Microbiology, 2006, 72, 4839-4844.	1.4	38
38	Multiple biosynthetic and uptake systems mediate siderophore-dependent iron acquisition in Streptomyces coelicolor A3(2) and Streptomyces ambofaciens ATCC 23877. Microbiology (United Kingdom), 2007, 147, 1010-1015.	1.7	10
39	Methyltransferase Erm(37) Slips on rRNA to Confer Atypical Resistance in Mycobacterium tuberculosis. Journal of Biological Chemistry, 2005, 280, 38942-38947.	1.6	65
40	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
41	Molecular Basis of Intrinsic Macrolide Resistance in the Mycobacterium tuberculosis Complex. Antimicrobial Agents and Chemotherapy, 2004, 48, 143-150.	1.4	135
42	Conjugal immunity of Streptomyces strains carrying the integrative element pSAM2 is due to the pif gene (pSAM2 immunity factor). Molecular Microbiology, 2003, 47, 1385-1393.	1.2	17
43	Recombinant Environmental Libraries Provide Access to Microbial Diversity for Drug Discovery from Natural Products. Applied and Environmental Microbiology, 2003, 69, 49-55.	1.4	305
44	Natural and Acquired Macrolide Resistance in Mycobacteria. Current Drug Targets Infectious Disorders, 2002, 2, 355-370.	2.1	38
45	The Albonoursin Gene Cluster of S. noursei. Chemistry and Biology, 2002, 9, 1355-1364.	6.2	133
46	Erratum to "Development of a conditional lethal system for a Streptomyces lividans strain and its use to investigate conjugative transfer in soil" [FEMS Microbiology Ecology 38 (2001) 115-121]. FEMS Microbiology Ecology, 2002, 40, 83-84.	1.3	0
47	Erratum to "Development of a conditional lethal system for a Streptomyces lividans strain and its use to investigate conjugative transfer in soil" [FEMS Microbiology Ecology, 2002, 40, 83-84].	1.3	0
48	Characterization of the attP site of the integrative element pSAM2 from Streptomyces ambofaciens. Microbiology (United Kingdom), 2002, 148, 61-67.	0.7	13
49	Development of a conditional lethal system for a Streptomyces lividans strain and its use to investigate conjugative transfer in soil. FEMS Microbiology Ecology, 2001, 38, 115-121.	1.3	1
50	KorSA from the Streptomyces Integrative Element pSAM2 Is a Central Transcriptional Repressor: Target Genes and Binding Sites. Journal of Bacteriology, 2000, 182, 1243-1250.	1.0	33
51	Dispensable ribosomal resistance to spiramycin conferred by srmA in the spiramycin producer Streptomyces ambofaciens The EMBL/GenBank accession number for the nucleotide sequence described in this paper is AJ223970.. Microbiology (United Kingdom), 1999, 145, 2355-2364.	0.7	10
52	Structure of the chromosomal insertion site for pSAM2: functional analysis in Escherichia coli. Molecular Microbiology, 1998, 28, 333-342.	1.2	37
53	Characterization of a Glycosyl Transferase Inactivating Macrolides, Encoded by <i>gimA</i> from <i>Streptomyces ambofaciens</i> . Antimicrobial Agents and Chemotherapy, 1998, 42, 2612-2619.	1.4	29
54	Replicase, Excisionase, and Integrase Genes of the <i>Streptomyces</i> Element pSAM2 Constitute an Operon Positively Regulated by the <i>pra</i> Gene. Journal of Bacteriology, 1998, 180, 3056-3061.	1.0	25

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55	Antibiotic resistance gene cassettes derived from the \int interposon for use in <i>E. coli</i> and <i>Streptomyces</i> . <i>Gene</i> , 1997, 190, 315-317.	1.0	193
56	Complete conversion of antibiotic precursor to pristinamycin IIA by overexpression of <i>Streptomyces pristinaespiralis</i> biosynthetic genes. <i>Nature Biotechnology</i> , 1997, 15, 349-353.	9.4	43
57	The macrolide-lincosamide-streptogramin B resistance phenotypes characterized by using a specifically deleted, antibiotic-sensitive strain of <i>Streptomyces lividans</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 1996, 40, 581-585.	1.4	32
58	Characterization of <i>pra</i> , a gene for replication control in pSAM2, the integrating element of <i>Streptomyces ambofaciens</i> . <i>Molecular Microbiology</i> , 1995, 17, 533-544.	1.2	20
59	Identification of a Gene Encoding the Replication Initiator Protein of the <i>Streptomyces</i> Integrating Element, pSAM2. <i>Plasmid</i> , 1994, 31, 166-183.	0.4	23
60	Cloning of <i>Frankia</i> species putative tRNA(Pro) genes and their efficacy for pSAM2 site-specific integration in <i>Streptomyces lividans</i> . <i>Applied and Environmental Microbiology</i> , 1994, 60, 4279-4283.	1.4	11
61	Mode and origin of replication of pSAM2, a conjugative integrating element of <i>Streptomyces ambofaciens</i> . <i>Molecular Microbiology</i> , 1993, 10, 799-812.	1.2	33
62	Transfer functions of the conjugative integrating element pSAM2 from <i>Streptomyces ambofaciens</i> : characterization of a <i>kil-kor</i> system associated with transfer. <i>Journal of Bacteriology</i> , 1993, 175, 5529-5538.	1.0	52
63	Functional analysis of the <i>Streptomyces ambofaciens</i> element pSAM2. <i>Plasmid</i> , 1991, 25, 40-52.	0.4	36
64	Organization and nucleotide sequence analysis of a ribosomal RNA gene cluster from <i>Streptomyces ambofaciens</i> . <i>Gene</i> , 1989, 79, 33-46.	1.0	137
65	Structural analysis of loci involved in pSAM2 site-specific integration in <i>Streptomyces</i> . <i>Plasmid</i> , 1989, 21, 59-70.	0.4	52
66	Site-specific integration of plasmid pSAM2 in <i>Streptomyces lividans</i> and <i>S. ambofaciens</i> . <i>Molecular Genetics and Genomics</i> , 1988, 212, 432-439.	2.4	26
67	Excision and integration of a self-transmissible replicon of <i>Streptomyces ambofaciens</i> . <i>Gene</i> , 1987, 59, 137-144.	1.0	27
68	Plasmids in different strains of <i>Streptomyces ambofaciens</i> : free and integrated form of plasmid pSAM2. <i>Molecular Genetics and Genomics</i> , 1984, 198, 35-41.	2.4	108
69	Isolation and physical characterization of streptomycete plasmids. <i>Molecular Genetics and Genomics</i> , 1981, 182, 53-59.	2.4	18