

Andriy Luzhetskyy

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/716247/publications.pdf>

Version: 2024-02-01

134
papers

5,976
citations

81743

39
h-index

88477

70
g-index

138
all docs

138
docs citations

138
times ranked

5027
citing authors

#	ARTICLE	IF	CITATIONS
1	Total In Vitro Biosynthesis of the Thioamide Thioholgamide and Investigation of the Pathway. <i>Journal of the American Chemical Society</i> , 2022, 144, 5136-5144.	6.6	19
2	Novel Biosynthetic Route to the Isoquinoline Scaffold. <i>ACS Chemical Biology</i> , 2022, 17, 598-608.	1.6	3
3	Scandium- ⁴⁵ microorganism interactions in new biotechnologies. <i>Trends in Biotechnology</i> , 2022, 40, 1088-1101.	4.9	7
4	Multiple copies of the oxytetracycline gene cluster in selected <i>Streptomyces rimosus</i> strains can provide significantly increased titers. <i>Microbial Cell Factories</i> , 2021, 20, 47.	1.9	5
5	Genetically engineered <i>rpsL</i> merodiploidy impacts secondary metabolism and antibiotic resistance in <i>Streptomyces</i> . <i>World Journal of Microbiology and Biotechnology</i> , 2021, 37, 62.	1.7	5
6	Microparticles enhance the formation of seven major classes of natural products in native and metabolically engineered actinobacteria through accelerated morphological development. <i>Biotechnology and Bioengineering</i> , 2021, 118, 3076-3093.	1.7	15
7	Discovery and Heterologous Production of New Cyclic Depsibosamycins. <i>Microorganisms</i> , 2021, 9, 1396.	1.6	2
8	Superior production of heavy pamamycin derivatives using a <i>bkdR</i> deletion mutant of <i>Streptomyces albus</i> J1074/R2. <i>Microbial Cell Factories</i> , 2021, 20, 111.	1.9	11
9	Cyclofaulknamycin with the Rare Amino Acid D-capreomycinidine Isolated from a Well-Characterized <i>Streptomyces albus</i> Strain. <i>Microorganisms</i> , 2021, 9, 1609.	1.6	3
10	Bonsecamin: A New Cyclic Pentapeptide Discovered through Heterologous Expression of a Cryptic Gene Cluster. <i>Microorganisms</i> , 2021, 9, 1640.	1.6	3
11	Towards the sustainable discovery and development of new antibiotics. <i>Nature Reviews Chemistry</i> , 2021, 5, 726-749.	13.8	439
12	The diversity and antibacterial activity of culturable actinobacteria isolated from the rhizosphere soil of <i>Deschampsia antarctica</i> (Galindez Island, Maritime Antarctic). <i>Polar Biology</i> , 2021, 44, 1859-1868.	0.5	5
13	Engineering the precursor pool to modulate the production of pamamycins in the heterologous host <i>S. albus</i> J1074. <i>Metabolic Engineering</i> , 2021, 67, 11-18.	3.6	7
14	New Scabimycins A-C Isolated from <i>Streptomyces acidiscabies</i> (Lu19992). <i>Molecules</i> , 2021, 26, 5922.	1.7	1
15	A Promiscuous Halogenase for the Derivatization of Flavonoids. <i>Molecules</i> , 2021, 26, 6220.	1.7	4
16	New Kendomycin Derivative Isolated from <i>Streptomyces</i> sp. Cl 58-27. <i>Molecules</i> , 2021, 26, 6834.	1.7	3
17	Engineering <i>Corynebacterium glutamicum</i> with a comprehensive genomic library and phage-based vectors. <i>Metabolic Engineering</i> , 2020, 62, 221-234.	3.6	4
18	Non-Heme Monooxygenase ThoJ Catalyzes Thioholgamide $\hat{1}^2$ -Hydroxylation. <i>ACS Chemical Biology</i> , 2020, 15, 2815-2819.	1.6	9

#	ARTICLE	IF	CITATIONS
19	Dudomycins: New Secondary Metabolites Produced after Heterologous Expression of an Nrps Cluster from <i>Streptomyces albus</i> ssp. <i>Chlorinus</i> Nr1 B-24108. <i>Microorganisms</i> , 2020, 8, 1800.	1.6	4
20	Identification of a Biosynthetic Gene Cluster Responsible for the Production of a New Pyrrolopyrimidine Natural Product—Huimycin. <i>Biomolecules</i> , 2020, 10, 1074.	1.8	11
21	Loseolamycins: A Group of New Bioactive Alkylresorcinols Produced after Heterologous Expression of a Type III PKS from <i>Micromonospora endolithica</i> . <i>Molecules</i> , 2020, 25, 4594.	1.7	7
22	Microparticles globally reprogram <i>Streptomyces albus</i> toward accelerated morphogenesis, streamlined carbon core metabolism, and enhanced production of the antituberculosis polyketide pamamycin. <i>Biotechnology and Bioengineering</i> , 2020, 117, 3858-3875.	1.7	22
23	Targeted Genome Mining—From Compound Discovery to Biosynthetic Pathway Elucidation. <i>Microorganisms</i> , 2020, 8, 2034.	1.6	3
24	Baikalomycins A-C, New Aquayamycin-Type Angucyclines Isolated from Lake Baikal Derived <i>Streptomyces</i> sp. IB201691-2A. <i>Microorganisms</i> , 2020, 8, 680.	1.6	19
25	Novel Fredericamycin Variant Overproduced by a Streptomycin-Resistant <i>Streptomyces albus</i> subsp. <i>chlorinus</i> Strain. <i>Marine Drugs</i> , 2020, 18, 284.	2.2	10
26	The bottromycin epimerase BotH defines a group of atypical β -hydrolase-fold enzymes. <i>Nature Chemical Biology</i> , 2020, 16, 1013-1018.	3.9	18
27	Identification and Heterologous Expression of the Albucidin Gene Cluster from the Marine Strain <i>Streptomyces Albus</i> Subsp. <i>Chlorinus</i> NRRL B-24108. <i>Microorganisms</i> , 2020, 8, 237.	1.6	18
28	Engineering of <i>Streptomyces lividans</i> for heterologous expression of secondary metabolite gene clusters. <i>Microbial Cell Factories</i> , 2020, 19, 5.	1.9	68
29	Thioholgamide A, a New Anti-Proliferative Anti-Tumor Agent, Modulates Macrophage Polarization and Metabolism. <i>Cancers</i> , 2020, 12, 1288.	1.7	22
30	Back Cover Image, Volume 117, Number 12, December 2020. <i>Biotechnology and Bioengineering</i> , 2020, 117, .	1.7	0
31	Perquinoline...: neuartige bakterielle Tetrahydroisochinoline mit einer bemerkenswerten Biosynthese. <i>Angewandte Chemie</i> , 2019, 131, 13063-13068.	1.6	0
32	Effect of ϵ -ribosome engineering on the transcription level and production of <i>S. albus</i> indigenous secondary metabolites. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 7097-7110.	1.7	23
33	Perquinolines: Unprecedented Bacterial Tetrahydroisoquinolines Involving an Intriguing Biosynthesis. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 12930-12934.	7.2	10
34	Heterologous production of small molecules in the optimized <i>Streptomyces</i> hosts. <i>Natural Product Reports</i> , 2019, 36, 1281-1294.	5.2	65
35	Gene <i>miaA</i> for post-transcriptional modification of tRNA ^{XXA} is important for morphological and metabolic differentiation in <i>Streptomyces</i> . <i>Molecular Microbiology</i> , 2019, 112, 249-265.	1.2	26
36	Genome Engineering Approaches to Improve Nosokomycin A Production by <i>Streptomyces ghanaensis</i> B38.3. <i>Indian Journal of Microbiology</i> , 2019, 59, 109-111.	1.5	3

#	ARTICLE	IF	CITATIONS
37	Benzantronic Acid, a Novel Metabolite From <i>Streptomyces albus</i> Del14 Expressing the Nybomycin Gene Cluster. <i>Frontiers in Chemistry</i> , 2019, 7, 896.	1.8	15
38	Gene <i>ssfg_01967</i> (<i>miaB</i>) for tRNA modification influences morphogenesis and moenomycin biosynthesis in <i>Streptomyces ghanaensis</i> ATCC14672. <i>Microbiology (United Kingdom)</i> , 2019, 165, 233-245.	0.7	11
39	A set of synthetic versatile genetic control elements for the efficient expression of genes in Actinobacteria. <i>Scientific Reports</i> , 2018, 8, 491.	1.6	50
40	Design, development and application of whole-cell based antibiotic-specific biosensor. <i>Metabolic Engineering</i> , 2018, 47, 263-270.	3.6	33
41	Enabling the valorization of guaiacol-based lignin: Integrated chemical and biochemical production of <i>cis,cis</i> -muconic acid using metabolically engineered <i>Amycolatopsis</i> sp ATCC 39116. <i>Metabolic Engineering</i> , 2018, 45, 200-210.	3.6	125
42	Heterologous Expression of the Nybomycin Gene Cluster from the Marine Strain <i>Streptomyces albus</i> subsp. <i>chlorinus</i> NRRL B-24108. <i>Marine Drugs</i> , 2018, 16, 435.	2.2	22
43	Monitoring Protein Secretion in <i>Streptomyces</i> Using Fluorescent Proteins. <i>Frontiers in Microbiology</i> , 2018, 9, 3019.	1.5	11
44	Characterization of Sigma Factor Genes in <i>Streptomyces lividans</i> TK24 Using a Genomic Library-Based Approach for Multiple Gene Deletions. <i>Frontiers in Microbiology</i> , 2018, 9, 3033.	1.5	23
45	Generation of a cluster-free <i>Streptomyces albus</i> chassis strains for improved heterologous expression of secondary metabolite clusters. <i>Metabolic Engineering</i> , 2018, 49, 316-324.	3.6	140
46	Secondary metabolites overproduction through transcriptional gene cluster refactoring. <i>Metabolic Engineering</i> , 2018, 49, 299-315.	3.6	63
47	Heterologous AdpA transcription factors enhance landomycin production in <i>Streptomyces cyanogenus</i> S136 under a broad range of growth conditions. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 8419-8428.	1.7	22
48	Multi-Omics and Targeted Approaches to Determine the Role of Cellular Proteases in <i>Streptomyces</i> Protein Secretion. <i>Frontiers in Microbiology</i> , 2018, 9, 1174.	1.5	29
49	New Alpiniamides From <i>Streptomyces</i> sp. IB2014/011-12 Assembled by an Unusual Hybrid Non-ribosomal Peptide Synthetase Trans-AT Polyketide Synthase Enzyme. <i>Frontiers in Microbiology</i> , 2018, 9, 1959.	1.5	19
50	Development of a Biosensor Concept to Detect the Production of Cluster-Specific Secondary Metabolites. <i>ACS Synthetic Biology</i> , 2017, 6, 1026-1033.	1.9	28
51	Chromosomal position effect influences the heterologous expression of genes and biosynthetic gene clusters in <i>Streptomyces albus</i> J1074. <i>Microbial Cell Factories</i> , 2017, 16, 5.	1.9	34
52	New natural products identified by combined genomics-metabolomics profiling of marine <i>Streptomyces</i> sp. MP131-18. <i>Scientific Reports</i> , 2017, 7, 42382.	1.6	86
53	Cloning and Expression of Metagenomic DNA in <i>Streptomyces lividans</i> and Subsequent Fermentation for Optimized Production. <i>Methods in Molecular Biology</i> , 2017, 1539, 99-144.	0.4	36
54	Properties of <i>Streptomyces albus</i> J1074 mutant deficient in tRNA ^{Leu} UAA gene <i>bldA</i> . <i>Archives of Microbiology</i> , 2017, 199, 1175-1183.	1.0	24

#	ARTICLE	IF	CITATIONS
55	Complete Draft Genome Sequence of the Actinobacterium <i>Nocardiosis sinuspersici</i> UTMC102 (DSM) Tj ETQq1 1 0,784314 1gBT /Over	0,8	1gBT /Over
56	Identification of butenolide regulatory system controlling secondary metabolism in <i>Streptomyces albus</i> J1074. <i>Scientific Reports</i> , 2017, 7, 9784.	1.6	34
57	<i>Streptomyces albus</i> : A New Cell Factory for Non-Canonical Amino Acids Incorporation into Ribosomally Synthesized Natural Products. <i>ACS Chemical Biology</i> , 2017, 12, 2362-2370.	1.6	38
58	Cloning and Heterologous Expression of the Grecoacycline Biosynthetic Gene Cluster. <i>PLoS ONE</i> , 2016, 11, e0158682.	1.1	61
59	A gene cluster for the biosynthesis of moenomycin family antibiotics in the genome of teicoplanin producer <i>Actinoplanes teichomyceticus</i> . <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 7629-7638.	1.7	12
60	Metabolic engineering of natural product biosynthesis in actinobacteria. <i>Current Opinion in Biotechnology</i> , 2016, 42, 98-107.	3.3	66
61	Dual control system – A novel scaffolding architecture of an inducible regulatory device for the precise regulation of gene expression. <i>Metabolic Engineering</i> , 2016, 37, 11-23.	3.6	31
62	Native and engineered promoters in natural product discovery. <i>Natural Product Reports</i> , 2016, 33, 1006-1019.	5.2	97
63	Generation of new compounds through unbalanced transcription of landomycin A cluster. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 9175-9186.	1.7	24
64	An influence of the copy number of biosynthetic gene clusters on the production level of antibiotics in a heterologous host. <i>Journal of Biotechnology</i> , 2016, 232, 110-117.	1.9	18
65	Characterization of the Post-Assembly Line Tailoring Processes in Teicoplanin Biosynthesis. <i>ACS Chemical Biology</i> , 2016, 11, 2254-2264.	1.6	18
66	New Simocyclinones: Surprising Evolutionary and Biosynthetic Insights. <i>ACS Chemical Biology</i> , 2016, 11, 241-250.	1.6	26
67	The isolation and characterization of actinobacteria from dominant benthic macroinvertebrates endemic to Lake Baikal. <i>Folia Microbiologica</i> , 2016, 61, 159-168.	1.1	15
68	The <i>adpA</i> -like regulatory gene from <i>Actinoplanes teichomyceticus</i> : in silico analysis and heterologous expression. <i>World Journal of Microbiology and Biotechnology</i> , 2015, 31, 1297-1301.	1.7	10
69	Insights into the Pamamycin Biosynthesis. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 2280-2284.	7.2	27
70	Endophytic <i>Streptomyces</i> in the traditional medicinal plant <i>Arnica montana</i> L.: secondary metabolites and biological activity. <i>Antonie Van Leeuwenhoek</i> , 2015, 108, 391-402.	0.7	13
71	Minimum Information about a Biosynthetic Gene cluster. <i>Nature Chemical Biology</i> , 2015, 11, 625-631.	3.9	715
72	Novel and tightly regulated resorcinol and cumate-inducible expression systems for <i>Streptomyces</i> and other actinobacteria. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 8641-8655.	1.7	61

#	ARTICLE	IF	CITATIONS
73	The pathway-specific regulatory genes, <i>tei15*</i> and <i>tei16*</i> , are the master switches of teicoplanin production in <i>Actinoplanes teichomyceticus</i> . <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 9295-9309.	1.7	36
74	Complete genome sequence of producer of the glycopeptide antibiotic Aculeximycin <i>Kutzneria albida</i> DSM 43870T, a representative of minor genus of Pseudonocardiaceae. <i>BMC Genomics</i> , 2014, 15, 885.	1.2	26
75	Actinomycetes biosynthetic potential: how to bridge in silico and in vivo?. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2014, 41, 387-402.	1.4	63
76	Unusual site-specific DNA integration into the highly active pseudo-attB of the <i>Streptomyces albus</i> J1074 genome. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 5095-5104.	1.7	42
77	Genome rearrangements of <i>Streptomyces albus</i> J1074 lead to the carotenoid gene cluster activation. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 795-806.	1.7	30
78	Iterative marker excision system. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 4557-4570.	1.7	49
79	Insights into naturally minimised <i>Streptomyces albus</i> J1074 genome. <i>BMC Genomics</i> , 2014, 15, 97.	1.2	137
80	Amycomycins C and D, new angucyclines from <i>Kitasatospora</i> sp.. <i>Tetrahedron Letters</i> , 2014, 55, 5771-5773.	0.7	4
81	Oleamycins A and B: new antibacterial cyclic hexadepsipeptides isolated from a terrestrial <i>Streptomyces</i> sp.. <i>Journal of Antibiotics</i> , 2014, 67, 339-343.	1.0	21
82	Regio- and Stereoselective Intermolecular Oxidative Phenol Coupling in <i>Streptomyces</i> . <i>Journal of the American Chemical Society</i> , 2014, 136, 6195-6198.	6.6	63
83	Lorneic acids C and D, new trialkyl-substituted aromatic acids isolated from a terrestrial <i>Streptomyces</i> sp.. <i>Journal of Antibiotics</i> , 2013, 66, 347-349.	1.0	10
84	Design, construction and characterisation of a synthetic promoter library for fine-tuned gene expression in actinomycetes. <i>Metabolic Engineering</i> , 2013, 19, 98-106.	3.6	172
85	In vivo random mutagenesis of streptomycetes using mariner-based transposon Himar1. <i>Applied Microbiology and Biotechnology</i> , 2013, 97, 351-359.	1.7	27
86	Tracking Down Biotransformation to the Genetic Level: Identification of a Highly Flexible Glycosyltransferase from <i>Saccharothrix espanaensis</i> . <i>Applied and Environmental Microbiology</i> , 2013, 79, 5224-5232.	1.4	13
87	Rubimycinone A, a new anthraquinone from a terrestrial <i>Streptomyces</i> sp.. <i>Tetrahedron Letters</i> , 2013, 54, 900-902.	0.7	8
88	Evaluation of heterologous promoters for genetic analysis of <i>Actinoplanes teichomyceticus</i> – Producer of teicoplanin, drug of last defense. <i>Journal of Biotechnology</i> , 2013, 168, 367-372.	1.9	26
89	A transposon-based strategy to identify the regulatory gene network responsible for landomycin E biosynthesis. <i>FEMS Microbiology Letters</i> , 2013, 342, 138-146.	0.7	16
90	Genome engineering in actinomycetes using site-specific recombinases. <i>Applied Microbiology and Biotechnology</i> , 2013, 97, 4701-4712.	1.7	16

#	ARTICLE	IF	CITATIONS
91	Oleaceran: A Novel Spiro[isobenzofuran-1,2- <i>napho</i> [1,8- <i>bc</i>]furan] Isolated from a Terrestrial <i>Streptomyces</i> sp.. <i>Organic Letters</i> , 2013, 15, 3487-3489.	2.4	23
92	<i>MoeH</i> : a natural glycorandomizer from the moenomycin biosynthetic pathway. <i>Molecular Microbiology</i> , 2013, 90, 1324-1338.	1.2	13
93	Pleiotropic regulatory genes <i>bldA</i> , <i>adpA</i> and <i>absB</i> are implicated in production of phosphoglycolipid antibiotic moenomycin. <i>Open Biology</i> , 2013, 3, 130121.	1.5	56
94	Site-Specific Recombination Strategies for Engineering Actinomycete Genomes. <i>Applied and Environmental Microbiology</i> , 2012, 78, 1804-1812.	1.4	88
95	Juniperolide A: A New Polyketide Isolated from a Terrestrial Actinomycete, <i>Streptomyces</i> sp.. <i>Organic Letters</i> , 2012, 14, 5860-5863.	2.4	15
96	Leopolic acid A, isolated from a terrestrial actinomycete, <i>Streptomyces</i> sp.. <i>Tetrahedron Letters</i> , 2012, 53, 6300-6301.	0.7	18
97	Actinomycetes genome engineering approaches. <i>Antonie Van Leeuwenhoek</i> , 2012, 102, 503-516.	0.7	26
98	SimReg1 is a master switch for biosynthesis and export of simocyclinone D8 and its precursors. <i>AMB Express</i> , 2012, 2, 1.	1.4	89
99	Pimprinols ¹⁴ C, from the terrestrial actinomycete, <i>Streptomyces</i> sp.. <i>Tetrahedron Letters</i> , 2012, 53, 3009-3011.	0.7	23
100	Rational Design of an Aryl-C-Glycoside Catalyst from a Natural Product O-Glycosyltransferase. <i>Chemistry and Biology</i> , 2011, 18, 520-530.	6.2	58
101	β -Glucuronidase as a Sensitive and Versatile Reporter in Actinomycetes. <i>Applied and Environmental Microbiology</i> , 2011, 77, 5370-5383.	1.4	182
102	Characterization and analysis of the regulatory network involved in control of lipomycin biosynthesis in <i>Streptomyces aureofaciens</i> TÅ¼117. <i>Applied Microbiology and Biotechnology</i> , 2010, 85, 1069-1079.	1.7	17
103	I-SceI endonuclease: a new tool for DNA repair studies and genetic manipulations in streptomycetes. <i>Applied Microbiology and Biotechnology</i> , 2010, 87, 1525-1532.	1.7	40
104	Cloning and Sequencing of the Biosynthetic Gene Cluster for Saquayamycin Z and Galtamycin B and the Elucidation of the Assembly of Their Saccharide Chains. <i>ChemBioChem</i> , 2009, 10, 1392-1401.	1.3	35
105	Differences in the substrate specificity of glycosyltransferases involved in landomycins A and E biosynthesis. <i>Applied Microbiology and Biotechnology</i> , 2009, 83, 1067-1076.	1.7	14
106	In vivo Tn5-based transposon mutagenesis of Streptomycetes. <i>Applied Microbiology and Biotechnology</i> , 2009, 83, 979-986.	1.7	26
107	Engineering a Function into a Glycosyltransferase. <i>Chemistry and Biology</i> , 2009, 16, 28-35.	6.2	38
108	Function of <i>lanI</i> in regulation of landomycin A biosynthesis in <i>Streptomyces cyanogenus</i> S136 and cross-complementation studies with <i>Streptomyces</i> antibiotic regulatory proteins encoding genes. <i>Archives of Microbiology</i> , 2008, 189, 111-120.	1.0	27

#	ARTICLE	IF	CITATIONS
109	Functional expression of the Cre recombinase in actinomycetes. <i>Applied Microbiology and Biotechnology</i> , 2008, 78, 1065-1070.	1.7	56
110	Aranciamycin analogs generated by combinatorial biosynthesis show improved antitumor activity. <i>Applied Microbiology and Biotechnology</i> , 2008, 80, 15-19.	1.7	17
111	Features and applications of bacterial glycosyltransferases: current state and prospects. <i>Applied Microbiology and Biotechnology</i> , 2008, 80, 945-952.	1.7	38
112	Coordination of export and glycosylation of landomycins in <i>Streptomyces cyanogenus</i> S136. <i>FEMS Microbiology Letters</i> , 2008, 285, 195-202.	0.7	41
113	Marker removal from actinomycetes genome using Flp recombinase. <i>Gene</i> , 2008, 419, 43-47.	1.0	52
114	Glycosyltransferases, Important Tools for Drug Design. <i>Current Topics in Medicinal Chemistry</i> , 2008, 8, 680-709.	1.0	70
115	Deoxysugars in Bioactive Natural Products: Development of Novel Derivatives by Altering the Sugar Pattern. <i>Current Topics in Medicinal Chemistry</i> , 2008, 8, 710-724.	1.0	43
116	Type II polyketide synthases: gaining a deeper insight into enzymatic teamwork. <i>Natural Product Reports</i> , 2007, 24, 162-190.	5.2	513
117	Generation of New Landomycins with Altered Saccharide Patterns through Over-expression of the Glycosyltransferase GenelanGT3 in the Biosynthetic Gene Cluster of Landomycin A in <i>Streptomyces cyanogenus</i> S-136. <i>ChemBioChem</i> , 2007, 8, 83-88.	1.3	51
118	Cloning and Heterologous Expression of the Aranciamycin Biosynthetic Gene Cluster Revealed a New Flexible Glycosyltransferase. <i>ChemBioChem</i> , 2007, 8, 599-602.	1.3	48
119	A strategy for cloning glycosyltransferase genes involved in natural product biosynthesis. <i>Applied Microbiology and Biotechnology</i> , 2007, 75, 1367-1375.	1.7	18
120	Cloning, purification and characterization of a functional anthracycline glycosyltransferase. <i>Journal of Biotechnology</i> , 2006, 125, 425-433.	1.9	10
121	A putative proteinase gene is involved in regulation of landomycin E biosynthesis in <i>Streptomyces globisporus</i> 1912. <i>FEMS Microbiology Letters</i> , 2006, 255, 280-285.	0.7	9
122	Biosynthesis of the Terpene Phenalinolactone in <i>Streptomyces</i> sp. TÅ¼46071: Analysis of the Gene Cluster and Generation of Derivatives. <i>Chemistry and Biology</i> , 2006, 13, 365-377.	6.2	71
123	On the Acceptor Substrate of C-Glycosyltransferase UrdGT2: Three Prejadomycin C-Glycosides from an Engineered Mutant of <i>Streptomyces globisporus</i> 1912 Δ IndE(urdGT2). <i>Angewandte Chemie - International Edition</i> , 2006, 45, 7842-7846.	7.2	36
124	Iteratively Acting Glycosyltransferases Involved in the Hexasaccharide Biosynthesis of Landomycin A. <i>Chemistry and Biology</i> , 2005, 12, 725-729.	6.2	54
125	It works: combinatorial biosynthesis for generating novel glycosylated compounds. <i>Molecular Microbiology</i> , 2005, 58, 3-5.	1.2	18
126	Generation of Novel Landomycins M and O through Targeted Gene Disruption. <i>ChemBioChem</i> , 2005, 6, 675-678.	1.3	43

#	ARTICLE	IF	CITATIONS
127	LanGT2 Catalyzes the First Glycosylation Step during Landomycin A Biosynthesis. <i>ChemBioChem</i> , 2005, 6, 1406-1410.	1.3	47
128	LanV, a Bifunctional Enzyme: Aromatase and Ketoreductase during Landomycin A Biosynthesis. <i>ChemBioChem</i> , 2005, 6, 2312-2315.	1.3	19
129	Identification of the Function of GeneLndM2Encoding a Bifunctional Oxygenase-Reductase Involved in the Biosynthesis of the Antitumor Antibiotic Landomycin E by <i>Streptomyces globisporus</i> 1912 Supports the Originally Assigned Structure for Landomycinone. <i>Journal of Organic Chemistry</i> , 2005, 70, 631-638.	1.7	52
130	Glycosyltransferases involved in the biosynthesis of biologically active natural products that contain oligosaccharides. <i>Molecular BioSystems</i> , 2005, 1, 117.	2.9	51
131	Expression of the landomycin biosynthetic gene cluster in a PKS mutant of <i>Streptomyces fradiae</i> is dependent on the coexpression of a putative transcriptional activator gene. <i>FEMS Microbiology Letters</i> , 2004, 230, 91-97.	0.7	43
132	Function of lanGT3, a Glycosyltransferase Gene Involved in Landomycin A Biosynthesis. <i>ChemBioChem</i> , 2004, 5, 1567-1570.	1.3	29
133	Generation of New Landomycins by Combinatorial Biosynthetic Manipulation of the LndGT4 Gene of the Landomycin E Cluster in <i>S. globisporus</i> . <i>Chemistry and Biology</i> , 2004, 11, 547-555.	6.2	63
134	Generation of <i>Streptomyces globisporus</i> SMY622 Strain with Increased Landomycin E Production and It's Initial Characterization. <i>Journal of Antibiotics</i> , 2004, 57, 383-389.	1.0	20