Mohamed A Marahiel

List of Publications by Year
in descending order

238 papers	26,781 citations	4960 84 h-index	6996 154 g-index
252 all docs	252 docs citations	252 times ranked	15358 citing authors

#	Article	IF	Citations
1	Communication Breakdown: Dissecting the COM Interfaces between the Subunits of Nonribosomal Peptide Synthetases. ACS Catalysis, 2021, 11, 10802-10813.	11.2	14
2	Genome Mining and Heterologous Expression Reveal Two Distinct Families of Lasso Peptides Highly Conserved in Endofungal Bacteria. ACS Chemical Biology, 2020, 15, 1169-1176.	3.4	20
3	Enterobacter bugandensis: a novel enterobacterial species associated with severe clinical infection. Scientific Reports, 2018, 8, 5392.	3.3	61
4	Structural Basis for Natural Product Selection and Export by Bacterial ABC Transporters. ACS Chemical Biology, 2018, 13, 1598-1609.	3.4	33
5	Structural and mutational analysis of the nonribosomal peptide synthetase heterocyclization domain provides insight into catalysis. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 95-100.	7.1	75
6	Signatures of Mechanically Interlocked Topology of Lasso Peptides by Ion Mobilityâ€“Mass Spectrometry: Lessons from a Collection of Representatives. Journal of the American Society for Mass Spectrometry, 2017, 28, 315-322.	2.8	17
7	Crystal Structure of Bacillus subtilis Cysteine Desulfurase SufS and Its Dynamic Interaction with Frataxin and Scaffold Protein SufU. PLoS ONE, 2016, 11, e0158749.	2.5	24
8	Insights into the Unique Phosphorylation of the Lasso Peptide Paeninodin. Journal of Biological Chemistry, 2016, 291, 13662-13678.	3.4	100
9	Dual substrateâ€controlled kinase activity leads to polyphosphorylated lasso peptides. FEBS Letters, 2016, 590, 3323-3334.	2.8	23
10	Structure and Mechanism of the Sphingopyxinâ€…I Lasso Peptide Isopeptidase. Angewandte Chemie - International Edition, 2016, 55, 12717-12721.	13.8	27
11	Structure and Mechanism of the Sphingopyxinâ€…I Lasso Peptide Isopeptidase. Angewandte Chemie, 2016, 128, 12909-12913.	2.0	2
12	The B1 Protein Guides the Biosynthesis of a Lasso Peptide. Scientific Reports, 2016, 6, 35604.	3.3	48
13	The ring residue proline 8 is crucial for the thermal stability of the lasso peptide caulosegnin II. Molecular BioSystems, 2016, 12, 1106-1109.	2.9	35
14	A structural model for multimodular NRPS assembly lines. Natural Product Reports, 2016, 33, 136-140.	10.3	110
15	Rational and combinatorial tailoring of bioactive cyclic dipeptides. Frontiers in Microbiology, 2015, 6, 785.	3.5	67
16	Molecular Insights into Frataxin-Mediated Iron Supply for Heme Biosynthesis in Bacillus subtilis. PLoS ONE, 2015, 10, e0122538.	2.5	23
17	The PqqD homologous domain of the radical SAM enzyme ThnB is required for thioether bond formation during thurincin H maturation. FEBS Letters, 2015, 589, 1802-1806.	2.8	60
18	Ion Mobilityâ€“Mass Spectrometry of Lasso Peptides: Signature of a Rotaxane Topology. Analytical Chemistry, 2015, 87, 1166-1172.	6.5	48

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19	Lasso Peptides: An Intriguing Class of Bacterial Natural Products. Accounts of Chemical Research, 2015, 48, 1909-1919.	15.6	290
20	Catalytic mechanism and allosteric regulation of an oligomeric (p)ppGpp synthetase by an alarmone. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 13348-13353.	7.1	111
21	Peptide Antibiotics. , 2014, , 897-916.		45
22	Xanthomoninsâ€…Iâ€“III: A New Class of Lasso Peptides with a Sevenâ€Residue Macrolactam Ring. Angewandte Chemie - International Edition, 2014, 53, 2230-2234.	13.8	72
23	The tRNA-Dependent Biosynthesis of Modified Cyclic Dipeptides. International Journal of Molecular Sciences, 2014, 15, 14610-14631.	4.1	68
24	Crystal Structure of a PCP/Sfp Complex Reveals the Structural Basis for Carrier Protein Posttranslational Modification. Chemistry and Biology, 2014, 21, 552-562.	6.0	37
25	Characterization of caulonodin lasso peptides revealed unprecedented N-terminal residues and a precursor motif essential for peptide maturation. Chemical Science, 2014, 5, 4032-4043.	7.4	40
26	Rational Improvement of the Affinity and Selectivity of Integrin Binding of Grafted Lasso Peptides. Journal of Medicinal Chemistry, 2014, 57, 5829-5834.	6.4	68
27	Structural Characterization of the Heterobactin Siderophores from Rhodococcus erythropolis PR4 and Elucidation of Their Biosynthetic Machinery. Journal of Natural Products, 2013, 76, 2282-2290.	3.0	52
28	Caulosegnins Iâ€“III: A Highly Diverse Group of Lasso Peptides Derived from a Single Biosynthetic Gene Cluster. Journal of the American Chemical Society, 2013, 135, 210-222.	13.7	99
29	Ribosomally synthesized and post-translationally modified peptide natural products: overview and recommendations for a universal nomenclature. Natural Product Reports, 2013, 30, 108-160.	10.3	1,692
30	The Bacillus subtilis EfeUOB transporter is essential for high-affinity acquisition of ferrous and ferric iron. Biochimica Et Biophysica Acta - Molecular Cell Research, 2013, 1833, 2267-2278.	4.1	84
31	Radical S-adenosylmethionine enzyme catalyzed thioether bond formation in sactipeptide biosynthesis. Current Opinion in Chemical Biology, 2013, 17, 605-612.	6.1	78
32	Uptake of xenosiderophores in Bacillus subtilis occurs with high affinity and enhances the folding stabilities of substrate binding proteins. FEBS Letters, 2013, 587, 206-213.	2.8	24
33	The Astexin-1 Lasso Peptides: Biosynthesis, Stability, and Structural Studies. Chemistry and Biology, 2013, 20, 558-569.	6.0	79
34	Insights into the Generation of Structural Diversity in a tRNA-Dependent Pathway for Highly Modified Bioactive Cyclic Dipeptides. Chemistry and Biology, 2013, 20, 828-838.	6.0	62
35	A tRNA-Dependent Two-Enzyme Pathway for the Generation of Singly and Doubly Methylated Ditryptophan 2,5-Diketopiperazines. Biochemistry, 2013, 52, 4274-4283.	2.5	67
36	Two [4Fe-4S] Clusters Containing Radical SAM Enzyme SkfB Catalyze Thioether Bond Formation during the Maturation of the Sporulation Killing Factor. Journal of the American Chemical Society, 2013, 135, 959-962.	13.7	89

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37	Nonribosomal Peptide Synthesis. , 2013, , 138-149.		7
38	Lasso peptides from proteobacteria: Genome mining employing heterologous expression and mass spectrometry. Biopolymers, 2013, 100, 527-542.	2.4	111
39	The radical SAM enzyme AlbA catalyzes thioether bond formation in subtilosin A. Nature Chemical Biology, 2012, 8, 350-357.	8.0	166
40	An Enzymatic Pathway for the Biosynthesis of the Formylhydroxyornithine Required for Rhodochelin Iron Coordination. Biochemistry, 2012, 51, 3059-3066.	2.5	31
41	Determination of Peptide Topology through Time-Resolved Double-Resonance under Electron Capture Dissociation Conditions. Analytical Chemistry, 2012, 84, 4957-4964.	6.5	20
42	Isolation, Structure Elucidation, and Biosynthesis of an Unusual Hydroxamic Acid Ester-Containing Siderophore from Actinosynnema mirum. Journal of Natural Products, 2012, 75, 905-914.	3.0	52
43	NMR as an Effective Tool for the Structure Determination of Lasso Peptides. ChemBioChem, 2012, 13, 621-625.	2.6	37
44	Dissecting the Maturation Steps of the Lasso Peptide Microcin J25 in vitro. ChemBioChem, 2012, 13, 1046-1052.	2.6	106
45	Exploring the mechanism of lipid transfer during biosynthesis of the acidic lipopeptide antibiotic CDA. FEBS Letters, 2012, 586, 283-288.	2.8	26
46	Ribosomeâ€independent biosynthesis of biologically active peptides: Application of synthetic biology to generate structural diversity. FEBS Letters, 2012, 586, 2065-2075.	2.8	50
47	Consecutive Enzymatic Modification of Ornithine Generates the Hydroxamate Moieties of the Siderophore Erythrochelin. Biochemistry, 2011, 50, 6073-6080.	2.5	20
48	Identification and Characterization of a Novel-type Ferric Siderophore Reductase from a Gram-positive Extremophile. Journal of Biological Chemistry, 2011, 286, 2245-2260.	3.4	26
49	The Siderophore-Interacting Protein YqjH Acts as a Ferric Reductase in Different Iron Assimilation Pathways of Escherichia coli. Biochemistry, 2011, 50, 10951-10964.	2.5	70
50	Biosynthesis of the Siderophore Rhodochelin Requires the Coordinated Expression of Three Independent Gene Clusters in Rhodococcus jostii RHA1. Journal of the American Chemical Society, 2011, 133, 4587-4595.	13.7	55
51	A Four-Enzyme Pathway for 3,5-Dihydroxy-4-methylanthranilic Acid Formation and Incorporation into the Antitumor Antibiotic Sibiromycin. Biochemistry, 2011, 50, 5680-5692.	2.5	29
52	The Antibacterial Threaded-lasso Peptide Capistruin Inhibits Bacterial RNA Polymerase. Journal of Molecular Biology, 2011, 412, 842-848.	4.2	82
53	Environmental Salinity Determines the Specificity and Need for Tat-Dependent Secretion of the YwbN Protein in Bacillus subtilis. PLoS ONE, 2011, 6, e18140.	2.5	36
54	Mechanistic characterization of sulfur transfer from cysteine desulfurase SufS to the iron-sulfur scaffold SufU in Bacillus subtilis. FEBS Letters, 2011, 585, 465-470.	2.8	28

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55	Topoisomer Differentiation of Molecular Knots by FTICR MS: Lessons from Class II Lasso Peptides. Journal of the American Society for Mass Spectrometry, 2011, 22, 467-479.	2.8	38
56	Introducing Lasso Peptides as Molecular Scaffolds for Drug Design: Engineering of an Integrin Antagonist. Angewandte Chemie - International Edition, 2011, 50, 8714-8717.	13.8	108
57	The Frataxin Homologue Fra Plays a Key Role in Intracellular Iron Channeling in Bacillus subtilis. ChemBioChem, 2011, 12, 2052-2061.	2.6	17
58	Identification and Characterization of the Lysobactin Biosynthetic Gene Cluster Reveals Mechanistic Insights into an Unusual Termination Module Architecture. Chemistry and Biology, 2011, 18, 655-664.	6.0	68
59	The Siderophore Binding Protein FeuA Shows Limited Promiscuity toward Exogenous Triscatecholates. Chemistry and Biology, 2011, 18, 907-919.	6.0	57
60	Nonribosomal peptide synthetases: structures and dynamics. Current Opinion in Structural Biology, 2010, 20, 234-240.	5.7	366
61	The glucagon receptor antagonist Blâ€32169 constitutes a new class of lasso peptides. FEBS Letters, 2010, 584, 785-789.	2.8	70
62	Direct Identification of a Siderophore Import Protein Using Synthetic Petrobactin Ligands. Angewandte Chemie - International Edition, 2010, 49, 10210-10213.	13.8	20
63	Functional Dissection of Surfactin Synthetase Initiation Module Reveals Insights into the Mechanism of Lipoinitiation. Chemistry and Biology, 2010, 17, 872-880.	6.0	113
64	Erythrochelin â€“ a hydroxamateâ€type siderophore predicted from the genome of Saccharopolysporaâ€ferythraea. FEBS Journal, 2010, 277, 663-676.	4.7	50
65	Elucidation of the complete ferrichrome A biosynthetic pathway in Ustilago maydis. Molecular Microbiology, 2010, 75, 1260-1271.	2.5	68
66	Copper Stress Affects Iron Homeostasis by Destabilizing Iron-Sulfur Cluster Formation in Bacillus subtilis. Journal of Bacteriology, 2010, 192, 2512-2524.	2.2	200
67	SufU Is an Essential Iron-Sulfur Cluster Scaffold Protein in Bacillus subtilis. Journal of Bacteriology, 2010, 192, 1643-1651.	2.2	83
68	Daptomycin, a Bacterial Lipopeptide Synthesized by a Nonribosomal Machinery. Journal of Biological Chemistry, 2010, 285, 27501-27508.	3.4	176
69	The reversible macrocyclization of Tyrocidine A aldehyde: a hemiaminal reminiscent of the tetrahedral intermediate of macrolactamization. Organic and Biomolecular Chemistry, 2010, 8, 559-563.	2.8	18
70	Copper Acquisition Is Mediated by YcnJ and Regulated by YcnK and CsoR in Bacillus subtilis. Journal of Bacteriology, 2009, 191, 2362-2370.	2.2	88
71	The Structural Diversity of Acidic Lipopeptide Antibiotics. ChemBioChem, 2009, 10, 607-616.	2.6	113
72	The Synechocystis sp. PCC6803 Sfpâ€Type Phosphopantetheinyl Transferase Does Not Possess Characteristic Broadâ€Range Activity. ChemBioChem, 2009, 10, 1869-1877.	2.6	18

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73	Structural Basis and Stereochemistry of Triscatecholate Siderophore Binding by FeuA. Angewandte Chemie - International Edition, 2009, 48, 7924-7927.	13.8	51
74	Working outside the proteinâ€synthesis rules: insights into nonâ€ribosomal peptide synthesis. Journal of Peptide Science, 2009, 15, 799-807.	1.4	119
75	TioSâ€ƒTâ€TEâ€ƒâ€“â€ƒa prototypical thioesterase responsible for cyclodimerization of the quinolineâ€and quinoxalineâ€type class of chromodepsipeptides. FEBS Journal, 2009, 276, 1641-1653.	4.7	21
76	Structural basis for the erythroâ€stereospecificity of the <scp>l</scp>â€arginine oxygenase VioC in viomycin biosynthesis. FEBS Journal, 2009, 276, 3669-3682.	4.7	64
77	Crosslinking Studies of Protein-Protein Interactions in Nonribosomal Peptide Biosynthesis. Chemistry and Biology, 2009, 16, 372-381.	6.0	42
78	Insights into the Biosynthesis and Stability of the Lasso Peptide Capistruin. Chemistry and Biology, 2009, 16, 1290-1298.	6.0	118
79	Stereospecific Synthesis of threo- and erythro-Î²-Hydroxyglutamic Acid During Kutzneride Biosynthesis. Journal of the American Chemical Society, 2009, 131, 13523-13530.	13.7	68
80	Post-Translational Modification and folding of A Lasso-Type Gene-Encoded Antimicrobial Peptide Require Two Enzymes only in Escherichia coli. Advances in Experimental Medicine and Biology, 2009, 611, 35-36.	1.6	7
81	Nonâ€Heme Hydroxylase Engineering For Simple Enzymatic Synthesis of <scp>L</scp>â€threoâ€Hydroxyaspartic Acid. ChemBioChem, 2008, 9, 374-376.	2.6	24
82	The Entropy Balance of Nostocyclopeptide Macrocyclization Analysed by NMR Spectroscopy. ChemBioChem, 2008, 9, 2597-2601.	2.6	16
83	Role of DptE and DptF in the lipidation reaction of daptomycin. FEBS Journal, 2008, 275, 5343-5354.	4.7	66
84	Structural basis for the selectivity of the external thioesterase of the surfactin synthetase. Nature, 2008, 454, 907-911.	27.8	112
85	Î´-Amino group hydroxylation of l-ornithine during coelichelin biosynthesis. Organic and Biomolecular Chemistry, 2008, 6, 1843.	2.8	31
86	How to tailor non-ribosomal peptide productsâ€”new clues about the structures and mechanisms of modifying enzymes. Molecular BioSystems, 2008, 4, 387.	2.9	36
87	Harnessing the Chemical Activation Inherent to Carrier Protein-Bound Thioesters for the Characterization of Lipopeptide Fatty Acid Tailoring Enzymes. Journal of the American Chemical Society, 2008, 130, 2656-2666.	13.7	32
88	Isolation and Structural Characterization of Capistruin, a Lasso Peptide Predicted from the Genome Sequence of Burkholderia thailandensis E264. Journal of the American Chemical Society, 2008, 130, 11446-11454.	13.7	220
89	Crystal Structure of the Termination Module of a Nonribosomal Peptide Synthetase. Science, 2008, 321, 659-663.	12.6	311
90	Crystal Structure of DltA. Journal of Biological Chemistry, 2008, 283, 32484-32491.	3.4	117

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91	The Major Facilitator Superfamily-Type Transporter YmfE and the Multidrug-Efflux Activator Mta Mediate Bacillibactin Secretion in Bacillus subtilis. Journal of Bacteriology, 2008, 190, 5143-5152.	2.2	51
92	Cloning and characterization of the biosynthetic gene cluster for kutznerides. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 16498-16503.	7.1	144
93	Aminoacyl-coenzyme A synthesis catalyzed by adenylation domains. FEBS Letters, 2007, 581, 905-910.	2.8	31
94	Macrocyclization strategies in polyketide and nonribosomal peptide biosynthesis. Natural Product Reports, 2007, 24, 735.	10.3	197
95	Stereospecific Enzymatic Transformation of Î±-Ketoglutarate to (2S,3R)-3-Methyl Glutamate during Acidic Lipopeptide Biosynthesis. Journal of the American Chemical Society, 2007, 129, 12011-12018.	13.7	75
96	Mechanistic and Structural Basis of Stereospecific CÎ²-Hydroxylation in Calcium-Dependent Antibiotic, a Daptomycin-Type Lipopeptide. ACS Chemical Biology, 2007, 2, 187-196.	3.4	107
97	Siderophore-Based Iron Acquisition and Pathogen Control. Microbiology and Molecular Biology Reviews, 2007, 71, 413-451.	6.6	1,342
98	Synthesis of a 2-indolylphosphonamide derivative with inhibitory activity against yersiniabactin biosynthesis. Tetrahedron Letters, 2007, 48, 6080-6083.	1.4	26
99	Where chemistry meets biology: the chemoenzymatic synthesis of nonribosomal peptides and polyketides. Current Opinion in Biotechnology, 2007, 18, 513-520.	6.6	55
100	The Iterative Gramicidin S Thioesterase Catalyzes Peptide Ligation and Cyclization. Chemistry and Biology, 2007, 14, 13-22.	6.0	81
101	Characterization of a mutation in the acetolactate synthase ofBacillus subtilisthat causes a cold-sensitive phenotype. FEMS Microbiology Letters, 2007, 272, 30-34.	1.8	9
102	Detection of nonribosomal peptide synthetase genes inXylariasp. BCC1067 and cloning ofXyNRPSA. FEMS Microbiology Letters, 2007, 274, 260-268.	1.8	5
103	Structural and Functional Insights into a Peptide Bond-Forming Bidomain from a Nonribosomal Peptide Synthetase. Structure, 2007, 15, 781-792.	3.3	152
104	A biosynthetic gene cluster for a secreted cellobiose lipid with antifungal activity from Ustilago maydis. Molecular Microbiology, 2007, 66, 525-533.	2.5	148
105	Conformational Switches Modulate Protein Interactions in Peptide Antibiotic Synthetases. Science, 2006, 312, 273-276.	12.6	149
106	Peptide Macrocyclization:Â The Reductase of the Nostocyclopeptide Synthetase Triggers the Self-Assembly of a Macrocyclic Imine. Journal of the American Chemical Society, 2006, 128, 16478-16479.	13.7	70
107	Nonribosomally Synthesized Microbial Macrocyclic Peptides. , 2006, , 89-96.		2
108	Cold-Induced Putative DEAD Box RNA Helicases CshA and CshB Are Essential for Cold Adaptation and Interact with Cold Shock Protein B in Bacillus subtilis. Journal of Bacteriology, 2006, 188, 240-248.	2.2	114

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109	Rational Design of Bacitracin A Derivatives by Incorporating Natural Product Derived Heterocycles. Journal of the American Chemical Society, 2006, 128, 10513-10520.	13.7	39
110	Chemoenzymatic Design of Acidic Lipopeptide Hybrids:â€‰ New Insights into the Structureâ^'Activity Relationship of Daptomycin and A54145. Biochemistry, 2006, 45, 10474-10481.	2.5	52
111	Formylation Domain:Â An Essential Modifying Enzyme for the Nonribosomal Biosynthesis of Linear Gramicidin. Journal of the American Chemical Society, 2006, 128, 7406-7407.	13.7	46
112	The Thioesterase Domain of the Fengycin Biosynthesis Cluster: A Structural Base for the Macrocyclization of a Non-ribosomal Lipopeptide. Journal of Molecular Biology, 2006, 359, 876-889.	4.2	110
113	Histidine 109 in peptidyl-prolyl cis-trans isomerase of Bacillus subtilis plays an important role in catalysis and in cyclosporin A binding. FEMS Microbiology Letters, 2006, 154, 139-144.	1.8	4
114	Inhibition of aryl acid adenylation domains involved in bacterial siderophore synthesis. FEBS Journal, 2006, 273, 409-419.	4.7	84
115	Ferriâ€bacillibactin uptake and hydrolysis in Bacillus subtilis. Molecular Microbiology, 2006, 61, 1413-1427.	2.5	160
116	Solvent Engineering Substantially Enhances the Chemoenzymatic Production of Surfactin. ChemBioChem, 2006, 7, 595-597.	2.6	10
117	Impact of Epimerization Domains on the Intermodular Transfer of Enzyme-Bound Intermediates in Nonribosomal Peptide Synthesis. ChemBioChem, 2006, 7, 1807-1814.	2.6	28
118	Iron Starvation Triggers the Stringent Response and Induces Amino Acid Biosynthesis for Bacillibactin Production in Bacillus subtilis. Journal of Bacteriology, 2006, 188, 8655-8657.	2.2	39
119	Sigma L Is Important for Cold Shock Adaptation of Bacillus subtilis. Journal of Bacteriology, 2006, 188, 3130-3133.	2.2	30
120	Identification of a Gene Cluster for Biosynthesis of Mannosylerythritol Lipids in the Basidiomycetous Fungus Ustilago maydis. Applied and Environmental Microbiology, 2006, 72, 5469-5477.	3.1	145
121	Chemoenzymatic and Template-Directed Synthesis of Bioactive Macrocyclic Peptides. Microbiology and Molecular Biology Reviews, 2006, 70, 121-146.	6.6	199
122	Peptidantibiotika vom molekularen FlieÃŸband. Nachrichten Aus Der Chemie, 2005, 53, 507-513.	0.0	4
123	Fluorescence Resonance Energy Transfer as a Probe of Peptide Cyclization Catalyzed by Nonribosomal Thioesterase Domains. Chemistry and Biology, 2005, 12, 873-881.	6.0	17
124	Inhibition of the D-alanine:D-alanyl carrier protein ligase from Bacillus subtilis increases the bacterium's susceptibility to antibiotics that target the cell wall. FEBS Journal, 2005, 272, 2993-3003.	4.7	93
125	Utility of epimerization domains for the redesign of nonribosomal peptide synthetases. FEBS Journal, 2005, 272, 4506-4520.	4.7	37
126	Molecular Mechanisms Underlying Nonribosomal Peptide Synthesis: Approaches to New Antibiotics. ChemInform, 2005, 36, no.	0.0	3

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127	Chemoenzymatic Approach to Enantiopure Streptogramin B Variants:Â Characterization of Stereoselective Pristinamycin I Cyclase fromStreptomyces pristinaespiralis. Journal of the American Chemical Society, 2005, 127, 9571-9580.	13.7	32
128	Synthesis of Linear Gramicidin Requires the Cooperation of Two Independent Reductases. Biochemistry, 2005, 44, 8507-8513.	2.5	29
129	Molecular Mechanisms Underlying Nonribosomal Peptide Synthesis:â€‰ Approaches to New Antibiotics. Chemical Reviews, 2005, 105, 715-738.	47.7	523
130	Reactions Catalyzed by Mature and Recombinant Nonribosomal Peptide Synthetases. Methods in Enzymology, 2004, 388, 293-315.	1.0	44
131	The Linear Pentadecapeptide Gramicidin Is Assembled by Four Multimodular Nonribosomal Peptide Synthetases That Comprise 16 Modules with 56 Catalytic Domains. Journal of Biological Chemistry, 2004, 279, 7413-7419.	3.4	92
132	Mutational analysis of a type II thioesterase associated with nonribosomal peptide synthesis. FEBS Journal, 2004, 271, 1536-1545.	0.2	32
133	Role of the Bacillus subtilis fatty acid desaturase in membrane adaptation during cold shock. Molecular Microbiology, 2004, 39, 1321-1329.	2.5	100
134	Genetic evidence for the temperature-sensing ability of the membrane domain of the Bacillus subtilis histidine kinase DesK. FEMS Microbiology Letters, 2004, 230, 41-46.	1.8	16
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136	Enzymatic Cyclisation of Peptidomimetics with Incorporated (E)-Alkene Dipeptide Isosteres. ChemBioChem, 2004, 5, 1000-1003.	2.6	18
137	Rational Design of a Bimodular Model System for the Investigation of Heterocyclization in Nonribosomal Peptide Biosynthesis. Chemistry and Biology, 2004, 11, 261-271.	6.0	55
138	Mutational Analysis of Peptidyl Carrier Protein and Acyl Carrier Protein Synthase Unveils Residues Involved in Proteinâ'Protein Recognitionâ€. Biochemistry, 2004, 43, 8946-8956.	2.5	32
139	Chemo- and Regioselective Peptide Cyclization Triggered by the N-Terminal Fatty Acid Chain Length:Â The Recombinant Cyclase of the Calcium-Dependent Antibiotic fromStreptomyces coelicolorâ€. Biochemistry, 2004, 43, 2915-2925.	2.5	53
140	Structure-Based Mutational Analysis of the 4â€˜-Phosphopantetheinyl Transferases Sfp from Bacillus subtilis:â€‰ Carrier Protein Recognition and Reaction Mechanism,. Biochemistry, 2004, 43, 4128-4136.	2.5	62
141	Biosynthesis of Nonribosomal Peptides. Annual Review of Microbiology, 2004, 58, 453-488.	7.3	775
142	Synthesis and Derivatization of Daptomycin:ÂÂA Chemoenzymatic Route to Acidic Lipopeptide Antibiotics. Journal of the American Chemical Society, 2004, 126, 17025-17031.	13.7	118
143	Nonribosomal Peptides: From Genes to Products. ChemInform, 2003, 34, no.	0.0	0
144	Aminoacyl Adenylate Substrate Analogues for the Inhibition of Adenylation Domains of Nonribosomal Peptide Synthetases. ChemBioChem, 2003, 4, 903-906.	2.6	66

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145	Fourier-transform mass spectrometry for detection of thioester-bound intermediates in unfractionated proteolytic mixtures of 80 and 191 kDa portions of Bacitracin A synthetase. Analytica Chimica Acta, 2003, 496, 217-224.	5.4	10
146	Construction of hybrid peptide synthetases for the production of alpha-l-aspartyl-l-phenylalanine, a precursor for the high-intensity sweetener aspartame. FEBS Journal, 2003, 270, 4555-4563.	0.2	45
147	Chirality of Peptide Bond-Forming Condensation Domains in Nonribosomal Peptide Synthetases:Â The C5Domain of Tyrocidine Synthetase Is aDCLCatalystâ€. Biochemistry, 2003, 42, 12095-12104.	2.5	83
148	Systematic and Quantitative Analysis of Proteinâ^'Protein Recognition between Nonribosomal Peptide Synthetases Investigated in the Tyrocidine Biosynthetic Templateâ€. Biochemistry, 2003, 42, 5114-5124.	2.5	26
149	Nonribosomal peptides: from genes to products. Natural Product Reports, 2003, 20, 275.	10.3	503
150	Loading Peptidyl-Coenzyme A onto Peptidyl Carrier Proteins: A Novel Approach in Characterizing Macrocyclization by Thioesterase Domains. Journal of the American Chemical Society, 2003, 125, 10862-10866.	13.7	66
151	Learning from Nature's Drug Factories: Nonribosomal Synthesisof MacrocyclicPeptides. Journal of Bacteriology, 2003, 185, 7036-7043.	2.2	103
152	Bacterial Cold Shock Responses. Science Progress, 2003, 86, 9-75.	1.9	171
153	Ebony, a Novel Nonribosomal Peptide Synthetase for Î²-Alanine Conjugation with Biogenic Amines in Drosophila. Journal of Biological Chemistry, 2003, 278, 41160-41166.	3.4	123
154	Coping with the cold: the cold shock response in the Gram-positive soil bacterium Bacillus subtilis. Philosophical Transactions of the Royal Society B: Biological Sciences, 2002, 357, 895-907.	4.0	55
155	Characterization of a New Type of Phosphopantetheinyl Transferase for Fatty Acid and Siderophore Synthesis in Pseudomonas aeruginosa. Journal of Biological Chemistry, 2002, 277, 50293-50302.	3.4	88
156	Genomewide Transcriptional Analysis of the Cold Shock Response in Bacillus subtilis. Journal of Bacteriology, 2002, 184, 6395-6402.	2.2	113
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