

Yasushi Ogasawara

List of Publications by Year in descending order

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56
papers

1,106
citations

394390

19
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454934

30
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58
all docs

58
docs citations

58
times ranked

1196
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Cyclopropane Formation in the Biosyntheses of Hormaomycins and Belactosins: Sequential Nitration and Cyclopropanation by Metalloenzymes. <i>Angewandte Chemie</i> , 2022, 134, .	2.0	3
2	Identification of Cyclopropane Formation in the Biosyntheses of Hormaomycins and Belactosins: Sequential Nitration and Cyclopropanation by Metalloenzymes. <i>Angewandte Chemie - International Edition</i> , 2022, 61, e202113189.	13.8	18
3	Biosynthetic Gene Cluster of Linaridin Peptides Contains Epimerase Gene. <i>ChemBioChem</i> , 2022, 23, .	2.6	10
4	Identification of the peptide epimerase MslH responsible for d-amino acid introduction at the C-terminus of ribosomal peptides. <i>Chemical Science</i> , 2021, 12, 2567-2574.	7.4	13
5	Discovery of an alternative pathway of peptidoglycan biosynthesis: A new target for pathway specific inhibitors. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2021, 48, .	3.0	4
6	Identification of pulvomycin as an inhibitor of the futasoline pathway. <i>Journal of Antibiotics</i> , 2021, 74, 825-829.	2.0	2
7	Identification of actinomycin D as a specific inhibitor of the alternative pathway of peptidoglycan biosynthesis. <i>Journal of Antibiotics</i> , 2020, 73, 125-127.	2.0	10
8	High Production of Ergothioneine in <i>Escherichia coli</i> using the Sulfoxide Synthase from <i>Methylobacterium</i> strains. <i>Journal of Agricultural and Food Chemistry</i> , 2020, 68, 6390-6394.	5.2	16
9	Off-Loading Mechanism of Products in Polyunsaturated Fatty Acid Synthases. <i>ACS Chemical Biology</i> , 2020, 15, 651-656.	3.4	11
10	Characterization of the coformycin biosynthetic gene cluster in <i>Streptomyces kaniharaensis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 10265-10270.	7.1	8
11	Recent advances in functional analysis of polyunsaturated fatty acid synthases. <i>Current Opinion in Chemical Biology</i> , 2020, 59, 30-36.	6.1	14
12	In vitro characterization of MitE and MitB: Formation of N-acetylglucosaminyl-3-amino-5-hydroxybenzoyl-MmcB as a key intermediate in the biosynthesis of antitumor antibiotic mitomycins. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2019, 29, 2076-2078.	2.2	6
13	Identification of the C-Glycoside Synthases during Biosynthesis of the PyrazoleC-Nucleosides Formycin and Pyrazofurin. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 16512-16516.	13.8	25
14	Subtle Control of Carbon Chain Length in Polyunsaturated Fatty Acid Synthases. <i>ACS Chemical Biology</i> , 2019, 14, 2553-2556.	3.4	9
15	Identification of the C-Glycoside Synthases during Biosynthesis of the PyrazoleC-Nucleosides Formycin and Pyrazofurin. <i>Angewandte Chemie</i> , 2019, 131, 16664-16668.	2.0	6
16	The Amipurimycin and Miharamycin Biosynthetic Gene Clusters: Unraveling the Origins of 2-Aminopurinylyl Peptidyl Nucleoside Antibiotics. <i>Journal of the American Chemical Society</i> , 2019, 141, 14152-14159.	13.7	25
17	Involvement of Peptide Epimerization in Poly- γ -glutamic Acid Biosynthesis. <i>Organic Letters</i> , 2019, 21, 3972-3975.	4.6	11
18	Control Mechanism for Carbon Chain Length in Polyunsaturated Fatty Acid Synthases. <i>Angewandte Chemie</i> , 2019, 131, 6677-6682.	2.0	2

#	ARTICLE	IF	CITATIONS
19	Control Mechanism for Carbon-Chain Length in Polyunsaturated Fatty Acid Synthases. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 6605-6610.	13.8	31
20	Amino Acid Residues Recognizing Isomeric Glutamate Substrates in UDP-N-acetylmuramic acid-alanine-glutamate Synthetases. <i>ACS Chemical Biology</i> , 2019, 14, 975-978.	3.4	5
21	Identification of the Formycin A Biosynthetic Gene Cluster from <i>Streptomyces kaniharaensis</i> Illustrates the Interplay between Biological Pyrazolopyrimidine Formation and <i>de Novo</i> Purine Biosynthesis. <i>Journal of the American Chemical Society</i> , 2019, 141, 6127-6131.	13.7	38
22	New enzymes for peptide biosynthesis in microorganisms. <i>Bioscience, Biotechnology and Biochemistry</i> , 2019, 83, 589-597.	1.3	3
23	Searching for potent and specific antibiotics against pathogenic <i>Helicobacter</i> and <i>Campylobacter</i> strains. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2019, 46, 409-414.	3.0	3
24	Control Mechanism for <i>cis</i> Double Bond Formation by Polyunsaturated Fatty Acid Synthases. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 2326-2330.	13.8	33
25	Control Mechanism for <i>cis</i> Double Bond Formation by Polyunsaturated Fatty Acid Synthases. <i>Angewandte Chemie</i> , 2019, 131, 2348-2352.	2.0	3
26	Enzymatic Formation of a Skipped Methyl-Substituted Octaprenyl Side Chain of Longestin (KS505a): Involvement of Homo-PP as a Common Extender Unit. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 6629-6632.	13.8	27
27	Enzymatic Formation of a Skipped Methyl-Substituted Octaprenyl Side Chain of Longestin (KS505a): Involvement of Homo-PP as a Common Extender Unit. <i>Angewandte Chemie</i> , 2018, 130, 6739-6742.	2.0	7
28	Functional analysis of methyltransferases participating in streptothricin-related antibiotic biosynthesis. <i>Journal of Bioscience and Bioengineering</i> , 2018, 125, 148-154.	2.2	1
29	Total Biosynthesis of Brassicicenes: Identification of a Key Enzyme for Skeletal Diversification. <i>Organic Letters</i> , 2018, 20, 6178-6182.	4.6	21
30	Biosynthetic Gene Cluster of a <i>d</i> -Tryptophan-Containing Lasso Peptide, MS271. <i>ChemBioChem</i> , 2018, 19, 2045-2048.	2.6	40
31	Peptide Epimerization Machineries Found in Microorganisms. <i>Frontiers in Microbiology</i> , 2018, 9, 156.	3.5	19
32	Aplasmomycin and boromycin are specific inhibitors of the futasoline pathway. <i>Journal of Antibiotics</i> , 2018, 71, 968-970.	2.0	22
33	æ”¼ç-šèEâEç”Ÿã;â‡ªã-ãŸç-‘ã¼¼ãfšãf-ãfãf%âE-âç%©. <i>Kagaku To Seibutsu</i> , 2018, 56, 76-78.	0.0	0
34	Biosynthesis of the Carbonylmethylene Structure Found in the Ketomemycin Class of Pseudotripeptides. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 2026-2029.	13.8	17
35	Biosynthesis of the Carbonylmethylene Structure Found in the Ketomemycin Class of Pseudotripeptides. <i>Angewandte Chemie</i> , 2017, 129, 2058-2061.	2.0	2
36	Identification and Characterization of Enzymes Catalyzing Pyrazolopyrimidine Formation in the Biosynthesis of Formycin A. <i>Organic Letters</i> , 2017, 19, 1426-1429.	4.6	20

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37	Identification of tirandamycins as specific inhibitors of the futasine pathway. <i>Journal of Antibiotics</i> , 2017, 70, 798-800.	2.0	20
38	Biosynthesis of Oligopeptides Using ATP-Grasp Enzymes. <i>Chemistry - A European Journal</i> , 2017, 23, 10714-10724.	3.3	22
39	<i>N</i> -Phenylacetylation and Nonribosomal Peptide Synthetases with Substrate Promiscuity for Biosynthesis of Heptapeptide Variants, JBIR-78 and JBIR-95. <i>ACS Chemical Biology</i> , 2017, 12, 1813-1819.	3.4	11
40	A Glycopeptidyl-Glutamate Epimerase for Bacterial Peptidoglycan Biosynthesis. <i>Journal of the American Chemical Society</i> , 2017, 139, 4243-4245.	13.7	11
41	Synthesis of Acylborons by Ozonolysis of Alkenylboronates: Preparation of an Enantioenriched Amino Acid Acylboronate. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 13847-13851.	13.8	64
42	Synthesis of Acylborons by Ozonolysis of Alkenylboronates: Preparation of an Enantioenriched Amino Acid Acylboronate. <i>Angewandte Chemie</i> , 2017, 129, 14035-14039.	2.0	33
43	Frontispiece: Biosynthesis of Oligopeptides Using ATP-Grasp Enzymes. <i>Chemistry - A European Journal</i> , 2017, 23, .	3.3	0
44	Exploring Peptide Ligase Orthologs in Actinobacteria—Discovery of Pseudopeptide Natural Products, Ketomemcins. <i>ACS Chemical Biology</i> , 2016, 11, 1686-1692.	3.4	20
45	Characterization of three amidinotransferases involved in the biosynthesis of ketomemcins. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2016, 26, 3662-3664.	2.2	9
46	Structure and activity relationships of the anti-Mycobacterium antibiotics resorcinomycin and pheganomycin. <i>Journal of Antibiotics</i> , 2016, 69, 119-120.	2.0	5
47	Expanding our Understanding of Sequence-Function Relationships of Type II Polyketide Biosynthetic Gene Clusters: Bioinformatics-Guided Identification of Frankiamicin A from <i>Frankia</i> sp. EAN1pec. <i>PLoS ONE</i> , 2015, 10, e0121505.	2.5	25
48	Identification and analysis of the resorcinomycin biosynthetic gene cluster. <i>Bioscience, Biotechnology and Biochemistry</i> , 2015, 79, 1833-1837.	1.3	12
49	High-Quality Draft Genome Sequence of <i>Actinobacterium Kibdelosporangium</i> sp. MJ126-NF4, Producer of Type II Polyketide Azicemcins, Using Illumina and PacBio Technologies. <i>Genome Announcements</i> , 2015, 3, .	0.8	6
50	GenK-Catalyzed C-6 Methylolation in the Biosynthesis of Gentamicin: Isolation and Characterization of a Cobalamin-Dependent Radical SAM Enzyme. <i>Journal of the American Chemical Society</i> , 2013, 135, 8093-8096.	13.7	110
51	Radical SAM enzymes in the biosynthesis of sugar-containing natural products. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2012, 1824, 1231-1244.	2.3	39
52	A Biosynthetic Pathway for BE-7585A, a 2-Thiosugar-Containing Angucycline-Type Natural Product. <i>Journal of the American Chemical Society</i> , 2010, 132, 7405-7417.	13.7	63
53	Biosynthesis of Spinosyn in <i>Saccharopolyspora spinosa</i> : Synthesis of Permethylated Rhamnose and Characterization of the Functions of SpnH, SpnI, and SpnK. <i>Journal of the American Chemical Society</i> , 2010, 132, 2901-2903.	13.7	46
54	Biosynthetic Studies of Aziridine Formation in Azicemcins. <i>Journal of the American Chemical Society</i> , 2009, 131, 18066-18068.	13.7	47

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55	Involvement of Glutamate Mutase in the Biosynthesis of the Unique Starter Unit of the Macrolactam Polyketide Antibiotic Vicenistatin. <i>Journal of Antibiotics</i> , 2005, 58, 468-472.	2.0	24
56	Cloning, Sequencing, and Functional Analysis of the Biosynthetic Gene Cluster of Macrolactam Antibiotic Vicenistatin in <i>Streptomyces halstedii</i> . <i>Chemistry and Biology</i> , 2004, 11, 79-86.	6.0	54