

Koji Ichinose

List of Publications by Year in descending order

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papers

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citations

471061

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#	ARTICLE	IF	CITATIONS
1	Characterization of stereospecific enoyl reductase ActVI-ORF2 for pyran ring formation in the actinorhodin biosynthesis of <i>Streptomyces coelicolor</i> A3(2). <i>Bioorganic and Medicinal Chemistry Letters</i> , 2022, 66, 128727.	1.0	3
2	Identification of a C-Glycosyltransferase Involved in Medermycin Biosynthesis. <i>ACS Chemical Biology</i> , 2021, 16, 1059-1069.	1.6	7
3	Total Synthesis of 6-Deoxydihydrokalafungin, a Key Biosynthetic Precursor of Actinorhodin, and Its Epimer. <i>Molecules</i> , 2021, 26, 6397.	1.7	3
4	Unveiling Two Consecutive Hydroxylations: Mechanisms of Aromatic Hydroxylations Catalyzed by Flavin-Dependent Monooxygenases for the Biosynthesis of Actinorhodin and Related Antibiotics. <i>ChemBioChem</i> , 2020, 21, 623-627.	1.3	12
5	Discovery of C-Glycosylpyranonaphthoquinones in <i>Streptomyces</i> sp. MBT76 by a Combined NMR-Based Metabolomics and Bioinformatics Workflow. <i>Journal of Natural Products</i> , 2017, 80, 269-277.	1.5	36
6	Bifunctionality of ActIV as a Cyclase-Thioesterase Revealed by in Vitro Reconstitution of Actinorhodin Biosynthesis in <i>Streptomyces coelicolor</i> A3(2). <i>ChemBioChem</i> , 2017, 18, 316-323.	1.3	30
7	Structure and biosynthetic implication of 5R-(N-acetyl-L-cysteinyl)-14S-hydroxy-dihydrokalafungin from a mutant of the actVA-ORF4 gene for actinorhodin biosynthesis in <i>Streptomyces coelicolor</i> A3(2). <i>Journal of Antibiotics</i> , 2015, 68, 481-483.	1.0	6
8	Distinct intraspecific variations of garlic (<i>Allium sativum</i> L.) revealed by the exon-intron sequences of the alliinase gene. <i>Journal of Natural Medicines</i> , 2014, 68, 442-447.	1.1	5
9	Natural Product Proteomining, a Quantitative Proteomics Platform, Allows Rapid Discovery of Biosynthetic Gene Clusters for Different Classes of Natural Products. <i>Chemistry and Biology</i> , 2014, 21, 707-718.	6.2	51
10	Biosynthesis of pyranonaphthoquinone polyketides reveals diverse strategies for enzymatic carbon-carbon bond formation. <i>Current Opinion in Chemical Biology</i> , 2013, 17, 562-570.	2.8	29
11	Biosynthetic Conclusions from the Functional Dissection of Oxygenases for Biosynthesis of Actinorhodin and Related <i>Streptomyces</i> Antibiotics. <i>Chemistry and Biology</i> , 2013, 20, 510-520.	6.2	45
12	Identification of the actinorhodin monomer and its related compound from a deletion mutant of the actVA-ORF4 gene of <i>Streptomyces coelicolor</i> A3(2). <i>Bioorganic and Medicinal Chemistry Letters</i> , 2012, 22, 5041-5045.	1.0	39
13	Epoxyquinone Formation Catalyzed by a Two-Component Flavin-Dependent Monooxygenase Involved in Biosynthesis of the Antibiotic Actinorhodin. <i>ChemBioChem</i> , 2011, 12, 2767-2773.	1.3	17
14	Practical procedures for genetic manipulation systems for medermycin-producing <i>Streptomyces</i> sp. AM7161. <i>Journal of Basic Microbiology</i> , 2010, 50, 299-301.	1.8	3
15	Biosynthesis of Actinorhodin and Related Antibiotics: Discovery of Alternative Routes for Quinone Formation Encoded in the act Gene Cluster. <i>Chemistry and Biology</i> , 2009, 16, 226-236.	6.2	88
16	Actinoperlyone, a novel perylenequinone-type shunt product, from a deletion mutant of the actVA-ORF5 and ORF6 genes for actinorhodin biosynthesis in <i>Streptomyces coelicolor</i> A3(2). <i>Tetrahedron Letters</i> , 2008, 49, 1208-1211.	0.7	11
17	Actinorhodin Biosynthesis: Structural Requirements for Post-PKS Tailoring Intermediates Revealed by Functional Analysis of ActVI-ORF1 Reductase. <i>Biochemistry</i> , 2007, 46, 8181-8188.	1.2	28
18	Possible involvement of ActVI-ORFA in transcriptional regulation of actVI tailoring-step genes for actinorhodin biosynthesis. <i>FEMS Microbiology Letters</i> , 2007, 269, 234-239.	0.7	12

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19	Functional studies on a ketoreductase gene from <i>Streptomyces</i> sp. AM-7161 to control the stereochemistry in medermycin biosynthesis. <i>Bioorganic and Medicinal Chemistry</i> , 2005, 13, 6856-6863.	1.4	15
20	The Glycosyltransferase UrdGT2 Catalyzes Both C- and O-Glycosidic Sugar Transfers. <i>Angewandte Chemie - International Edition</i> , 2004, 43, 2962-2965.	7.2	109
21	Remarkably different structures and reaction mechanisms of ketoreductases for the opposite stereochemical control in the biosynthesis of BIQ antibiotics. <i>Bioorganic and Medicinal Chemistry</i> , 2004, 12, 5917-5927.	1.4	18
22	Cloning, sequencing and heterologous expression of the medermycin biosynthetic gene cluster of <i>Streptomyces</i> sp. AM-7161: towards comparative analysis of the benzoisochromanequinone gene clusters. <i>Microbiology (United Kingdom)</i> , 2003, 149, 1633-1645.	0.7	112
23	A New Mode of Stereochemical Control Revealed by Analysis of the Biosynthesis of Dihydrogranaticin in <i>Streptomyces violaceoruber</i> TÅ¼422. <i>Journal of the American Chemical Society</i> , 2001, 123, 11376-11380.	6.6	31
24	Functional Complementation of Pyran Ring Formation in Actinorhodin Biosynthesis in <i>Streptomyces coelicolor</i> A3(2) by Ketoreductase Genes for Granaticin Biosynthesis. <i>Journal of Bacteriology</i> , 2001, 183, 3247-3250.	1.0	26
25	Chemical Characterisation of Disruptants of the <i>Streptomyces coelicolor</i> A3(2). ActVI Genes Involved in Actinorhodin Biosynthesis.. <i>Journal of Antibiotics</i> , 2000, 53, 144-152.	1.0	61
26	Proof that the actVI genetic region of <i>Streptomyces coelicolor</i> A3(2) is involved in stereospecific pyran ring formation in the biosynthesis of actinorhodin. <i>Bioorganic and Medicinal Chemistry Letters</i> , 1999, 9, 395-400.	1.0	47
27	The granaticin biosynthetic gene cluster of <i>Streptomyces violaceoruber</i> TÅ¼422: sequence analysis and expression in a heterologous host. <i>Chemistry and Biology</i> , 1998, 5, 647-659.	6.2	141