Bin Hong

List of Publications by Year in descending order

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414414 394421 1,260 62 19 32 h-index citations g-index papers 65 65 65 1677 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	MicroRNAs 185, 96, and 223 Repress Selective High-Density Lipoprotein Cholesterol Uptake through Posttranscriptional Inhibition. Molecular and Cellular Biology, 2013, 33, 1956-1964.	2.3	144
2	Butyrate protects against highâ€fat dietâ€induced atherosclerosis via upâ€regulating ABCA1 expression in apolipoprotein Eâ€deficiency mice. British Journal of Pharmacology, 2020, 177, 1754-1772.	5.4	96
3	Berberine attenuates choline-induced atherosclerosis by inhibiting trimethylamine and trimethylamine-N-oxide production via manipulating the gut microbiome. Npj Biofilms and Microbiomes, 2021, 7, 36.	6.4	81
4	Salvianolic acid B inhibits macrophage uptake of modified low density lipoprotein (mLDL) in a scavenger receptor CD36-dependent manner. Atherosclerosis, 2012, 223, 152-159.	0.8	72
5	microRNA-185 modulates low density lipoprotein receptor expression as a key posttranscriptional regulator. Atherosclerosis, 2015, 243, 523-532.	0.8	60
6	Rutaecarpine suppresses atherosclerosis in ApoE \hat{a} '/ \hat{a} ' mice through upregulating ABCA1 and SR-BI within RCT. Journal of Lipid Research, 2014, 55, 1634-1647.	4.2	53
7	Discovery of Antagonists for Human Scavenger Receptor CD36 via an ELISA-Like High-Throughput Screening Assay. Journal of Biomolecular Screening, 2010, 15, 239-250.	2.6	47
8	SsaA, a Member of a Novel Class of Transcriptional Regulators, Controls Sansanmycin Production in Streptomyces sp. Strain SS through a Feedback Mechanism. Journal of Bacteriology, 2013, 195, 2232-2243.	2.2	36
9	A small-molecule inhibitor of PCSK9 transcription ameliorates atherosclerosis through the modulation of FoxO1/3 and HNF1 $\hat{l}\pm$. EBioMedicine, 2020, 52, 102650.	6.1	36
10	Identification of Upregulators of Human ATP-Binding Cassette Transporter A1 via High-Throughput Screening of a Synthetic and Natural Compound Library. Journal of Biomolecular Screening, 2008, 13, 648-656.	2.6	35
11	Streptomycin production by Streptomyces griseus can be modulated by a mechanism not associated with change in the adpA component of the A-factor cascade. Biotechnology Letters, 2006, 29, 57-64.	2.2	31
12	Identification of trichostatin A as a novel transcriptional up-regulator of scavenger receptor BI both in HepG2 and RAW 264.7 cells. Atherosclerosis, 2009, 204, 127-135.	0.8	28
13	Binding of a biosynthetic intermediate to <scp>AtrA</scp> modulates the production of lidamycin by <scp><i>S</i></scp> <i>treptomyces globisporus</i>	2.5	28
14	Identification of novel mureidomycin analogues via rational activation of a cryptic gene cluster in Streptomyces roseosporus NRRL 15998. Scientific Reports, 2015, 5, 14111.	3.3	27
15	Identification of an anti-Gram-negative bacteria agent disrupting the interaction between lipopolysaccharide transporters LptA and LptC. International Journal of Antimicrobial Agents, 2019, 53, 442-448.	2.5	27
16	NRPS Substrate Promiscuity Leads to More Potent Antitubercular Sansanmycin Analogues. Journal of Natural Products, 2014, 77, 1744-1748.	3.0	26
17	Production of C-Terminal Amidated Recombinant Salmon Calcitonin in Streptomyces lividans. Applied Biochemistry and Biotechnology, 2003, 110, 113-123.	2.9	24
18	Biosynthesis of antibiotic chuangxinmycin from Actinoplanes tsinanensis. Acta Pharmaceutica Sinica B, 2018, 8, 283-294.	12.0	22

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19	Entecavir therapy reverses gut microbiota dysbiosis induced by hepatitis B virus infection in a mouse model. International Journal of Antimicrobial Agents, 2020, 56, 106000.	2.5	21
20	The Cytochrome P450 Catalyzing Câ^'S Bond Formation in <i>S</i> â€Heterocyclization of Chuangxinmycin Biosynthesis. Angewandte Chemie - International Edition, 2021, 60, 15399-15404.	13.8	20
21	Complete genome sequence of Streptomyces globisporus C-1027, the producer of an enediyne antibiotic lidamycin. Journal of Biotechnology, 2016, 222, 9-10.	3.8	19
22	Improving the N-terminal diversity of sansanmycin through mutasynthesis. Microbial Cell Factories, 2016, 15, 77.	4.0	17
23	Role of sgcR3 in positive regulation of enediyne antibiotic C-1027 production of Streptomyces globisporus C-1027. BMC Microbiology, 2009, 9, 14.	3.3	16
24	Identification of two antagonists of the scavenger receptor CD36 using a high-throughput screening model. Analytical Biochemistry, 2010, 400, 207-212.	2.4	15
25	Precursor-directed biosynthesis of new sansanmycin analogs bearing para-substituted-phenylalanines with high yields. Journal of Antibiotics, 2016, 69, 765-768.	2.0	14
26	Exploring novel herbicidin analogues by transcriptional regulator overexpression and MS/MS molecular networking. Microbial Cell Factories, 2019, 18, 175.	4.0	14
27	Draft Genome Sequence of Streptomyces globisporus C-1027, Which Produces an Antitumor Antibiotic Consisting of a Nine-Membered Enediyne with a Chromoprotein. Journal of Bacteriology, 2012, 194, 4144-4144.	2.2	13
28	Identification of trichostatin derivatives from Streptomyces sp. CPCC 203909. Bioorganic and Medicinal Chemistry Letters, 2015, 25, 562-565.	2.2	13
29	Structure–activity relationship and biological evaluation of berberine derivatives as PCSK9 down-regulating agents. Bioorganic Chemistry, 2021, 113, 104994.	4.1	13
30	Identifying Small-Molecule Inhibitors of SARS-CoV-2 RNA-Dependent RNA Polymerase by Establishing a Fluorometric Assay. Frontiers in Immunology, 2022, 13, 844749.	4.8	13
31	Mycophenolic acid induces ATP-binding cassette transporter A1 (ABCA1) expression through the PPARγ–LXRα–ABCA1 pathway. Biochemical and Biophysical Research Communications, 2011, 414, 779-782.	2.1	12
32	Application of untargeted tandem mass spectrometry with molecular networking for detection of enniatins and beauvericins from complex samples. Journal of Chromatography A, 2020, 1634, 461626.	3.7	12
33	Hangtaimycin, a peptide secondary metabolite discovered from Streptomyces spectabilis CPCC 200148 by chemical screening. Journal of Antibiotics, 2016, 69, 835-838.	2.0	11
34	Heterologous expression of human interleukin-6 in Streptomyces lividans TK24 using novel secretory expression vectors. Biotechnology Letters, 2011, 33, 253-261.	2.2	10
35	The human gut microbiome – a new and exciting avenue in cardiovascular drug discovery. Expert Opinion on Drug Discovery, 2019, 14, 1037-1052.	5.0	10
36	Disruption of <i>cagA </i> , the apoprotein gene of chromoprotein antibiotic C-1027, eliminates holo-antibiotic production, but not the cytotoxic chromophore. FEMS Microbiology Letters, 2009, 301, 57-68.	1.8	9

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37	Synthesis and structure–activity relationship of N-(2-arylethyl) isoquinoline derivatives as human scavenger receptor CD36 antagonists. European Journal of Medicinal Chemistry, 2011, 46, 1066-1073.	5.5	9
38	Metformin alleviates choline diet-induced TMAO elevation in C57BL/6J mice by influencing gut-microbiota composition and functionality. Nutrition and Diabetes, 2021, 11, 27.	3.2	9
39	MicroRNA-185 modulates CYP7A1 mediated cholesterol-bile acid metabolism through post-transcriptional and post-translational regulation of FoxO1. Atherosclerosis, 2022, 348, 56-67.	0.8	9
40	High-density lipoprotein-based drug discovery for treatment of atherosclerosis. Expert Opinion on Drug Discovery, 2015, 10, 841-855.	5.0	8
41	Low-density lipoprotein upregulate SR-BI through Sp1 Ser702 phosphorylation in hepatic cells. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2016, 1861, 1066-1075.	2.4	8
42	Structure-based manual screening and automatic networking for systematically exploring sansanmycin analogues using high performance liquid chromatography tandem mass spectroscopy. Journal of Pharmaceutical and Biomedical Analysis, 2018, 158, 94-105.	2.8	8
43	E17241 as a Novel ABCA1 (ATP-Binding Cassette Transporter A1) Upregulator Ameliorates Atherosclerosis in Mice. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, e284-e298.	2.4	8
44	A novel compound 4010B-30 upregulates apolipoprotein A-I gene expression through activation of PPAR \hat{I}^3 in HepG2 cells. Atherosclerosis, 2015, 239, 589-598.	0.8	7
45	Complete genome sequence of Amycolatopsis orientalis CPCC200066, the producer of norvancomycin. Journal of Biotechnology, 2017, 247, 6-10.	3.8	7
46	Substituted Benzamides Containing Azaspiro Rings as Upregulators of Apolipoprotein A-I Transcription. Molecules, 2012, 17, 7379-7386.	3.8	6
47	Three structurally-related impurities in norvancomycin drug substance. Journal of Antibiotics, 2017, 70, 158-165.	2.0	6
48	Comparative genomics and transcriptomics analyses provide insights into the high yield and regulatory mechanism of Norvancomycin biosynthesis in Amycolatopsis orientalis NCPC 2-48. Microbial Cell Factories, 2021, 20, 28.	4.0	6
49	Antisense microRNA185 loaded liposome for efficient inhibition of the hepatic endogenous microRNA185 level. European Journal of Pharmaceutical Sciences, 2021, 161, 105803.	4.0	6
50	How can high-throughput screening deliver drugs to treat atherosclerosis? Expert Opinion on Drug Discovery, 2010, 5, 1175-1188.	5.0	5
51	Draft Genome Sequence of Streptomyces sp. Strain SS, Which Produces a Series of Uridyl Peptide Antibiotic Sansanmycins. Journal of Bacteriology, 2012, 194, 6988-6989.	2.2	5
52	Advances in the role of microRNAs in lipid metabolism-related anti-atherosclerotic drug discovery. Expert Opinion on Drug Discovery, 2013, 8, 977-990.	5.0	5
53	Rescrutiny of the sansanmycin biosynthetic gene cluster leads to the discovery of a novel sansanmycin analogue with more potency against Mycobacterium tuberculosis. Journal of Antibiotics, 2019, 72, 769-774.	2.0	5
54	Cytochrome P450 Monooxygenase for Catalyzing C-42 Hydroxylation of the Glycine-Derived Fragment in Hangtaimycin Biosynthesis. Organic Letters, 2022, 24, 1388-1393.	4.6	5

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55	Draft Genome Sequence of Norvancomycin-Producing Strain Amycolatopsis orientalis CPCC200066. Genome Announcements, 2015, 3, .	0.8	4
56	Identification of Novel Compounds Enhancing SR-BI mRNA Stability through High-Throughput Screening. SLAS Discovery, 2020, 25, 397-408.	2.7	2
57	The Cytochrome P450 Catalyzing Câ°'S Bond Formation in S â€Heterocyclization of Chuangxinmycin Biosynthesis. Angewandte Chemie, 2021, 133, 15527-15532.	2.0	2
58	Mintaimycins, a Group of Novel Peptide Metabolites from Micromonospora sp. C-3509. Molecules, 2022, 27, 1150.	3.8	2
59	Multi-Omics-Guided Discovery of Omicsynins Produced by Streptomyces sp. 1647: Pseudo-Tetrapeptides Active Against Influenza A Viruses and Coronavirus HCoV-229E. Engineering, 2022, 16, 176-186.	6.7	1
60	Draft Genome Sequence of Teicoplanin-Producing Strain Actinoplanes teichomyceticus CPCC 203265. Microbiology Resource Announcements, 2019, 8, .	0.6	1
61	Treating chronic diseases by regulating the gut microbiota. Engineering, 2021, , .	6.7	1
62	Draft Genome Sequence of Streptomyces sp. Strain 105A-00742, Isolated in Shangri-La, China. Microbiology Resource Announcements, 2020, 9, .	0.6	0