

Changhao Bi

List of Publications by Year in descending order

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Version: 2024-02-01

52
papers

1,931
citations

257101

24
h-index

276539

41
g-index

61
all docs

61
docs citations

61
times ranked

2146
citing authors

#	ARTICLE	IF	CITATIONS
1	Imperfect guide-RNA (igRNA) enables CRISPR single-base editing with ABE and CBE. <i>Nucleic Acids Research</i> , 2022, 50, 4161-4170.	6.5	13
2	Reconstructed glycosylase base editors GBE2.0 with enhanced C-to-G base editing efficiency and purity. <i>Molecular Therapy</i> , 2022, 30, 2452-2463.	3.7	17
3	Engineering Circularized mRNAs for the Production of Spider Silk Proteins. <i>Applied and Environmental Microbiology</i> , 2022, 88, e0002822.	1.4	6
4	Glycosylase base editors enable C-to-A and C-to-G base changes. <i>Nature Biotechnology</i> , 2021, 39, 35-40.	9.4	277
5	Engineering an efficient H ₂ utilizing <i>Escherichia coli</i> platform by modulation of endogenous hydrogenases. <i>Biochemical Engineering Journal</i> , 2021, 166, 107851.	1.8	0
6	CRISPR-based metabolic pathway engineering. <i>Metabolic Engineering</i> , 2021, 63, 148-159.	3.6	24
7	Characterization of JEN family carboxylate transporters from the acid-tolerant yeast <i>Pichia kudriavzevii</i> and their applications in succinic acid production. <i>Microbial Biotechnology</i> , 2021, 14, 1130-1147.	2.0	23
8	Helicase-AID: A novel molecular device for base editing at random genomic loci. <i>Metabolic Engineering</i> , 2021, 67, 396-402.	3.6	6
9	Multiple strategies for metabolic engineering of <i>Escherichia coli</i> for efficient production of glycolate. <i>Biotechnology and Bioengineering</i> , 2021, 118, 4699-4707.	1.7	12
10	Molecular Mechanism of the Cytosine CRISPR Base Editing Process and the Roles of Translesion DNA Polymerases. <i>ACS Synthetic Biology</i> , 2021, 10, 3353-3358.	1.9	10
11	Identification of <i>Absidia orchidis</i> steroid 11 β -hydroxylation system and its application in engineering <i>Saccharomyces cerevisiae</i> for one-step biotransformation to produce hydrocortisone. <i>Metabolic Engineering</i> , 2020, 57, 31-42.	3.6	42
12	Constructing a Novel Biosynthetic Pathway for the Production of Glycolate from Glycerol in <i>Escherichia coli</i> . <i>ACS Synthetic Biology</i> , 2020, 9, 2600-2609.	1.9	19
13	Manipulating the position of DNA expression cassettes using location tags fused to dCas9 (Cas9-Lag) to improve metabolic pathway efficiency. <i>Microbial Cell Factories</i> , 2020, 19, 229.	1.9	5
14	Engineering the Calvin-Benson-Bassham cycle and hydrogen utilization pathway of <i>Ralstonia eutropha</i> for improved autotrophic growth and polyhydroxybutyrate production. <i>Microbial Cell Factories</i> , 2020, 19, 228.	1.9	39
15	Coordinated Expression of Astaxanthin Biosynthesis Genes for Improved Astaxanthin Production in <i>Escherichia coli</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2020, 68, 14917-14927.	2.4	38
16	A novel gene expression system for <i>Ralstonia eutropha</i> based on the T7 promoter. <i>BMC Microbiology</i> , 2020, 20, 121.	1.3	6
17	Nonclassical Biofilms Induced by DNA Breaks in <i>Klebsiella pneumoniae</i> . <i>MSphere</i> , 2020, 5, .	1.3	6
18	CRISPR-dCas9 Mediated Cytosine Deaminase Base Editing in <i>Bacillus subtilis</i> . <i>ACS Synthetic Biology</i> , 2020, 9, 1781-1789.	1.9	38

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19	Construction of a carbon-conserving pathway for glycolate production by synergetic utilization of acetate and glucose in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2020, 61, 152-159.	3.6	19
20	A programmable CRISPR/Cas9-based phage defense system for <i>Escherichia coli</i> BL21(DE3). <i>Microbial Cell Factories</i> , 2020, 19, 136.	1.9	9
21	Production of 14 β -hydroxysteroids by a recombinant <i>Saccharomyces cerevisiae</i> biocatalyst expressing of a fungal steroid 14 β -hydroxylation system. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 8363-8374.	1.7	22
22	Construction of <i>Escherichia coli</i> cell factories for crocin biosynthesis. <i>Microbial Cell Factories</i> , 2019, 18, 120.	1.9	39
23	Combinatorial modulation of initial codons for improved zeaxanthin synthetic pathway efficiency in <i>Escherichia coli</i> . <i>MicrobiologyOpen</i> , 2019, 8, e930.	1.2	11
24	CRISPR-Cas9-assisted native end-joining editing offers a simple strategy for efficient genetic engineering in <i>Escherichia coli</i> . <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 8497-8509.	1.7	25
25	Engineering an electroactive <i>Escherichia coli</i> for the microbial electrosynthesis of succinate from glucose and CO ₂ . <i>Microbial Cell Factories</i> , 2019, 18, 15.	1.9	66
26	Engineering an Artificial Membrane Vesicle Trafficking System (AMVTS) for the Excretion of β -Carotene in <i>Escherichia coli</i> . <i>ACS Synthetic Biology</i> , 2019, 8, 1037-1046.	1.9	36
27	Engineering an electroactive <i>Escherichia coli</i> for the microbial electrosynthesis of succinate by increasing the intracellular FAD pool. <i>Biochemical Engineering Journal</i> , 2019, 146, 132-142.	1.8	12
28	Development of an autotrophic fermentation technique for the production of fatty acids using an engineered <i>Ralstonia eutropha</i> cell factory. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2019, 46, 783-790.	1.4	17
29	Double-Check Base Editing for Efficient A to G Conversions. <i>ACS Synthetic Biology</i> , 2019, 8, 2629-2634.	1.9	14
30	Optimizing the localization of astaxanthin enzymes for improved productivity. <i>Biotechnology for Biofuels</i> , 2018, 11, 278.	6.2	49
31	CRISPR/Cas9 Assisted Multiplex Genome Editing Technique in <i>Escherichia coli</i> . <i>Biotechnology Journal</i> , 2018, 13, e1700604.	1.8	44
32	Engineering membrane morphology and manipulating synthesis for increased lycopene accumulation in <i>Escherichia coli</i> cell factories. <i>3 Biotech</i> , 2018, 8, 269.	1.1	34
33	Genome editing of <i>Ralstonia eutropha</i> using an electroporation-based CRISPR-Cas9 technique. <i>Biotechnology for Biofuels</i> , 2018, 11, 172.	6.2	50
34	Engineering <i>Saccharomyces cerevisiae</i> for the production of the valuable monoterpene ester geranyl acetate. <i>Microbial Cell Factories</i> , 2018, 17, 85.	1.9	25
35	Type II _s restriction based combinatory modulation technique for metabolic pathway optimization. <i>Microbial Cell Factories</i> , 2017, 16, 47.	1.9	7
36	Balanced activation of IspG and IspH to eliminate MEP intermediate accumulation and improve isoprenoids production in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2017, 44, 13-21.	3.6	51

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37	Improving Succinate Productivity by Engineering a Cyanobacterial CO ₂ Concentrating System (CCM) in <i>Escherichia coli</i> . <i>Biotechnology Journal</i> , 2017, 12, 1700199.	1.8	8
38	The CRISPR/Cas9-facilitated multiplex pathway optimization (CFPO) technique and its application to improve the <i>Escherichia coli</i> xylose utilization pathway. <i>Metabolic Engineering</i> , 2017, 43, 37-45.	3.6	57
39	CRISPR/Cas9-assisted gRNA-free one-step genome editing with no sequence limitations and improved targeting efficiency. <i>Scientific Reports</i> , 2017, 7, 16624.	1.6	29
40	Membrane engineering - A novel strategy to enhance the production and accumulation of β -carotene in <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2017, 43, 85-91.	3.6	89
41	Construction of a novel anaerobic pathway in <i>Escherichia coli</i> for propionate production. <i>BMC Biotechnology</i> , 2017, 17, 38.	1.7	21
42	A novel point mutation in RpoB improves osmotolerance and succinic acid production in <i>Escherichia coli</i> . <i>BMC Biotechnology</i> , 2017, 17, 10.	1.7	28
43	Development of a fast and easy method for <i>Escherichia coli</i> genome editing with CRISPR/Cas9. <i>Microbial Cell Factories</i> , 2016, 15, 205.	1.9	96
44	Combinatory optimization of chromosomal integrated mevalonate pathway for β -carotene production in <i>Escherichia coli</i> . <i>Microbial Cell Factories</i> , 2016, 15, 202.	1.9	29
45	Engineering <i>Corynebacterium glutamicum</i> for violacein hyper production. <i>Microbial Cell Factories</i> , 2016, 15, 148.	1.9	46
46	Development of a modularized two-step (M2S) chromosome integration technique for integration of multiple transcription units in <i>Saccharomyces cerevisiae</i> . <i>Biotechnology for Biofuels</i> , 2016, 9, 232.	6.2	22
47	End-to-end automated microfluidic platform for synthetic biology: from design to functional analysis. <i>Journal of Biological Engineering</i> , 2016, 10, 3.	2.0	54
48	PR-PR: Cross-Platform Laboratory Automation System. <i>ACS Synthetic Biology</i> , 2014, 3, 515-524.	1.9	41
49	Development of a broad-host synthetic biology toolbox for <i>Ralstonia eutropha</i> and its application to engineering hydrocarbon biofuel production. <i>Microbial Cell Factories</i> , 2013, 12, 107.	1.9	103
50	PaR-PaR Laboratory Automation Platform. <i>ACS Synthetic Biology</i> , 2013, 2, 216-222.	1.9	46
51	Engineering of <i>Ralstonia eutropha</i> H16 for Autotrophic and Heterotrophic Production of Methyl Ketones. <i>Applied and Environmental Microbiology</i> , 2013, 79, 4433-4439.	1.4	139
52	Cytotoxicity of HIV-gp41 segments expressed in <i>E. coli</i> . <i>Science Bulletin</i> , 2004, 49, 668-671.	1.7	1