

Liming Liu

List of Publications by Year in descending order

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205
papers

5,960
citations

71102

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all docs

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docs citations

225
times ranked

4956
citing authors

#	ARTICLE	IF	CITATIONS
1	The RAVEN Toolbox and Its Use for Generating a Genome-scale Metabolic Model for <i>Penicillium chrysogenum</i> . <i>PLoS Computational Biology</i> , 2013, 9, e1002980.	3.2	364
2	DCEO Biotechnology: Tools To Design, Construct, Evaluate, and Optimize the Metabolic Pathway for Biosynthesis of Chemicals. <i>Chemical Reviews</i> , 2018, 118, 4-72.	47.7	141
3	Engineering redox balance through cofactor systems. <i>Trends in Biotechnology</i> , 2014, 32, 337-343.	9.3	138
4	Reconstruction and analysis of a genome-scale metabolic model of the oleaginous fungus <i>Mortierella alpina</i> . <i>BMC Systems Biology</i> , 2015, 9, 1.	3.0	131
5	Genome-wide association study identifies new susceptibility loci for adolescent idiopathic scoliosis in Chinese girls. <i>Nature Communications</i> , 2015, 6, 8355.	12.8	104
6	ATP in current biotechnology: Regulation, applications and perspectives. <i>Biotechnology Advances</i> , 2009, 27, 94-101.	11.7	103
7	Engineering microbial membranes to increase stress tolerance of industrial strains. <i>Metabolic Engineering</i> , 2019, 53, 24-34.	7.0	94
8	Engineering Microorganisms for Enhanced CO ₂ Sequestration. <i>Trends in Biotechnology</i> , 2019, 37, 532-547.	9.3	86
9	Programmable biomolecular switches for rewiring flux in <i>Escherichia coli</i> . <i>Nature Communications</i> , 2019, 10, 3751.	12.8	84
10	Metabolic engineering of <i>Torulopsis glabrata</i> for malate production. <i>Metabolic Engineering</i> , 2013, 19, 10-16.	7.0	83
11	Identification of a critical determinant that enables efficient fatty acid synthesis in oleaginous fungi. <i>Scientific Reports</i> , 2015, 5, 11247.	3.3	83
12	Use of genome-scale metabolic models for understanding microbial physiology. <i>FEBS Letters</i> , 2010, 584, 2556-2564.	2.8	81
13	Engineering <i>Escherichia coli</i> for malate production by integrating modular pathway characterization with CRISPRi-guided multiplexed metabolic tuning. <i>Biotechnology and Bioengineering</i> , 2018, 115, 661-672.	3.3	77
14	Light-driven CO ₂ sequestration in <i>Escherichia coli</i> to achieve theoretical yield of chemicals. <i>Nature Catalysis</i> , 2021, 4, 395-406.	34.4	75
15	Reconstruction of cytosolic fumaric acid biosynthetic pathways in <i>Saccharomyces cerevisiae</i> . <i>Microbial Cell Factories</i> , 2012, 11, 24.	4.0	68
16	Enhancement of pyruvate production by osmotic-tolerant mutant of <i>Torulopsis glabrata</i> . <i>Biotechnology and Bioengineering</i> , 2007, 97, 825-832.	3.3	67
17	Screening of a thiamine-auxotrophic yeast for L-glutamic acid overproduction. <i>Letters in Applied Microbiology</i> , 2010, 51, 264-271.	2.2	67
18	Metabolic engineering of <i>Escherichia coli</i> W3110 to produce L-malate. <i>Biotechnology and Bioengineering</i> , 2017, 114, 656-664.	3.3	67

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19	Pathway dissection, regulation, engineering and application: lessons learned from biobutanol production by solventogenic clostridia. <i>Biotechnology for Biofuels</i> , 2020, 13, 39.	6.2	65
20	Acetoin production enhanced by manipulating carbon flux in a newly isolated <i>Bacillus amyloliquefaciens</i> . <i>Bioresource Technology</i> , 2013, 130, 256-260.	9.6	64
21	Redistribution of carbon flux in <i>Torulopsis glabrata</i> by altering vitamin and calcium level. <i>Metabolic Engineering</i> , 2007, 9, 21-29.	7.0	63
22	Enhanced hyaluronic acid production of <i>Streptococcus zooepidemicus</i> by an intermittent alkaline-stress strategy. <i>Letters in Applied Microbiology</i> , 2008, 46, 383-388.	2.2	62
23	Engineering <i>Escherichia coli</i> lifespan for enhancing chemical production. <i>Nature Catalysis</i> , 2020, 3, 307-318.	34.4	61
24	Manipulation of <i>B. megaterium</i> growth for efficient 2-KLG production by <i>K. vulgare</i> . <i>Process Biochemistry</i> , 2010, 45, 602-606.	3.7	59
25	Development of chemically defined media supporting high cell density growth of <i>Ketogulonigenium vulgare</i> and <i>Bacillus megaterium</i> . <i>Bioresource Technology</i> , 2011, 102, 4807-4814.	9.6	58
26	Engineering rTCA pathway and C4-dicarboxylate transporter for l-malic acid production. <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 4041-4052.	3.6	57
27	Manipulating the pyruvate dehydrogenase bypass of a multi-vitamin auxotrophic yeast <i>Torulopsis glabrata</i> enhanced pyruvate production. <i>Letters in Applied Microbiology</i> , 2004, 39, 199-206.	2.2	56
28	Enhancement of pyruvate productivity in <i>Torulopsis glabrata</i> : Increase of NAD ⁺ availability. <i>Journal of Biotechnology</i> , 2006, 126, 173-185.	3.8	55
29	Engineering synergetic CO ₂ -fixing pathways for malate production. <i>Metabolic Engineering</i> , 2018, 47, 496-504.	7.0	55
30	Enhancement of L-ketoglutarate production in <i>Torulopsis glabrata</i> : Redistribution of carbon flux from pyruvate to L-ketoglutarate. <i>Biotechnology and Bioprocess Engineering</i> , 2009, 14, 134-139.	2.6	53
31	Fumaric acid production by <i>Torulopsis glabrata</i> : Engineering the urea cycle and the purine nucleotide cycle. <i>Biotechnology and Bioengineering</i> , 2015, 112, 156-167.	3.3	52
32	Improved ATP supply enhances acid tolerance of <i>Candida glabrata</i> during pyruvic acid production. <i>Journal of Applied Microbiology</i> , 2011, 110, 44-53.	3.1	51
33	Fumaric acid production in <i>Saccharomyces cerevisiae</i> by simultaneous use of oxidative and reductive routes. <i>Bioresource Technology</i> , 2013, 148, 91-96.	9.6	51
34	Modular optimization of multi-gene pathways for fumarate production. <i>Metabolic Engineering</i> , 2016, 33, 76-85.	7.0	51
35	Light-powered <i>Escherichia coli</i> cell division for chemical production. <i>Nature Communications</i> , 2020, 11, 2262.	12.8	51
36	Fumaric Acid Production in <i>Saccharomyces cerevisiae</i> by In Silico Aided Metabolic Engineering. <i>PLoS ONE</i> , 2012, 7, e52086.	2.5	51

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37	Enzymatic production of $\hat{\pm}$ -ketoglutaric acid from l-glutamic acid via l-glutamate oxidase. <i>Journal of Biotechnology</i> , 2014, 179, 56-62.	3.8	50
38	Reconstruction and analysis of the genome-scale metabolic model of schizochytrium limacinum SR21 for docosaehaenoic acid production. <i>BMC Genomics</i> , 2015, 16, 799.	2.8	50
39	Isolation and Characterization of Three Antihypertension Peptides from the Mycelia of <i>Ganoderma Lucidum</i> (Agaricomycetes). <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 8149-8159.	5.2	49
40	Improving lysine production through construction of an <i>Escherichia coli</i> enzyme-constrained model. <i>Biotechnology and Bioengineering</i> , 2020, 117, 3533-3544.	3.3	47
41	Complete Genome Sequence of the Industrial Strain <i>Bacillus megaterium</i> WSH-002. <i>Journal of Bacteriology</i> , 2011, 193, 6389-6390.	2.2	46
42	Med15B Regulates Acid Stress Response and Tolerance in <i>Candida glabrata</i> by Altering Membrane Lipid Composition. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	46
43	Asymmetric assembly of high-value $\hat{\pm}$ -functionalized organic acids using a biocatalytic chiral-group-resetting process. <i>Nature Communications</i> , 2018, 9, 3818.	12.8	46
44	Genetic Circuit-Assisted Smart Microbial Engineering. <i>Trends in Microbiology</i> , 2019, 27, 1011-1024.	7.7	45
45	Compartmentalizing metabolic pathway in <i>Candida glabrata</i> for acetoin production. <i>Metabolic Engineering</i> , 2015, 28, 1-7.	7.0	43
46	Gelatin enhances 2-keto-l-gulonic acid production based on <i>Ketogulonigenium vulgare</i> genome annotation. <i>Journal of Biotechnology</i> , 2011, 156, 182-187.	3.8	42
47	Pyruvate production in <i>Candida glabrata</i> : manipulation and optimization of physiological function. <i>Critical Reviews in Biotechnology</i> , 2016, 36, 1-10.	9.0	42
48	Enhancing fructosylated chondroitin production in <i>Escherichia coli</i> K4 by balancing the UDP-precursors. <i>Metabolic Engineering</i> , 2018, 47, 314-322.	7.0	42
49	Production of bioactive metabolites by submerged fermentation of the medicinal mushroom <i>Antrodia cinnamomea</i> : recent advances and future development. <i>Critical Reviews in Biotechnology</i> , 2019, 39, 541-554.	9.0	42
50	Reconstruction and analysis of the genome-scale metabolic network of <i>Candida glabrata</i> . <i>Molecular BioSystems</i> , 2013, 9, 205-216.	2.9	41
51	Production of $\hat{2}$ -Alanine from Fumaric Acid Using a Dual-Enzyme Cascade. <i>ChemCatChem</i> , 2018, 10, 4984-4991.	3.7	39
52	Transcriptional engineering of <i>Escherichia coli</i> K4 for fructosylated chondroitin production. <i>Biotechnology Progress</i> , 2013, 29, 1140-1149.	2.6	37
53	Enhancement of malate production through engineering of the periplasmic rTCA pathway in <i>Escherichia coli</i> . <i>Biotechnology and Bioengineering</i> , 2018, 115, 1571-1580.	3.3	37
54	Lowering induction temperature for enhanced production of polygalacturonate lyase in recombinant <i>Pichia pastoris</i> . <i>Process Biochemistry</i> , 2009, 44, 949-954.	3.7	36

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55	Complete Genome Sequence of the Industrial Strain <i>Ketogulonicigenium vulgare</i> WSH-001. <i>Journal of Bacteriology</i> , 2011, 193, 6108-6109.	2.2	36
56	Reconstruction and analysis of a genome-scale metabolic model of the vitamin C producing industrial strain <i>Ketogulonicigenium vulgare</i> WSH-001. <i>Journal of Biotechnology</i> , 2012, 161, 42-48.	3.8	36
57	Open Gate of <i>Corynebacterium glutamicum</i> Threonine Deaminase for Efficient Synthesis of Bulky α -Keto Acids. <i>ACS Catalysis</i> , 2020, 10, 9994-10004.	11.2	36
58	A reusable method for construction of non-marker large fragment deletion yeast auxotroph strains: A practice in <i>Torulopsis glabrata</i> . <i>Journal of Microbiological Methods</i> , 2009, 76, 70-74.	1.6	35
59	Genome-scale reconstruction and in silico analysis of <i>Aspergillus terreus</i> metabolism. <i>Molecular BioSystems</i> , 2013, 9, 1939.	2.9	35
60	Engineering of the Conformational Dynamics of Lipase To Increase Enantioselectivity. <i>ACS Catalysis</i> , 2017, 7, 7593-7599.	11.2	35
61	Metabolic model reconstruction and analysis of an artificial microbial ecosystem for vitamin C production. <i>Journal of Biotechnology</i> , 2014, 182-183, 61-67.	3.8	34
62	Rewiring carbon flux in <i>Escherichia coli</i> using a bifunctional molecular switch. <i>Metabolic Engineering</i> , 2020, 61, 47-57.	7.0	34
63	Enhancement of alkaline polygalacturonate lyase production in recombinant <i>Pichia pastoris</i> according to the ratio of methanol to cell concentration. <i>Bioresource Technology</i> , 2009, 100, 1343-1349.	9.6	33
64	Reconstruction and analysis of the genome-scale metabolic model of <i>Lactobacillus casei</i> LC2W. <i>Gene</i> , 2015, 554, 140-147.	2.2	33
65	A constraint-based model of <i>Scheffersomyces stipitis</i> for improved ethanol production. <i>Biotechnology for Biofuels</i> , 2012, 5, 72.	6.2	32
66	Metabolic engineering of carbohydrate metabolism systems in <i>Corynebacterium glutamicum</i> for improving the efficiency of L-lysine production from mixed sugar. <i>Microbial Cell Factories</i> , 2020, 19, 39.	4.0	32
67	Arginine: A novel compatible solute to protect <i>Candida glabrata</i> against hyperosmotic stress. <i>Process Biochemistry</i> , 2011, 46, 1230-1235.	3.7	31
68	Enhancement of pyruvate productivity by inducible expression of a FOF1-ATPase inhibitor INH1 in <i>Torulopsis glabrata</i> CCTCC M202019. <i>Journal of Biotechnology</i> , 2009, 144, 120-126.	3.8	30
69	Structure, mechanism and regulation of an artificial microbial ecosystem for vitamin C production. <i>Critical Reviews in Microbiology</i> , 2013, 39, 247-255.	6.1	30
70	Engineering of carbonylase activity reaction in <i>Candida glabrata</i> for acetoin production. <i>Metabolic Engineering</i> , 2014, 22, 32-39.	7.0	30
71	Crz1p Regulates pH Homeostasis in <i>Candida glabrata</i> by Altering Membrane Lipid Composition. <i>Applied and Environmental Microbiology</i> , 2016, 82, 6920-6929.	3.1	30
72	Metabolic engineering of glucose uptake systems in <i>Corynebacterium glutamicum</i> for improving the efficiency of L-lysine production. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2019, 46, 937-949.	3.0	30

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73	Mitochondrial engineering of the TCA cycle for fumarate production. <i>Metabolic Engineering</i> , 2015, 31, 62-73.	7.0	29
74	Enhancing l-malate production of <i>Aspergillus oryzae</i> FMME218-37 by improving inorganic nitrogen utilization. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 8739-8751.	3.6	29
75	Mitochondrial DNA Heteroplasmy in <i>Candida glabrata</i> after Mitochondrial Transformation. <i>Eukaryotic Cell</i> , 2010, 9, 806-814.	3.4	28
76	Reconstruction and analysis of a genome-scale metabolic network of <i>Corynebacterium glutamicum</i> S9114. <i>Gene</i> , 2016, 575, 615-622.	2.2	27
77	Morphology engineering of <i>Aspergillus oryzae</i> for l-malate production. <i>Biotechnology and Bioengineering</i> , 2019, 116, 2662-2673.	3.3	27
78	Engineering of membrane phospholipid component enhances salt stress tolerance in <i>Saccharomyces cerevisiae</i> . <i>Biotechnology and Bioengineering</i> , 2020, 117, 710-720.	3.3	27
79	Significant increase of glycolytic flux in <i>Torulopsis glabrata</i> by inhibition of oxidative phosphorylation. <i>FEMS Yeast Research</i> , 2006, 6, 1117-1129.	2.3	26
80	<i>CgMED3</i> Changes Membrane Sterol Composition To Help <i>Candida glabrata</i> Tolerate Low-pH Stress. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	26
81	Reconstruction and Analysis of a Genome-Scale Metabolic Model of <i>Ganoderma lucidum</i> for Improved Extracellular Polysaccharide Production. <i>Frontiers in Microbiology</i> , 2018, 9, 3076.	3.5	26
82	Engineering microbial cell morphology and membrane homeostasis toward industrial applications. <i>Current Opinion in Biotechnology</i> , 2020, 66, 18-26.	6.6	26
83	Relationship Between Morphology and Itaconic Acid Production by <i>Aspergillus terreus</i> . <i>Journal of Microbiology and Biotechnology</i> , 2014, 24, 168-176.	2.1	26
84	Reconstruction and analysis of the industrial strain <i>Bacillus megaterium</i> WSH002 genome-scale in silico metabolic model. <i>Journal of Biotechnology</i> , 2013, 164, 503-509.	3.8	25
85	Production, structure and morphology of exopolysaccharides yielded by submerged fermentation of <i>Antrodia cinnamomea</i> . <i>Carbohydrate Polymers</i> , 2019, 205, 271-278.	10.2	25
86	Enhancement of Sphingolipid Synthesis Improves Osmotic Tolerance of <i>Saccharomyces cerevisiae</i> . <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	25
87	Increasing glycolytic flux in <i>Torulopsis glabrata</i> by redirecting ATP production from oxidative phosphorylation to substrate-level phosphorylation. <i>Journal of Applied Microbiology</i> , 2006, 100, 1043-1053.	3.1	24
88	Transcription factors Asg1p and Hal9p regulate pH homeostasis in <i>Candida glabrata</i> . <i>Frontiers in Microbiology</i> , 2015, 6, 843.	3.5	24
89	Enzymatic production of l-citrulline by hydrolysis of the guanidinium group of l-arginine with recombinant arginine deiminase. <i>Journal of Biotechnology</i> , 2015, 208, 37-43.	3.8	24
90	Enhancement of acetoin production in <i>Candida glabrata</i> by in silico-aided metabolic engineering. <i>Microbial Cell Factories</i> , 2014, 13, 55.	4.0	23

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91	A multifunctional tag with the ability to benefit the expression, purification, thermostability and activity of recombinant proteins. <i>Journal of Biotechnology</i> , 2018, 283, 1-10.	3.8	23
92	Comprehensive understanding of <i>Saccharomyces cerevisiae</i> phenotypes with whole-cell model WM_S288C. <i>Biotechnology and Bioengineering</i> , 2020, 117, 1562-1574.	3.3	23
93	Redirection of the NADH oxidation pathway in <i>Torulopsis glabrata</i> leads to an enhanced pyruvate production. <i>Applied Microbiology and Biotechnology</i> , 2006, 72, 377-385.	3.6	22
94	Engineering protonation conformation of aspartate decarboxylase to relieve mechanism-based inactivation. <i>Biotechnology and Bioengineering</i> , 2020, 117, 1607-1614.	3.3	22
95	Proline enhances <i>Torulopsis glabrata</i> growth during hyperosmotic stress. <i>Biotechnology and Bioprocess Engineering</i> , 2010, 15, 285-292.	2.6	21
96	Rational modification of <i>Corynebacterium glutamicum</i> dihydrodipicolinate reductase to switch the nucleotide cofactor specificity for increasing lysine production. <i>Biotechnology and Bioengineering</i> , 2018, 115, 1764-1777.	3.3	21
97	Spatial modulation and cofactor engineering of key pathway enzymes for fumarate production in <i>Candida glabrata</i> . <i>Biotechnology and Bioengineering</i> , 2019, 116, 622-630.	3.3	21
98	Comparison of covalent immobilization of amylase on polystyrene pellets with pentaethylenehexamine and pentaethylene glycol spacers. <i>Bioresource Technology</i> , 2011, 102, 9374-9379.	9.6	20
99	Development of a minimal chemically defined medium for <i>Ketogulonicigenium vulgare</i> WSH001 based on its genome-scale metabolic model. <i>Journal of Biotechnology</i> , 2014, 169, 15-22.	3.8	20
100	Kick-starting evolution efficiency with an autonomous evolution mutation system. <i>Metabolic Engineering</i> , 2019, 54, 127-136.	7.0	20
101	Dynamic consolidated bioprocessing for direct production of xylonate and shikimate from xylan by <i>Escherichia coli</i> . <i>Metabolic Engineering</i> , 2020, 60, 128-137.	7.0	20
102	Enhancing biofuels production by engineering the actin cytoskeleton in <i>Saccharomyces cerevisiae</i> . <i>Nature Communications</i> , 2022, 13, 1886.	12.8	20
103	Redirecting Carbon Flux in <i>Torulopsis glabrata</i> from Pyruvate to \pm -Ketoglutaric Acid by Changing Metabolic Co-factors. <i>Biotechnology Letters</i> , 2006, 28, 95-98.	2.2	19
104	KfoE encodes a fructosyltransferase involved in capsular polysaccharide biosynthesis in <i>Escherichia coli</i> K4. <i>Biotechnology Letters</i> , 2014, 36, 1469-1477.	2.2	19
105	Gene Circuits for Dynamically Regulating Metabolism. <i>Trends in Biotechnology</i> , 2018, 36, 751-754.	9.3	19
106	Regulation of thiamine synthesis in <i>Saccharomyces cerevisiae</i> for improved pyruvate production. <i>Yeast</i> , 2012, 29, 209-217.	1.7	18
107	Synergistic function of four novel thermostable glycoside hydrolases from a long-term enriched thermophilic methanogenic digester. <i>Frontiers in Microbiology</i> , 2015, 6, 509.	3.5	18
108	Enhancement of alpha-ketoglutaric acid production from l-glutamic acid by high-cell-density cultivation. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2016, 126, 10-17.	1.8	18

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109	Development of an <i>Escherichia coli</i> -based biocatalytic system for the efficient synthesis of N-acetyl-D-neuraminic acid. <i>Metabolic Engineering</i> , 2018, 47, 374-382.	7.0	18
110	A multi-enzyme cascade for efficient production of d-p-hydroxyphenylglycine from l-tyrosine. <i>Bioresources and Bioprocessing</i> , 2021, 8, .	4.2	18
111	Enzymatic production of l-ornithine from l-arginine with recombinant thermophilic arginase. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2014, 110, 1-7.	1.8	17
112	Pathway engineering of <i>Escherichia coli</i> for α -ketoglutaric acid production. <i>Biotechnology and Bioengineering</i> , 2020, 117, 2791-2801.	3.3	17
113	Microbial engineering for the production of C ₂ –C ₆ organic acids. <i>Natural Product Reports</i> , 2021, 38, 1518-1546.	10.3	17
114	Enhancing tryptophan production by balancing precursors in <i>Escherichia coli</i> . <i>Biotechnology and Bioengineering</i> , 2022, 119, 983-993.	3.3	17
115	Introduction of heterogeneous NADH reoxidation pathways into <i>Torulopsis glabrata</i> significantly increases pyruvate production efficiency. <i>Korean Journal of Chemical Engineering</i> , 2011, 28, 1078-1084.	2.7	16
116	Urea enhances cell growth and pyruvate production in <i>Torulopsis glabrata</i> . <i>Biotechnology Progress</i> , 2014, 30, 19-27.	2.6	16
117	<i>Cg</i> Hog1-Mediated <i>Cg</i> Rds2 Phosphorylation Alters Glycerophospholipid Composition To Coordinate Osmotic Stress in <i>Candida glabrata</i> . <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	16
118	Synergistic Metabolism of Glucose and Formate Increases the Yield of Short-Chain Organic Acids in <i>Escherichia coli</i> . <i>ACS Synthetic Biology</i> , 2022, 11, 135-143.	3.8	16
119	Production of polyvinyl alcohol-degrading enzyme with <i>Janthinobacterium</i> sp. and its application in cotton fabric desizing. <i>Biotechnology Journal</i> , 2007, 2, 752-758.	3.5	15
120	Genome-scale metabolic modelling common cofactors metabolism in microorganisms. <i>Journal of Biotechnology</i> , 2017, 251, 1-13.	3.8	15
121	Biocatalytic derivatization of proteinogenic amino acids for fine chemicals. <i>Biotechnology Advances</i> , 2020, 40, 107496.	11.7	15
122	Microbial cell engineering to improve cellular synthetic capacity. <i>Biotechnology Advances</i> , 2020, 45, 107649.	11.7	15
123	Engineering the Cad pathway in <i>Escherichia coli</i> to produce glutarate from l-lysine. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 3587-3599.	3.6	15
124	Enzymatic production of agmatine by recombinant arginine decarboxylase. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2015, 121, 1-8.	1.8	14
125	Genome-scale biological models for industrial microbial systems. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 3439-3451.	3.6	14
126	Hacking an Algal Transcription Factor for Lipid Biosynthesis. <i>Trends in Plant Science</i> , 2018, 23, 181-184.	8.8	14

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127	High-Throughput Screening of a 2-Keto-L-Gulonic Acid-Producing <i>Gluconobacter oxydans</i> Strain Based on Related Dehydrogenases. <i>Frontiers in Bioengineering and Biotechnology</i> , 2019, 7, 385.	4.1	14
128	Microbial physiological engineering increases the efficiency of microbial cell factories. <i>Critical Reviews in Biotechnology</i> , 2021, 41, 339-354.	9.0	14
129	Metabolic Engineering of <i>Candida glabrata</i> for Diacetyl Production. <i>PLoS ONE</i> , 2014, 9, e89854.	2.5	13
130	Genome Sequencing of the Pyruvate-producing Strain <i>Candida glabrata</i> CCTCC M202019 and Genomic Comparison with Strain CBS138. <i>Scientific Reports</i> , 2016, 6, 34893.	3.3	13
131	Production of α -Ketoisocaproate and α -Keto β -Methylvalerate by Engineered α -Amino Acid Deaminase. <i>ChemCatChem</i> , 2019, 11, 2464-2472.	3.7	13
132	A biosynthesis pathway for 3-hydroxypropionic acid production in genetically engineered <i>Saccharomyces cerevisiae</i> . <i>Green Chemistry</i> , 2021, 23, 4502-4509.	9.0	13
133	Reprogramming microbial populations using a programmed lysis system to improve chemical production. <i>Nature Communications</i> , 2021, 12, 6886.	12.8	13
134	Water-forming NADH oxidase protects <i>Torulopsis glabrata</i> against hyperosmotic stress. <i>Yeast</i> , 2010, 27, 207-216.	1.7	12
135	Glutathione enhances 2-keto-l-gulonic acid production based on <i>Ketogulonigenium vulgare</i> model iWZ663. <i>Journal of Biotechnology</i> , 2013, 164, 454-460.	3.8	12
136	Engineering the transmission efficiency of the noncyclic glyoxylate pathway for fumarate production in <i>Escherichia coli</i> . <i>Biotechnology for Biofuels</i> , 2020, 13, 132.	6.2	12
137	A novel high-yield process of phospholipase D-mediated phosphatidylserine production with cyclopentyl methyl ether. <i>Process Biochemistry</i> , 2018, 66, 146-149.	3.7	11
138	Reconstruction and in silico analysis of an <i>Actinoplanes</i> sp. SE50/110 genome-scale metabolic model for acarbose production. <i>Frontiers in Microbiology</i> , 2015, 6, 632.	3.5	10
139	Recycling of cooking oil fume condensate for the production of rhamnolipids by <i>Pseudomonas aeruginosa</i> WB505. <i>Bioprocess and Biosystems Engineering</i> , 2019, 42, 777-784.	3.4	10
140	Lsm12 Mediates Deubiquitination of DNA Polymerase β To Help <i>Saccharomyces cerevisiae</i> Resist Oxidative Stress. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	3.1	10
141	One-Pot Enzymatic "Chemical Cascade Route for Synthesizing Aromatic β -Hydroxy Ketones. <i>ACS Catalysis</i> , 2021, 11, 2808-2818.	11.2	10
142	Enhancing L-malate production of <i>Aspergillus oryzae</i> by nitrogen regulation strategy. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 3101-3113.	3.6	10
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