Ting Lu

List of Publications by Year in descending order

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257450 223800 2,408 47 24 46 citations h-index g-index papers 49 49 49 2658 all docs docs citations times ranked citing authors

#	Article	IF	Citations
1	Synthetic Biology Moving into the Clinic. Science, 2011, 333, 1248-1252.	12.6	348
2	Designing microbial consortia with defined social interactions. Nature Chemical Biology, 2018, 14, 821-829.	8.0	250
3	Stochastic Turing patterns in a synthetic bacterial population. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 6572-6577.	7.1	154
4	Markerless chromosomal gene deletion in Clostridium beijerinckii using CRISPR/Cas9 system. Journal of Biotechnology, 2015, 200, 1-5.	3.8	153
5	Bacterial Genome Editing with CRISPR-Cas9: Deletion, Integration, Single Nucleotide Modification, and Desirable "Clean―Mutant Selection in <i>Clostridium beijerinckii</i> as an Example. ACS Synthetic Biology, 2016, 5, 721-732.	3.8	143
6	Nonequilibrium physics in biology. Reviews of Modern Physics, 2019, 91, .	45.6	123
7	Autonomous production of 1,4-butanediol via a de novo biosynthesis pathway in engineered Escherichia coli. Metabolic Engineering, 2015, 29, 135-141.	7.0	109
8	Automatic Compilation from High-Level Biologically-Oriented Programming Language to Genetic Regulatory Networks. PLoS ONE, 2011, 6, e22490.	2.5	87
9	An integrative circuit–host modelling framework for predicting synthetic gene network behaviours. Nature Microbiology, 2017, 2, 1658-1666.	13.3	84
10	Bacterial social interactions drive the emergence of differential spatial colony structures. BMC Systems Biology, 2015, 9, 59.	3.0	62
11	Integrated, systems metabolic picture of acetone-butanol-ethanol fermentation by <i>Clostridium acetobutylicum</i> . Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8505-8510.	7.1	61
12	Composition and Metabolic Functions of the Microbiome in Fermented Grain during Light-Flavor Baijiu Fermentation. Microorganisms, 2020, 8, 1281.	3.6	52
13	Engineering robust and tunable spatial structures with synthetic gene circuits. Nucleic Acids Research, 2017, 45, 1005-1014.	14.5	48
14	Gene transcription repression in <i>Clostridium beijerinckii</i> using CRISPRâ€dCas9. Biotechnology and Bioengineering, 2016, 113, 2739-2743.	3.3	46
15	Cloning and Optimization of a Nisin Biosynthesis Pathway for Bacteriocin Harvest. ACS Synthetic Biology, 2014, 3, 439-445.	3.8	41
16	Identification and characterization of core sludge and biofilm microbiota in anaerobic membrane bioreactors. Environment International, 2019, 133, 105165.	10.0	40
17	Phenotypic variability of growing cellular populations. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 18982-18987.	7.1	39
18	Bacterial Consortium-Based Sensing System for Detecting Organophosphorus Pesticides. Analytical Chemistry, 2018, 90, 10577-10584.	6.5	39

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19	Interaction variability shapes succession of synthetic microbial ecosystems. Nature Communications, 2020, 11, 309.	12.8	33
20	Spatial interference scale as a determinant of microbial range expansion. Science Advances, 2018, 4, eaau0695.	10.3	32
21	Characterization of a <i>Clostridium beijerinckii spo0A</i> mutant and its application for butyl butyrate production. Biotechnology and Bioengineering, 2017, 114, 106-112.	3.3	31
22	Statistics of cellular signal transduction as a race to the nucleus by multiple random walkers in compartment/phosphorylation space. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 16752-16757.	7.1	30
23	Developing a Synthetic Biology Toolkit for <i>Comamonas testosteroni</i> , an Emerging Cellular Chassis for Bioremediation. ACS Synthetic Biology, 2018, 7, 1753-1762.	3.8	30
24	A molecular noise generator. Physical Biology, 2008, 5, 036006.	1.8	26
25	Population-Dynamic Modeling of Bacterial Horizontal Gene Transfer by Natural Transformation. Biophysical Journal, 2016, 110, 258-268.	0.5	24
26	A gene network engineering platform for lactic acid bacteria. Nucleic Acids Research, 2016, 44, e37-e37.	14.5	24
27	Engineered genetic information processing circuits. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2013, 5, 273-287.	6.6	23
28	Integrative Circuit-Host Modeling of a Genetic Switch in Varying Environments. Scientific Reports, 2020, 10, 8383.	3.3	23
29	Programming the group behaviors of bacterial communities with synthetic cellular communication. Bioresources and Bioprocessing, 2014, 1 , .	4.2	22
30	An Ecological Understanding of Quorum Sensing-Controlled Bacteriocin Synthesis. Cellular and Molecular Bioengineering, 2016, 9, 443-454.	2.1	22
31	Engineering microbial consortia with rationally designed cellular interactions. Current Opinion in Biotechnology, 2022, 76, 102730.	6.6	22
32	Slow and Steady Wins the Race: A Bacterial Exploitative Competition Strategy in Fluctuating Environments. ACS Synthetic Biology, 2015, 4, 240-248.	3.8	19
33	Circuit-Host Coupling Induces Multifaceted Behavioral Modulations of a Gene Switch. Biophysical Journal, 2018, 114, 737-746.	0.5	18
34	Extinction, coexistence, and localized patterns of a bacterial population with contact-dependent inhibition. BMC Systems Biology, 2014, 8, 23.	3.0	17
35	Development of an oxygen-independent flavin mononucleotide-based fluorescent reporter system in Clostridium beijerinckii and its potential applications. Journal of Biotechnology, 2018, 265, 119-126.	3.8	16
36	A Minimal Transcriptional Controlling Network of Regulatory T Cell Development. Journal of Physical Chemistry B, 2013, 117, 12995-13004.	2.6	15

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37	Genomic, Transcriptional, and Phenotypic Analysis of the Glucose Derepressed <i>Clostridium beijerinckii</i> Mutant Exhibiting Acid Crash Phenotype. Biotechnology Journal, 2017, 12, 1700182.	3.5	14
38	Engineering the bacterium Comamonas testosteroni CNB-1: Plasmid curing and genetic manipulation. Biochemical Engineering Journal, 2018, 133, 74-82.	3.6	13
39	Bacterial Genome Editing with CRISPR-Cas9: Taking Clostridium beijerinckii as an Example. Methods in Molecular Biology, 2018, 1772, 297-325.	0.9	13
40	Synthetic, Context-Dependent Microbial Consortium of Predator and Prey. ACS Synthetic Biology, 2019, 8, 1713-1722.	3.8	13
41	Efficacy of nisin derivatives with improved biochemical characteristics, alone and in combination with endolysin PlyP100 to control Listeria monocytogenes in laboratory-scale Queso Fresco. Food Microbiology, 2021, 94, 103668.	4.2	12
42	Harnessing lactic acid bacteria in synthetic microbial consortia. Trends in Biotechnology, 2022, 40, 8-11.	9.3	11
43	System-level modeling of acetone–butanol–ethanol fermentation. FEMS Microbiology Letters, 2016, 363, fnw074.	1.8	10
44	A comparative phenotypic and genomic analysis of Clostridium beijerinckii mutant with enhanced solvent production. Journal of Biotechnology, 2021, 329, 49-55.	3.8	9
45	Repulsive expansion dynamics in colony growth and gene expression. PLoS Computational Biology, 2021, 17, e1008168.	3.2	5
46	Precise and reliable control of gene expression in <i>Agrobacterium tumefaciens</i> . Biotechnology and Bioengineering, 2021, 118, 3962-3972.	3.3	2
47	MESOSCOPIC CIRCUIT WITH LINEAR DISSIPATION. , 2003, , .		O