

Fu Jing

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

878
citations

430874

18
h-index

677142

22
g-index

30
all docs

30
docs citations

30
times ranked

1057
citing authors

#	ARTICLE	IF	CITATIONS
1	Metabolic engineering of <i>Bacillus subtilis</i> for chiral pure meso-2,3-butanediol production. <i>Biotechnology for Biofuels</i> , 2016, 9, 90.	6.2	80
2	Enhanced anaerobic co-digestion of waste activated sludge and food waste by sulfidated microscale zerovalent iron: Insights in direct interspecies electron transfer mechanism. <i>Bioresource Technology</i> , 2020, 316, 123901.	9.6	67
3	NADH plays the vital role for chiral pure D-2,3-butanediol production in <i>Bacillus subtilis</i> under limited oxygen conditions. <i>Biotechnology and Bioengineering</i> , 2014, 111, 2126-2131.	3.3	63
4	Metabolic engineering of <i>Corynebacterium glutamicum</i> for efficient production of 5-aminolevulinic acid. <i>Biotechnology and Bioengineering</i> , 2016, 113, 1284-1293.	3.3	63
5	Efficient degradation of bisphenol A via peroxydisulfate activation using in-situ N-doped carbon nanoparticles: Structure-function relationship and reaction mechanism. <i>Journal of Colloid and Interface Science</i> , 2021, 586, 551-562.	9.4	52
6	Metabolic engineering of <i>Bacillus subtilis</i> for enhanced production of acetoin. <i>Biotechnology Letters</i> , 2012, 34, 1877-1885.	2.2	51
7	Synergistic adsorption and electrocatalytic reduction of bromate by Pd/N-doped loofah sponge-derived biochar electrode. <i>Journal of Hazardous Materials</i> , 2020, 386, 121651.	12.4	49
8	Evaluating the effect of biochar on mesophilic anaerobic digestion of waste activated sludge and microbial diversity. <i>Bioresource Technology</i> , 2019, 294, 122235.	9.6	48
9	Integrated whole-genome and transcriptome sequence analysis reveals the genetic characteristics of a riboflavin-overproducing <i>Bacillus subtilis</i> . <i>Metabolic Engineering</i> , 2018, 48, 138-149.	7.0	45
10	Engineering <i>Bacillus subtilis</i> for acetoin production from glucose and xylose mixtures. <i>Journal of Biotechnology</i> , 2013, 168, 499-505.	3.8	44
11	In silico metabolic engineering of <i>Bacillus subtilis</i> for improved production of riboflavin, Egl-237, (R,R)-2,3-butanediol and isobutanol. <i>Molecular BioSystems</i> , 2013, 9, 2034.	2.9	42
12	Deregulation of purine pathway in <i>Bacillus subtilis</i> and its use in riboflavin biosynthesis. <i>Microbial Cell Factories</i> , 2014, 13, 101.	4.0	39
13	Establishment of a Markerless Mutation Delivery System in <i>Bacillus subtilis</i> Stimulated by a Double-Strand Break in the Chromosome. <i>PLoS ONE</i> , 2013, 8, e81370.	2.5	37
14	Characterization of genome-reduced <i>Bacillus subtilis</i> strains and their application for the production of guanosine and thymidine. <i>Microbial Cell Factories</i> , 2016, 15, 94.	4.0	36
15	Systematic metabolic engineering of <i>Corynebacterium glutamicum</i> for the industrial-level production of optically pure D-acetoin. <i>Green Chemistry</i> , 2017, 19, 5691-5702.	9.0	36
16	Production of Acetoin through Simultaneous Utilization of Glucose, Xylose, and Arabinose by Engineered <i>Bacillus subtilis</i> . <i>PLoS ONE</i> , 2016, 11, e0159298.	2.5	29
17	Oligosaccharide polyester and triterpenoid saponins from the roots of <i>Polygala japonica</i> . <i>Phytochemistry</i> , 2008, 69, 1617-1624.	2.9	26
18	YAP mediates human decidualization of the uterine endometrial stromal cells. <i>Placenta</i> , 2017, 53, 30-35.	1.5	26

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19	Directed evolution of adenylosuccinate synthetase from <i>Bacillus subtilis</i> and its application in metabolic engineering. <i>Journal of Biotechnology</i> , 2016, 231, 115-121.	3.8	12
20	Engineering peptidoglycan degradation related genes of <i>Bacillus subtilis</i> for better fermentation processes. <i>Bioresource Technology</i> , 2018, 248, 238-247.	9.6	12
21	Three new xanthones from the roots of <i>Polygala japonica</i> Houtt.. <i>Journal of Asian Natural Products Research</i> , 2006, 8, 41-46.	1.4	10
22	Deregulation of purine pathway in. <i>Microbial Cell Factories</i> , 2014, 13, 101.	4.0	10
23	Study of a <i>CRISPR-Cas9</i>-Based Counterselective Method for Large-Scale Deletion of Genome Fragments in <i>Bacillus subtilis</i>. <i>Advanced Materials Research</i> , 0, 634-638, 1076-1080.	0.3	0
24	Title is missing!. , 2020, 15, e0234086.		0
25	Title is missing!. , 2020, 15, e0234086.		0
26	Title is missing!. , 2020, 15, e0234086.		0
27	Title is missing!. , 2020, 15, e0234086.		0
28	Title is missing!. , 2020, 15, e0234086.		0
29	Title is missing!. , 2020, 15, e0234086.		0