

Joshua R Elmore

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

1,300
citations

687363

13
h-index

996975

15
g-index

19
all docs

19
docs citations

19
times ranked

1344
citing authors

#	ARTICLE	IF	CITATIONS
1	Corrigendum to “Engineering glucose metabolism for enhanced muconic acid production in <i>Pseudomonas putida</i> KT2440” [Metab. Eng. 59 (2020) 64–75]. <i>Metabolic Engineering</i> , 2022, 72, 66-67.	7.0	0
2	Engineering <i>Citrobacter freundii</i> using CRISPR/Cas9 system. <i>Journal of Microbiological Methods</i> , 2022, 200, 106533.	1.6	3
3	Production of itaconic acid from alkali pretreated lignin by dynamic two stage bioconversion. <i>Nature Communications</i> , 2021, 12, 2261.	12.8	72
4	Tandem chemical deconstruction and biological upcycling of poly(ethylene terephthalate) to Î²-ketoadipic acid by <i>Pseudomonas putida</i> KT2440. <i>Metabolic Engineering</i> , 2021, 67, 250-261.	7.0	74
5	Metabolic engineering of <i>Pseudomonas putida</i> for increased polyhydroxyalkanoate production from lignin. <i>Microbial Biotechnology</i> , 2020, 13, 290-298.	4.2	120
6	Engineering glucose metabolism for enhanced muconic acid production in <i>Pseudomonas putida</i> KT2440. <i>Metabolic Engineering</i> , 2020, 59, 64-75.	7.0	76
7	Engineered <i>Pseudomonas putida</i> simultaneously catabolizes five major components of corn stover lignocellulose: Glucose, xylose, arabinose, p-coumaric acid, and acetic acid. <i>Metabolic Engineering</i> , 2020, 62, 62-71.	7.0	63
8	Evaluation of chromosomal insertion loci in the <i>Pseudomonas putida</i> KT2440 genome for predictable biosystems design. <i>Metabolic Engineering Communications</i> , 2020, 11, e00139.	3.6	18
9	Innovative Chemicals and Materials from Bacterial Aromatic Catabolic Pathways. <i>Joule</i> , 2019, 3, 1523-1537.	24.0	142
10	Engineered <i>Pseudomonas putida</i> KT2440 co-utilizes galactose and glucose. <i>Biotechnology for Biofuels</i> , 2019, 12, 295.	6.2	15
11	Development of a high efficiency integration system and promoter library for rapid modification of <i>Pseudomonas putida</i> KT2440. <i>Metabolic Engineering Communications</i> , 2017, 5, 1-8.	3.6	93
12	Bipartite recognition of target RNAs activates DNA cleavage by the Type III-B CRISPR-Cas system. <i>Genes and Development</i> , 2016, 30, 447-459.	5.9	212
13	DNA targeting by the type I-G and type I-A CRISPR-Cas systems of <i>Pyrococcus furiosus</i> . <i>Nucleic Acids Research</i> , 2015, 43, gkv1140.	14.5	38
14	Essential Structural and Functional Roles of the Cmr4 Subunit in RNA Cleavage by the Cmr CRISPR-Cas Complex. <i>Cell Reports</i> , 2014, 9, 1610-1617.	6.4	57
15	Programmable plasmid interference by the CRISPR-Cas system in <i>Thermococcus kodakarensis</i> . <i>RNA Biology</i> , 2013, 10, 828-840.	3.1	34
16	Essential Features and Rational Design of CRISPR RNAs that Function with the Cas RAMP Module Complex to Cleave RNAs. <i>Molecular Cell</i> , 2012, 45, 292-302.	9.7	275
17	The CRISPR-Cas system: small RNA-guided invader silencing in prokaryotes. <i>FASEB Journal</i> , 2012, 26, 353.3.	0.5	0