Jeffrey M Skerker

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8451224/publications.pdf

Version: 2024-02-01

361413 552781 26 1,624 20 citations h-index papers

g-index 38 38 38 2002 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	Selection of chromosomal DNA libraries using a multiplex CRISPR system. ELife, 2014, 3, .	6.0	314
2	Rhodosporidium toruloides: a new platform organism for conversion of lignocellulose into terpene biofuels and bioproducts. Biotechnology for Biofuels, 2017, 10, 241.	6.2	150
3	Engineering <i>Rhodosporidium toruloides</i> for increased lipid production. Biotechnology and Bioengineering, 2016, 113, 1056-1066.	3.3	143
4	Indirect and suboptimal control of gene expression is widespread in bacteria. Molecular Systems Biology, 2013, 9, 660.	7.2	111
5	Dissecting a complex chemical stress: chemogenomic profiling of plant hydrolysates. Molecular Systems Biology, 2013, 9, 674.	7.2	103
6	Metabolic engineering of the oleaginous yeast Rhodosporidium toruloides IFO0880 for lipid overproduction during high-density fermentation. Applied Microbiology and Biotechnology, 2016, 100, 9393-9405.	3.6	101
7	Functional genomics of lipid metabolism in the oleaginous yeast Rhodosporidium toruloides. ELife, 2018, 7, .	6.0	98
8	Towards an Informative Mutant Phenotype for Every Bacterial Gene. Journal of Bacteriology, 2014, 196, 3643-3655.	2.2	60
9	Monoterpene production by the carotenogenic yeast Rhodosporidium toruloides. Microbial Cell Factories, 2019, 18, 54.	4.0	59
10	Engineering Kluyveromyces marxianus as a Robust Synthetic Biology Platform Host. MBio, 2018, 9, .	4.1	58
11	A toolset of constitutive promoters for metabolic engineering of Rhodosporidium toruloides. Microbial Cell Factories, 2019, 18, 117.	4.0	50
12	Multiplexed CRISPR-Cas9-Based Genome Editing of <i>Rhodosporidium toruloides</i> . MSphere, 2019, 4,	2.9	47
13	Genetic dissection of interspecific differences in yeast thermotolerance. Nature Genetics, 2018, 50, 1501-1504.	21.4	43
14	Further engineering of R. toruloides for the production of terpenes from lignocellulosic biomass. Biotechnology for Biofuels, 2021, 14, 101.	6.2	31
15	Metabolic engineering of a haploid strain derived from a triploid industrial yeast for producing cellulosic ethanol. Metabolic Engineering, 2017, 40, 176-185.	7.0	27
16	Quantitative Trait Loci (QTL)-Guided Metabolic Engineering of a Complex Trait. ACS Synthetic Biology, 2017, 6, 566-581.	3.8	26
17	Multi-Omics Driven Metabolic Network Reconstruction and Analysis of Lignocellulosic Carbon Utilization in Rhodosporidium toruloides. Frontiers in Bioengineering and Biotechnology, 2020, 8, 612832.	4.1	25
18	Rapid and efficient galactose fermentation by engineered Saccharomyces cerevisiae. Journal of Biotechnology, 2016, 229, 13-21.	3.8	24

#	Article	IF	CITATIONS
19	Transposon insertional mutagenesis in <i>Saccharomyces uvarum</i> reveals <i>trans</i> -acting effects influencing species-dependent essential genes. Genome Research, 2019, 29, 396-406.	5.5	24
20	Complete Genome Sequence of Cupriavidus basilensis 4G11, Isolated from the Oak Ridge Field Research Center Site. Genome Announcements, $2015, 3, \ldots$	0.8	23
21	Fermentation of hydrolysate detoxified by pervaporation through block copolymer membranes. Green Chemistry, 2014, 16, 4206-4213.	9.0	22
22	Exploiting nonionic surfactants to enhance fatty alcohol production in <i>Rhodosporidium toruloides</i> . Biotechnology and Bioengineering, 2020, 117, 1418-1425.	3.3	21
23	Genomewide and Enzymatic Analysis Reveals Efficient <scp>d</scp> -Galacturonic Acid Metabolism in the Basidiomycete Yeast Rhodosporidium toruloides. MSystems, 2019, 4, .	3.8	20
24	Complete Genome Sequences of Four Escherichia coli ST95 Isolates from Bloodstream Infections. Genome Announcements, 2015, 3, .	0.8	18
25	Evolution, ecology and the engineered organism: lessons for synthetic biology. Genome Biology, 2009, 10, 114.	9.6	7
26	Barcoded reciprocal hemizygosity analysis <i>via</i> sequencing illuminates the complex genetic basis of yeast thermotolerance. G3: Genes, Genomes, Genetics, 2022, 12, .	1.8	7