Mian Zhou

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9205007/publications.pdf

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all docs

23 1,263 13 23 papers citations h-index g-index

24 24 24 1500

times ranked

citing authors

docs citations

#	Article	IF	CITATIONS
1	Non-optimal codon usage affects expression, structure and function of clock protein FRQ. Nature, 2013, 495, 111-115.	27.8	357
2	Codon usage is an important determinant of gene expression levels largely through its effects on transcription. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E6117-E6125.	7.1	326
3	Nonoptimal codon usage influences protein structure in intrinsically disordered regions. Molecular Microbiology, 2015, 97, 974-987.	2.5	99
4	Mit1 Transcription Factor Mediates Methanol Signaling and Regulates the Alcohol Oxidase 1 (AOX1) Promoter in Pichia pastoris. Journal of Biological Chemistry, 2016, 291, 6245-6261.	3.4	82
5	Codon usage affects the structure and function of the <i>Drosophila</i> circadian clock protein PERIOD. Genes and Development, 2016, 30, 1761-1775.	5.9	73
6	A novel methanol-free Pichia pastoris system for recombinant protein expression. Microbial Cell Factories, 2016, 15, 178.	4.0	53
7	Codon usage biases co-evolve with transcription termination machinery to suppress premature cleavage and polyadenylation. ELife, 2018, 7, .	6.0	50
8	PpNrg1 is a transcriptional repressor for glucose and glycerol repression of AOX1 promoter in methylotrophic yeast Pichia pastoris. Biotechnology Letters, 2016, 38, 291-298.	2.2	39
9	Identification and characterization of novel promoters for recombinant protein production in yeast <i>Pichia pastoris /i>. Yeast, 2018, 35, 379-385.</i>	1.7	32
10	Kinase Screening in Pichia pastoris Identified Promising Targets Involved in Cell Growth and Alcohol Oxidase 1 Promoter (PAOX1) Regulation. PLoS ONE, 2016, 11, e0167766.	2.5	28
11	Serine/threonine kinase PpkA coordinates the interplay between T6SS2 activation and quorum sensing in the marine pathogen <i>Vibrio alginolyticus</i> Environmental Microbiology, 2018, 20, 903-919.	3.8	25
12	Genome-Wide Determination of Gene Essentiality by Transposon Insertion Sequencing in Yeast Pichia pastoris. Scientific Reports, 2018, 8, 10223.	3.3	25
13	Codon usage bias regulates gene expression and protein conformation in yeast expression system P. pastoris. Microbial Cell Factories, 2021, 20, 91.	4.0	20
14	Heterologous expression and purification of a marine alginate lyase in Escherichia coli. Protein Expression and Purification, 2019, 153, 97-104.	1.3	12
15	EsrB negatively regulates expression of the glutamine sythetase GlnA in the fish pathogen Edwardsiella piscicida. FEMS Microbiology Letters, 2018, 365, .	1.8	7
16	PiggyBac transposon-mediated mutagenesis and application in yeast Komagataella phaffii. Biotechnology Letters, 2018, 40, 1365-1376.	2.2	7
17	Phosphorylation of PppA at threonine 253 controls T6SS2 expression and bacterial killing capacity in the marine pathogen Vibrio alginolyticus. Microbiological Research, 2018, 209, 70-78.	5.3	6
18	Methods to Study Molecular Mechanisms of the Neurospora Circadian Clock. Methods in Enzymology, 2015, 551, 137-151.	1.0	5

#	Article	IF	CITATION
19	Strong negative correlation between codon usage bias and protein structural disorder impedes protein expression after codon optimization. Journal of Biotechnology, 2022, 343, 15-24.	3.8	5
20	Codon usage bias affects α-amylase mRNA level by altering RNA stability and cytosine methylation patterns in <i>Escherichia coli</i> i>. Canadian Journal of Microbiology, 2020, 66, 521-528.	1.7	4
21	Online bioinformatics teaching practice: Comparison of popular docking programs using <scp>SARSâ€CoV</scp> â€2 spike <scp>RBD–ACE2</scp> complex as a benchmark. Biochemistry and Molecular Biology Education, 2021, 49, 833-840.	1.2	4
22	Identification and study of InV as an inverse autotransporter family representative in Edwardsiella piscicida. Archives of Microbiology, 2020, 202, 1107-1116.	2.2	2
23	Enhanced bioproduction of chitin in engineered Pichia pastoris. Food Bioscience, 2022, 47, 101606.	4.4	2