Kyle R Pomraning

List of Publications by Year in descending order

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687363 752698 1,087 19 13 20 citations g-index h-index papers 21 21 21 1808 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The Genomes of Three Uneven Siblings: Footprints of the Lifestyles of Three Trichoderma Species. Microbiology and Molecular Biology Reviews, 2016, 80, 205-327.	6.6	194
2	Genome-wide high throughput analysis of DNA methylation in eukaryotes. Methods, 2009, 47, 142-150.	3.8	178
3	Genome resequencing reveals multiscale geographic structure and extensive linkage disequilibrium in the forest tree <i>Populus trichocarpa</i> New Phytologist, 2012, 196, 713-725.	7.3	173
4	Heterochromatin Is Required for Normal Distribution of Neurospora crassa CenH3. Molecular and Cellular Biology, 2011, 31, 2528-2542.	2.3	107
5	Bulk Segregant Analysis Followed by High-Throughput Sequencing Reveals the Neurospora Cell Cycle Gene, <i> ndc-1 </i> , To Be Allelic with the Gene for Ornithine Decarboxylase, <i> spe-1 </i> . Eukaryotic Cell, 2011, 10, 724-733.	3.4	67
6	Multi-omics analysis reveals regulators of the response to nitrogen limitation in Yarrowia lipolytica. BMC Genomics, 2016, 17, 138.	2.8	62
7	A molecular genetic toolbox for Yarrowia lipolytica. Biotechnology for Biofuels, 2017, 10, 2.	6.2	62
8	Comprehensive Metabolomic, Lipidomic and Microscopic Profiling of Yarrowia lipolytica during Lipid Accumulation Identifies Targets for Increased Lipogenesis. PLoS ONE, 2015, 10, e0123188.	2.5	54
9	Bacterial Resistance to Antisense Peptide Phosphorodiamidate Morpholino Oligomers. Antimicrobial Agents and Chemotherapy, 2012, 56, 6147-6153.	3.2	41
10	Regulation of Nitrogen Metabolism by GATA Zinc Finger Transcription Factors in Yarrowia lipolytica. MSphere, 2017, 2, .	2.9	29
11	Transcriptomic analysis of the oleaginous yeast Lipomyces starkeyi during lipid accumulation on enzymatically treated corn stover hydrolysate. Biotechnology for Biofuels, 2019, 12, 162.	6.2	24
12	Omics Analyses of Trichoderma reesei CBS999.97 and QM6a Indicate the Relevance of Female Fertility to Carbohydrate-Active Enzyme and Transporter Levels. Applied and Environmental Microbiology, 2017, 83,	3.1	22
13	Library Preparation and Data Analysis Packages for Rapid Genome Sequencing. Methods in Molecular Biology, 2012, 944, 1-22.	0.9	17
14	Integration of Proteomics and Metabolomics Into the Design, Build, Test, Learn Cycle to Improve 3-Hydroxypropionic Acid Production in Aspergillus pseudoterreus. Frontiers in Bioengineering and Biotechnology, 2021, 9, 603832.	4.1	12
15	Deletion analysis of the itaconic acid biosynthesis gene cluster components in Aspergillus pseudoterreus ATCC32359. Applied Microbiology and Biotechnology, 2020, 104, 3981-3992.	3.6	10
16	Deletion of the KU70 homologue facilitates gene targeting in Lipomyces starkeyi strain NRRL Y-11558. Current Genetics, 2019, 65, 269-282.	1.7	9
17	Genetically Engineered Oleaginous Yeast <i>Lipomyces starkeyi</i> for Sesquiterpene α-Zingiberene Production. ACS Synthetic Biology, 2021, 10, 1000-1008.	3.8	5
18	Multimodal microfluidic platform for controlled culture and analysis of unicellular organisms. Biomicrofluidics, 2017, 11, 054104.	2.4	4

#	Article	IF	CITATIONS
19	Bayesian Inference for Integrating <i>Yarrowia lipolytica</i> Multiomics Datasets with Metabolic Modeling. ACS Synthetic Biology, 2021, 10, 2968-2981.	3.8	4