

Yo Suzuki

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3735125/publications.pdf>

Version: 2024-02-01

29
papers

2,794
citations

394421

19
h-index

501196

28
g-index

29
all docs

29
docs citations

29
times ranked

4718
citing authors

#	ARTICLE	IF	CITATIONS
1	Adaptive laboratory evolution in <i>S. cerevisiae</i> highlights role of transcription factors in fungal xenobiotic resistance. <i>Communications Biology</i> , 2022, 5, 128.	4.4	8
2	Blockade of endoplasmic reticulum stress-induced cell death by <i>Ureaplasma parvum</i> vacuolating factor. <i>Cellular Microbiology</i> , 2021, 23, e13392.	2.1	10
3	Biotechnology for secure biocontainment designs in an emerging bioeconomy. <i>Current Opinion in Biotechnology</i> , 2021, 71, 25-31.	6.6	23
4	Two inhibitors of yeast plasma membrane ATPase 1 (ScPma1p): toward the development of novel antifungal therapies. <i>Journal of Cheminformatics</i> , 2018, 10, 6.	6.1	17
5	Tuning Gene Activity by Inducible and Targeted Regulation of Gene Expression in Minimal Bacterial Cells. <i>ACS Synthetic Biology</i> , 2018, 7, 1538-1552.	3.8	30
6	The Human Microbiome and Cancer. <i>Cancer Prevention Research</i> , 2017, 10, 226-234.	1.5	230
7	Rapid Chagas Disease Drug Target Discovery Using Directed Evolution in Drug-Sensitive Yeast. <i>ACS Chemical Biology</i> , 2017, 12, 422-434.	3.4	26
8	Open Source Drug Discovery with the Malaria Box Compound Collection for Neglected Diseases and Beyond. <i>PLoS Pathogens</i> , 2016, 12, e1005763.	4.7	244
9	Comparative chemical genomics reveal that the spiroindolone antimalarial KAE609 (Cipargamin) is a P-type ATPase inhibitor. <i>Scientific Reports</i> , 2016, 6, 27806.	3.3	38
10	Design and synthesis of a minimal bacterial genome. <i>Science</i> , 2016, 351, aad6253.	12.6	1,077
11	Cloning Should Be Simple: <i>Escherichia coli</i> DH5 α -Mediated Assembly of Multiple DNA Fragments with Short End Homologies. <i>PLoS ONE</i> , 2015, 10, e0137466.	2.5	104
12	Bacterial genome reduction using the progressive clustering of deletions via yeast sexual cycling. <i>Genome Research</i> , 2015, 25, 435-444.	5.5	27
13	Strategies for cloning and manipulating natural and synthetic chromosomes. <i>Chromosome Research</i> , 2015, 23, 57-68.	2.2	30
14	Genomic and Transcriptomic Analyses of Colistin-Resistant Clinical Isolates of <i>Klebsiella pneumoniae</i> Reveal Multiple Pathways of Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 536-543.	3.2	185
15	Successful Diatom Transcription Factor Synthesis and Downstream Cloning Using the BioXp α 3200 System. <i>BioTechniques</i> , 2015, 59, 46-47.	1.8	0
16	Rescue of mutant fitness defects using in vitro reconstituted designer transposons in <i>Mycoplasma mycoides</i> . <i>Frontiers in Microbiology</i> , 2014, 5, 369.	3.5	12
17	The Insertion Green Monster (iGM) Method for Expression of Multiple Exogenous Genes in Yeast. <i>G3: Genes, Genomes, Genetics</i> , 2014, 4, 1183-1191.	1.8	14
18	Transferring whole genomes from bacteria to yeast spheroplasts using entire bacterial cells to reduce DNA shearing. <i>Nature Protocols</i> , 2014, 9, 743-750.	12.0	37

#	ARTICLE	IF	CITATIONS
19	Direct transfer of whole genomes from bacteria to yeast. <i>Nature Methods</i> , 2013, 10, 410-412.	19.0	64
20	The Green Monster Process for the Generation of Yeast Strains Carrying Multiple Gene Deletions. <i>Journal of Visualized Experiments</i> , 2012, , e4072.	0.3	12
21	Assembly of Large, High G+C Bacterial DNA Fragments in Yeast. <i>ACS Synthetic Biology</i> , 2012, 1, 267-273.	3.8	65
22	Systematic exploration of synergistic drug pairs. <i>Molecular Systems Biology</i> , 2011, 7, 544.	7.2	284
23	Knocking out multigene redundancies via cycles of sexual assortment and fluorescence selection. <i>Nature Methods</i> , 2011, 8, 159-164.	19.0	74
24	Reconstitution of human RNA interference in budding yeast. <i>Nucleic Acids Research</i> , 2011, 39, e43-e43.	14.5	26
25	Systematic genetics swims forward elegantly. <i>Molecular Systems Biology</i> , 2006, 2, 48.	7.2	2
26	Genetic redundancy masks diverse functions of the tumor suppressor gene PTEN during <i>C. elegans</i> development. <i>Genes and Development</i> , 2006, 20, 423-428.	5.9	25
27	Expression of the <i>C. elegans</i> labial orthologue <i>ceh-13</i> during male tail morphogenesis. <i>Developmental Biology</i> , 2003, 259, 137-149.	2.0	11
28	A <i>Caenorhabditis elegans</i> TGF-beta, DBL-1, controls the expression of LON-1, a PR-related protein, that regulates polyploidization and body length. <i>EMBO Journal</i> , 2002, 21, 1063-1073.	7.8	88
29	A Cuticle Collagen Encoded by the <i>lon-3</i> Gene May Be a Target of TGF- β^2 Signaling in Determining <i>Caenorhabditis elegans</i> Body Shape. <i>Genetics</i> , 2002, 162, 1631-1639.	2.9	31