

Maxim Kostylev

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10791012/publications.pdf>

Version: 2024-02-01

12
papers

651
citations

840776

11
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

1041
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamics of cheater invasion in a cooperating population of <i>Pseudomonas aeruginosa</i> . <i>Scientific Reports</i> , 2019, 9, 10190.	3.3	7
2	Evolution of the <i>Pseudomonas aeruginosa</i> quorum-sensing hierarchy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 7027-7032.	7.1	197
3	Modulation of <i>Pseudomonas aeruginosa</i> Quorum Sensing by Glutathione. <i>Journal of Bacteriology</i> , 2019, 201, .	2.2	12
4	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
5	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
6	Bacterial genome reduction using the progressive clustering of deletions via yeast sexual cycling. <i>Genome Research</i> , 2015, 25, 435-444.	5.5	27
7	A Distinct Model of Synergism between a Processive Endocellulase (TfCel9A) and an Exocellulase (TfCel48A) from <i>Thermobifida fusca</i> . <i>Applied and Environmental Microbiology</i> , 2014, 80, 339-344.	3.1	33
8	Cel48A from <i>Thermobifida fusca</i> : Structure and site directed mutagenesis of key residues. <i>Biotechnology and Bioengineering</i> , 2014, 111, 664-673.	3.3	35
9	Two-Parameter Kinetic Model Based on a Time-Dependent Activity Coefficient Accurately Describes Enzymatic Cellulose Digestion. <i>Biochemistry</i> , 2013, 52, 5656-5664.	2.5	34
10	Synergistic interactions in cellulose hydrolysis. <i>Biofuels</i> , 2012, 3, 61-70.	2.4	118
11	Determination of the molecular states of the processive endocellulase <i>Thermobifida fusca</i> Cel9A during crystalline cellulose depolymerization. <i>Biotechnology and Bioengineering</i> , 2012, 109, 295-299.	3.3	30
12	Determination of the Catalytic Base in Family 48 Glycosyl Hydrolases. <i>Applied and Environmental Microbiology</i> , 2011, 77, 6274-6276.	3.1	16