Matthieu Jules

List of Publications by Year in descending order

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Version: 2024-02-01

31	2,986	22	31
papers	citations	h-index	g-index
33	33	33	3388
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Condition-Dependent Transcriptome Reveals High-Level Regulatory Architecture in <i>Bacillus subtilis</i> . Science, 2012, 335, 1103-1106.	12.6	809
2	Global Network Reorganization During Dynamic Adaptations of <i>Bacillus subtilis</i> Metabolism. Science, 2012, 335, 1099-1103.	12.6	255
3	Virulence strategies for infecting phagocytes deduced from the in vivo transcriptional program of Legionella pneumophila. Cellular Microbiology, 2006, 8, 1228-1240.	2.1	241
4	Two small ncRNAs jointly govern virulence and transmission in <i>Legionella pneumophila</i> Molecular Microbiology, 2009, 72, 741-762.	2.5	166
5	A part toolbox to tune genetic expression in <i>Bacillus subtilis</i> . Nucleic Acids Research, 2016, 44, gkw624.	14.5	157
6	Transcriptional regulation is insufficient to explain substrateâ€induced flux changes in <i>Bacillus subtilis</i> . Molecular Systems Biology, 2013, 9, 709.	7.2	149
7	The Legionella pneumophila F-box protein Lpp2082 (AnkB) modulates ubiquitination of the host protein parvin B and promotes intracellular replication. Cellular Microbiology, 2010, 12, 1272-1291.	2.1	134
8	Quantitative prediction of genome-wide resource allocation in bacteria. Metabolic Engineering, 2015, 32, 232-243.	7.0	125
9	Metabolic Fluxes during Strong Carbon Catabolite Repression by Malate in Bacillus subtilis. Journal of Biological Chemistry, 2010, 285, 1587-1596.	3.4	100
10	Two Distinct Pathways for Trehalose Assimilation in the Yeast <i>Saccharomyces cerevisiae</i> Applied and Environmental Microbiology, 2004, 70, 2771-2778.	3.1	97
11	Isotopologue Profiling of Legionella pneumophila. Journal of Biological Chemistry, 2010, 285, 22232-22243.	3.4	95
12	New Insights into Trehalose Metabolism by <i>Saccharomyces cerevisiae</i> : <i>NTH2</i> Encodes a Functional Cytosolic Trehalase, and Deletion of <i>TPS1</i> Reveals Ath1p-Dependent Trehalose Mobilization. Applied and Environmental Microbiology, 2008, 74, 605-614.	3.1	73
13	Reconciling molecular regulatory mechanisms with noise patterns of bacterial metabolic promoters in induced and repressed states. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 155-160.	7.1	71
14	Acid trehalase in yeasts and filamentous fungi: Localization, regulation and physiological function. FEMS Yeast Research, 2005, 5, 503-511.	2.3	64
15	pBaSysBioII: an integrative plasmid generating gfp transcriptional fusions for high-throughput analysis of gene expression in Bacillus subtilis. Microbiology (United Kingdom), 2010, 156, 1600-1608.	1.8	56
16	Translation elicits a growth rateâ€dependent, genomeâ€wide, differential protein production in <i>Bacillus subtilis</i> . Molecular Systems Biology, 2016, 12, 870.	7.2	56
17	Legionella pneumophilaadaptation to intracellular life and the host response: Clues from genomics and transcriptomics. FEBS Letters, 2007, 581, 2829-2838.	2.8	53
18	Autonomous oscillations in Saccharomyces cerevisiae during batch cultures on trehalose. FEBS Journal, 2005, 272, 1490-1500.	4.7	39

#	Article	IF	CITATIONS
19	GamA is a eukaryotic-like glucoamylase responsible for glycogen- and starch-degrading activity of Legionella pneumophila. International Journal of Medical Microbiology, 2011, 301, 133-139.	3.6	36
20	Malate-Mediated Carbon Catabolite Repression in Bacillus subtilis Involves the HPrK/CcpA Pathway. Journal of Bacteriology, 2011, 193, 6939-6949.	2.2	36
21	Molecular and Physiological Logics of the Pyruvate-Induced Response of a Novel Transporter in <i>Bacillus subtilis</i> . MBio, 2017, 8, .	4.1	35
22	The <i>Legionella pneumophila</i> LetA/LetS Two-Component System Exhibits Rheostat-Like Behavior. Infection and Immunity, 2010, 78, 2571-2583.	2.2	30
23	Absolute quantification of gene expression in individual bacterial cells using two-photon fluctuation microscopy. Analytical Biochemistry, 2011, 419, 250-259.	2.4	22
24	Nicotinic Acid Modulates Legionella pneumophila Gene Expression and Induces Virulence Traits. Infection and Immunity, 2013, 81, 945-955.	2,2	19
25	The <i>Bacillus subtilis ywjl</i> (<i>glpX</i>) Gene Encodes a Class II Fructose-1,6-Bisphosphatase, Functionally Equivalent to the Class III Fbp Enzyme. Journal of Bacteriology, 2009, 191, 3168-3171.	2.2	18
26	Bacterial growth physiology and RNA metabolism. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2020, 1863, 194502.	1.9	13
27	Extrinsic noise prevents the independent tuning of gene expression noise and protein mean abundance in bacteria. Science Advances, 2020, 6, .	10.3	11
28	BasyLiCA: a tool for automatic processing of a Bacterial Live Cell Array. Bioinformatics, 2012, 28, 2705-2706.	4.1	8
29	Four species of bacteria deterministically assemble to form a stable biofilm in a millifluidic channel. Npj Biofilms and Microbiomes, 2021, 7, 64.	6.4	8
30	Differentiation of Vegetative Cells into Spores: a Kinetic Model Applied to Bacillus subtilis. Applied and Environmental Microbiology, 2019, 85, .	3.1	6
31	Sppl Forms a Membrane Protein Complex with SppA and Inhibits Its Protease Activity in Bacillus subtilis. MSphere, 2020, 5, .	2.9	3