Colin R. Harwood

List of Publications by Year in descending order

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41 papers

2,058 citations

26 h-index

218677

276875
41
g-index

42 all docs 42 docs citations 42 times ranked 2319 citing authors

#	Article	IF	Citations
1	Bacillus subtilis and its relatives: molecular biological and industrial workhorses. Trends in Biotechnology, 1992, 10, 247-256.	9.3	245
2	Secondary metabolite production and the safety of industrially important members of the Bacillus subtilis group. FEMS Microbiology Reviews, 2018, 42, 721-738.	8.6	162
3	Bacillus protein secretion: an unfolding story. Trends in Microbiology, 2008, 16, 73-79.	7.7	150
4	d-Alanine Substitution of Teichoic Acids as a Modulator of Protein Folding and Stability at the Cytoplasmic Membrane/Cell Wall Interface of Bacillus subtilis. Journal of Biological Chemistry, 2000, 275, 26696-26703.	3.4	100
5	<i>Bacillus subtilis,</i> the model Gramâ€positive bacterium: 20Âyears of annotation refinement. Microbial Biotechnology, 2018, 11, 3-17.	4.2	95
6	Heterologous Protein Secretion by Bacillus Species. Advances in Applied Microbiology, 2010, 73, 1-25.	2.4	93
7	Regulatory interactions between the Pho and $\ddot{l}fB$ -dependent general stress regulons of Bacillus subtilis. Microbiology (United Kingdom), 2002, 148, 1593-1602.	1.8	93
8	Post-translocational folding of secretory proteins in Gram-positive bacteria. Biochimica Et Biophysica Acta - Molecular Cell Research, 2004, 1694, 311-27.	4.1	89
9	Influence of a Cell-Wall-Associated Protease on Production of α-Amylase by <i>Bacillus subtilis</i> Applied and Environmental Microbiology, 1998, 64, 2875-2881.	3.1	77
10	Sequencing and functional analysis of the genome of Bacillus subtilisstrain 168. FEBS Letters, 1996, 389, 84-87.	2.8	75
11	Genome-Wide Transcriptional Analysis of the Phosphate Starvation Stimulon of Bacillus subtilis. Journal of Bacteriology, 2005, 187, 8063-8080.	2.2	69
12	Extracytoplasmic Proteases Determining the Cleavage and Release of Secreted Proteins, Lipoproteins, and Membrane Proteins in <i>Bacillus subtilis</i> Journal of Proteome Research, 2013, 12, 4101-4110.	3.7	64
13	Transcriptional Regulation of the phoPR Operon in Bacillus subtilis. Journal of Bacteriology, 2004, 186, 1182-1190.	2.2	61
14	Effect of Genome Position on Heterologous Gene Expression in <i>Bacillus subtilis</i> An Unbiased Analysis. ACS Synthetic Biology, 2016, 5, 942-947.	3.8	56
15	The extracellular and cytoplasmic proteomes of the non-virulentBacillus anthracis strain UM23C1-2. Proteomics, 2005, 5, 3684-3695.	2.2	54
16	Denitrification, anaerobic ammonium oxidation, and dissimilatory nitrate reduction to ammonium in an East African Great Lake (Lake Kivu). Limnology and Oceanography, 2018, 63, 687-701.	3.1	46
17	Proteomic analysis of <i>Bacillus subtilis</i> strains engineered for improved production of heterologous proteins. Proteomics, 2013, 13, 3298-3308.	2.2	44
18	The influence of protein folding on late stages of the secretion of α-amylases fromBacillus subtilis. FEBS Letters, 1998, 430, 385-389.	2.8	43

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19	Post-transcriptional regulation of the Bacillus subtilis pst operon encoding a phosphate-specific ABC transporter. Microbiology (United Kingdom), 2004, 150, 2619-2628.	1.8	39
20	Extracellular Self-Assembly of Functional and Tunable Protein Conjugates from <i>Bacillus subtilis</i> LACS Synthetic Biology, 2017, 6, 957-967.	3.8	38
21	Exploring the Nonconserved Sequence Space of Synthetic Expression Modules in <i>Bacillus subtilis</i> . ACS Synthetic Biology, 2018, 7, 1773-1784.	3.8	38
22	Environmental Salinity Determines the Specificity and Need for Tat-Dependent Secretion of the YwbN Protein in Bacillus subtilis. PLoS ONE, 2011, 6, e18140.	2.5	36
23	Emission and oxidation of methane in a meromictic, eutrophic and temperate lake (Dendre, Belgium). Chemosphere, 2017, 168, 756-764.	8.2	34
24	The ins and outs of <i>Bacillus</i> proteases: activities, functions and commercial significance. FEMS Microbiology Reviews, 2022, 46, .	8.6	33
25	Phosphate Starvation Induces the Sporulation Killing Factor of Bacillus subtilis. Journal of Bacteriology, 2006, 188, 5299-5303.	2.2	29
26	Combined proteomic and transcriptomic analysis of the response of $\langle i \rangle$ Bacillus anthracis $\langle i \rangle$ to oxidative stress. Proteomics, 2011, 11, 3036-3055.	2.2	28
27	The influence of secretory-protein charge on late stages of secretion from the Gram-positive bacterium Bacillus subtilis. Biochemical Journal, 2000, 350, 31-39.	3.7	25
28	e-Science: relieving bottlenecks in large-scale genome analyses. Nature Reviews Microbiology, 2008, 6, 948-954.	28.6	22
29	An Internal FK506-Binding Domain is the Catalytic Core of the Prolyl Isomerase Activity Associated with the Bacillus Subtilis Trigger Factor. FEBS Journal, 1997, 244, 59-65.	0.2	20
30	Comparative analysis of the responses of related pathogenic and environmental bacteria to oxidative stress. Microbiology (United Kingdom), 2012, 158, 636-647.	1.8	16
31	Impact of mass migrations on the clonal variation of clinical Staphylococcus aureus strains isolated from the Western region of Saudi Arabia. Journal of Infection and Public Health, 2019, 12, 317-322.	4.1	14
32	Effects of phosphate limitation on agarase production by Streptomyces lividans TK21. FEMS Microbiology Letters, 1998, 158, 107-113.	1.8	12
33	Binase-like guanyl-preferring ribonucleases are new members of Bacillus PhoP regulon. Microbiological Research, 2015, 170, 131-138.	5.3	10
34	From Gene Regulation to Gene Function: Regulatory Networks in Bacillus subtilis. Comparative and Functional Genomics, 2002, 3, 37-41.	2.0	9
35	Nitrate-dependent anaerobic methane oxidation and chemolithotrophic denitrification in a temperate eutrophic lake. FEMS Microbiology Ecology, 2021, 97, .	2.7	9
36	The Bacillus subtilis genome sequence: the molecular blueprint of a soil bacterium. FEMS Microbiology Ecology, 1999, 28, 1-9.	2.7	9

#	Article	IF	CITATIONS
37	From Genome to Function: Systematic Analysis of the Soil BacteriumBacillus subtilis. Comparative and Functional Genomics, 2001, 2, 22-24.	2.0	7
38	The possible occurrence of iron-dependent anaerobic methane oxidation in an Archean Ocean analogue. Scientific Reports, 2021, 11, 1597.	3. 3	6
39	Editorial for the thematic issue on "Industrial Microbiology― FEMS Microbiology Letters, 2018, 365, .	1.8	3
40	Whole genome microarray analysis of the expression profile of Escherichia coli in response to exposure to para-nitrophenol. Advances in Experimental Biology, 2008, 2, 221-248.	0.1	2
41	Studies on the influence of the cyclic AMP system on major outer membrane proteins of Escherichia coli k12. FEMS Microbiology Letters, 1980, 9, 95-98.	1.8	2