## John Chen

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3505358/publications.pdf

Version: 2024-02-01

41 3,396 26
papers citations h-index

39 g-index

44 44 all docs docs citations

44 times ranked 4301 citing authors

#	Article	IF	Citations
1	Shared signatures and divergence in skin microbiomes of children with atopic dermatitis and their caregivers. Journal of Allergy and Clinical Immunology, 2022, 150, 894-908.	2.9	14
2	inPhocus: Current State and Challenges of Phage Research in Singapore. Phage, 2022, 3, 6-11.	1.7	0
3	Phage-inducible chromosomal islands promote genetic variability by blocking phage reproduction and protecting transductants from phage lysis. PLoS Genetics, 2022, 18, e1010146.	3.5	8
4	Regulatory cascade in SaPI activation. Access Microbiology, 2022, 4, .	0.5	0
5	Characterisation of novel functionality within the Blastocystis tryptophanase gene. PLoS Neglected Tropical Diseases, 2021, 15, e0009730.	3.0	2
6	A regulatory cascade controls Staphylococcus aureus pathogenicity island activation. Nature Microbiology, 2021, 6, 1300-1308.	13.3	20
7	Staphylococcal phages and pathogenicity islands drive plasmid evolution. Nature Communications, 2021, 12, 5845.	12.8	26
8	Lateral transduction is inherent to the life cycle of the archetypical Salmonella phage P22. Nature Communications, 2021, 12, 6510.	12.8	30
9	Bacterial chromosomal mobility via lateral transduction exceeds that of classical mobile genetic elements. Nature Communications, 2021, 12, 6509.	12.8	46
10	Sequencing identifies multiple early introductions of SARS-CoV-2 to the New York City region. Genome Research, 2020, 30, 1781-1788.	5.5	66
11	Beyond the CRISPR-Cas safeguard: PICI-encoded innate immune systems protect bacteria from bacteriophage predation. Current Opinion in Microbiology, 2020, 56, 52-58.	5.1	28
12	Genetic transduction by phages and chromosomal islands: The new and noncanonical. PLoS Pathogens, 2019, 15, e1007878.	4.7	111
13	Hijacking the Hijackers: Escherichia coli Pathogenicity Islands Redirect Helper Phage Packaging for Their Own Benefit. Molecular Cell, 2019, 75, 1020-1030.e4.	9.7	45
14	Genome hypermobility by lateral transduction. Science, 2018, 362, 207-212.	12.6	187
15	Phage-inducible chromosomal islands are ubiquitous within the bacterial universe. ISME Journal, 2018, 12, 2114-2128.	9.8	115
16	Effect of food-related stress conditions and loss of agr and sigB on seb promoter activity in S.Âaureus. Food Microbiology, 2017, 65, 205-212.	4.2	15
17	Phage-inducible islands in the Gram-positive cocci. ISME Journal, 2017, 11, 1029-1042.	9.8	82
18	Tuning of the Lethal Response to Multiple Stressors with a Single-Site Mutation during Clinical Infection by <i>Staphylococcus aureus</i> ). MBio, 2017, 8, .	4.1	15

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19	Bacterial Hypoxic Responses Revealed as Critical Determinants of the Host-Pathogen Outcome by TnSeq Analysis of Staphylococcus aureus Invasive Infection. PLoS Pathogens, 2015, 11, e1005341.	4.7	118
20	Pathogenicity Island-Directed Transfer of Unlinked Chromosomal Virulence Genes. Molecular Cell, 2015, 57, 138-149.	9.7	52
21	An rpsL-based allelic exchange vector for Staphylococcus aureus. Plasmid, 2015, 79, 8-14.	1.4	11
22	An insight into staphylococcal pathogenicity island-mediated interference with phage late gene transcription. Bacteriophage, 2015, 5, e1028608.	1.9	4
23	Bacteriophage-mediated spread of bacterial virulence genes. Current Opinion in Microbiology, 2015, 23, 171-178.	5.1	268
24	Intra- and inter-generic transfer of pathogenicity island-encoded virulence genes by <i>cos</i> phages. ISME Journal, 2015, 9, 1260-1263.	9.8	49
25	Precisely modulated pathogenicity island interference with late phage gene transcription. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 14536-14541.	7.1	60
26	Single-copy vectors for integration at the SaPI1 attachment site for Staphylococcus aureus. Plasmid, 2014, 76, 1-7.	1.4	54
27	Allele-Dependent Differences in Quorum-Sensing Dynamics Result in Variant Expression of Virulence Genes in Staphylococcus aureus. Journal of Bacteriology, 2012, 194, 2854-2864.	2.2	54
28	Identification of ligand specificity determinants in AgrC, the Staphylococcus aureus quorum-sensing receptor Journal of Biological Chemistry, 2012, 287, 18588.	3.4	1
29	Staphylococcal pathogenicity island interference with helper phage reproduction is a paradigm of molecular parasitism. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 16300-16305.	7.1	113
30	Significant and vertically coherent seismic anisotropy beneath eastern Tibet. Journal of Geophysical Research, 2012, 117, .	3.3	46
31	<i>Staphylococcus aureus</i> leucocidin ED contributes to systemic infection by targeting neutrophils and promoting bacterial growth <i>in vivo</i> . Molecular Microbiology, 2012, 83, 423-435.	2.5	134
32	Control of <i>Staphylococcus aureus</i> pathogenicity island excision. Molecular Microbiology, 2012, 85, 833-845.	2.5	40
33	Rayleigh wave phase velocity maps of Tibet and the surrounding regions from ambient seismic noise tomography. Geochemistry, Geophysics, Geosystems, $2010,11,$ .	2.5	105
34	Underplating in the Himalaya-Tibet Collision Zone Revealed by the Hi-CLIMB Experiment. Science, 2009, 325, 1371-1374.	12.6	662
35	Noninvasive Realâ€Time Monitoring of Liverâ€Stage Development of Bioluminescent <i>Plasmodium</i> Parasites. Journal of Infectious Diseases, 2009, 200, 1470-1478.	4.0	52
36	Phage-Mediated Intergeneric Transfer of Toxin Genes. Science, 2009, 323, 139-141.	12.6	271

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37	Identification of Ligand Specificity Determinants in AgrC, the Staphylococcus aureus Quorum-sensing Receptor. Journal of Biological Chemistry, 2008, 283, 8930-8938.	3.4	88
38	svrA, a multi-drug exporter, does not control agr. Microbiology (United Kingdom), 2007, 153, 1604-1608.	1.8	14
39	Host cell-dependent secretion and translocation of the LepA and LepB effectors of Legionella pneumophila. Cellular Microbiology, 2007, 9, 1660-1671.	2.1	56
40	Legionella Effectors That Promote Nonlytic Release from Protozoa. Science, 2004, 303, 1358-1361.	12.6	271
41	Escherichia coli nusG mutations that block transcription termination by coliphage HK022 Nun protein. Molecular Microbiology, 1999, 31, 1783-1793.	2.5	26