Alex Wong

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/597824/publications.pdf

Version: 2024-02-01

233421 331670 2,371 49 21 45 citations h-index g-index papers 49 49 49 3578 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Hybrid Nanopore-Illumina Assemblies for Five Extraintestinal Pathogenic Escherichia coli Isolates. Microbiology Resource Announcements, 2021, 10, .	0.6	3
2	Folate Intake Alters Mutation Frequency and Profiles in a Tissue- and Dose-Specific Manner in MutaMouse Male Mice. Journal of Nutrition, 2021, 151, 800-809.	2.9	2
3	Mode of action of nisin on Escherichia coli. Canadian Journal of Microbiology, 2020, 66, 161-168.	1.7	8
4	Governing antimicrobial resistance: a narrative review of global governance mechanisms. Journal of Public Health Policy, 2020, 41, 515-528.	2.0	26
5	An sRNA Screen for Reversal of Quinolone Resistance in <i>Escherichia coli</i> . G3: Genes, Genomes, Genetics, 2020, 10, 79-88.	1.8	9
6	The effect of environmental heterogeneity on the fitness of antibiotic resistance mutations in Escherichia coli. Evolutionary Ecology, 2020, 34, 379-390.	1.2	13
7	Systematic Evaluation of Whole Genome Sequence-Based Predictions of Salmonella Serotype and Antimicrobial Resistance. Frontiers in Microbiology, 2020, $11,549$.	3.5	53
8	The mutational landscape of quinolone resistance in Escherichia coli. PLoS ONE, 2019, 14, e0224650.	2.5	25
9	The MTHFR 677C>T polymorphism is associated with unmetabolized folic acid in breast milk in a cohort of Canadian women. American Journal of Clinical Nutrition, 2019, 110, 401-409.	4.7	13
10	Unknown Risk on the Farm: Does Agricultural Use of Ionophores Contribute to the Burden of Antimicrobial Resistance?. MSphere, 2019, 4, .	2.9	20
11	Insights into the suitability of utilizing brown rats (Rattus norvegicus) as a model for healing spinal cord injury with epidermal growth factor and fibroblast growth factor-II by predicting protein-protein interactions. Computers in Biology and Medicine, 2019, 104, 220-226.	7.0	5
12	In Silico Engineering of Synthetic Binding Proteins from Random Amino Acid Sequences. IScience, 2019, 11, 375-387.	4.1	10
13	Fitness Tradeoffs of Antibiotic Resistance in Extraintestinal Pathogenic Escherichia coli. Genome Biology and Evolution, 2018, 10, 667-679.	2.5	49
14	Plasmid persistence: costs, benefits, and the plasmid paradox. Canadian Journal of Microbiology, 2018, 64, 293-304.	1.7	127
15	COMPASS: the COMPletely Arbitrary Sequence Simulator. Bioinformatics, 2017, 33, 3101-3103.	4.1	3
16	Implications of the gut microbiota in vulnerability to the social avoidance effects of chronic social defeat in male mice. Brain, Behavior, and Immunity, 2017, 66, 45-55.	4.1	83
17	Epistasis and the Evolution of Antimicrobial Resistance. Frontiers in Microbiology, 2017, 8, 246.	3.5	85
18	Multiplexed Single Intact Cell Droplet Digital PCR (MuSIC ddPCR) Method for Specific Detection of Enterohemorrhagic E. coli (EHEC) in Food Enrichment Cultures. Frontiers in Microbiology, 2017, 8, 332.	3.5	29

#	Article	IF	CITATIONS
19	Evolution of protein-protein interaction networks in yeast. PLoS ONE, 2017, 12, e0171920.	2.5	24
20	Predicting novel protein-protein interactions between the HIV-1 virus and homo sapiens. , 2016, , .		6
21	The evolution of gene expression and binding specificity of the largest transcription factor family in primates. Evolution; International Journal of Organic Evolution, 2016, 70, 167-180.	2.3	12
22	Clinical Isolates of <i>Pseudomonas aeruginosa</i> from Chronically Infected Cystic Fibrosis Patients Fail To Activate the Inflammasome during Both Stable Infection and Pulmonary Exacerbation. Journal of Immunology, 2016, 196, 3097-3108.	0.8	28
23	Effects of genotype on rates of substitution during experimental evolution. Evolution; International Journal of Organic Evolution, 2015, 69, 1772-1785.	2.3	14
24	Draft Genome Sequence of Hafnia paralvei Strain GTA-HAF03. Genome Announcements, 2015, 3, .	0.8	0
25	Quantifying polymorphism and divergence from epigenetic data: a framework for inferring the action of selection. Frontiers in Genetics, 2015, 6, 190.	2.3	0
26	Complete Genome Sequences of Citrobacter braakii Strains GTA-CB01 and GTA-CB04, Isolated from Ground Beef. Genome Announcements, 2015, 3, .	0.8	6
27	Genetic evidence for mixed broods and extra-pair matings in a socially monogamous biparental cichlidÂfish. Behaviour, 2015, 152, 1507-1526.	0.8	11
28	Don't pull the plug! the <i>Drosophila</i> mating plug preserves fertility. Fly, 2015, 9, 62-67.	1.7	22
29	The fitness costs of antibiotic resistance mutations. Evolutionary Applications, 2015, 8, 273-283.	3.1	490
30	Sexual Conflict and Seminal Fluid Proteins: A Dynamic Landscape of Sexual Interactions. Cold Spring Harbor Perspectives in Biology, 2015, 7, a017533.	5.5	123
31	Efficient prediction of human protein-protein interactions at a global scale. BMC Bioinformatics, 2014, 15, 383.	2.6	32
32	Identification and Characterization of Seminal Fluid Proteins in the Asian Tiger Mosquito, Aedes albopictus. PLoS Neglected Tropical Diseases, 2014, 8, e2946.	3.0	63
33	Covariance between Testes Size and Substitution Rates in Primates. Molecular Biology and Evolution, 2014, 31, 1432-1436.	8.9	21
34	Selection on the <i>Drosophila</i> seminal fluid protein Acp62F. Ecology and Evolution, 2013, 3, 1942-1950.	1.9	8
35	Genomics of Adaptation during Experimental Evolution of the Opportunistic Pathogen Pseudomonas aeruginosa. PLoS Genetics, 2012, 8, e1002928.	3.5	139
36	Temporally Variable Selection on Proteolysis-Related Reproductive Tract Proteins in Drosophila. Molecular Biology and Evolution, 2012, 29, 229-238.	8.9	12

#	ARTICLE	IF	CITATION
37	Evolutionary insight from wholeâ€genome sequencing of experimentally evolved microbes. Molecular Ecology, 2012, 21, 2058-2077.	3.9	128
38	Evolution of Drosophila seminal proteins and their networks. , 2012, , 144-152.		3
39	The Molecular Evolution of Animal Reproductive Tract Proteins: What Have We Learned from Mating-System Comparisons?. International Journal of Evolutionary Biology, 2011, 2011, 1-9.	1.0	28
40	Parallel evolution and local differentiation in quinolone resistance in Pseudomonas aeruginosa. Microbiology (United Kingdom), 2011, 157, 937-944.	1.8	52
41	TESTING THE EFFECTS OF MATING SYSTEM VARIATION ON RATES OF MOLECULAR EVOLUTION IN PRIMATES. Evolution; International Journal of Organic Evolution, 2010, 64, 2779-2785.	2.3	21
42	Immortal coils: Conserved dimerization motifs of the Drosophila ovulation prohormone ovulin. Insect Biochemistry and Molecular Biology, 2010, 40, 303-310.	2.7	3
43	Evidence for Positive Selection on Drosophila melanogaster Seminal Fluid Protease Homologs. Molecular Biology and Evolution, 2008, 25, 497-506.	8.9	54
44	A Role for Acp29AB, a Predicted Seminal Fluid Lectin, in Female Sperm Storage in <i>Drosophila melanogaster</i> . Genetics, 2008, 180, 921-931.	2.9	88
45	Evolution in the Fast Lane: Rapidly Evolving Sex-Related Genes in Drosophila. Genetics, 2007, 177, 1321-1335.	2.9	330
46	Phylogenetic incongruence in the Drosophila melanogaster species group. Molecular Phylogenetics and Evolution, 2007, 43, 1138-1150.	2.7	30
47	Sexual Behavior: A Seminal Peptide Stimulates Appetites. Current Biology, 2006, 16, R256-R257.	3.9	8
48	Evidence for structural constraint on ovulin, a rapidly evolving Drosophila melanogaster seminal protein. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 18644-18649.	7.1	21
49	Differentiation between subpopulations of a polychromatic damselfly with respect to morph frequencies, but not neutral genetic markers. Molecular Ecology, 2003, 12, 3505-3513.	3.9	31