## Alex Wong

## List of Publications by Year in descending order

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ALEX WONC

#	Article	IF	CITATIONS
1	The fitness costs of antibiotic resistance mutations. Evolutionary Applications, 2015, 8, 273-283.	3.1	490
2	Evolution in the Fast Lane: Rapidly Evolving Sex-Related Genes in Drosophila. Genetics, 2007, 177, 1321-1335.	2.9	330
3	Genomics of Adaptation during Experimental Evolution of the Opportunistic Pathogen Pseudomonas aeruginosa. PLoS Genetics, 2012, 8, e1002928.	3.5	139
4	Evolutionary insight from wholeâ€genome sequencing of experimentally evolved microbes. Molecular Ecology, 2012, 21, 2058-2077.	3.9	128
5	Plasmid persistence: costs, benefits, and the plasmid paradox. Canadian Journal of Microbiology, 2018, 64, 293-304.	1.7	127
6	Sexual Conflict and Seminal Fluid Proteins: A Dynamic Landscape of Sexual Interactions. Cold Spring Harbor Perspectives in Biology, 2015, 7, a017533.	5.5	123
7	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
8	Epistasis and the Evolution of Antimicrobial Resistance. Frontiers in Microbiology, 2017, 8, 246.	3.5	85
9	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
10	Identification and Characterization of Seminal Fluid Proteins in the Asian Tiger Mosquito, Aedes albopictus. PLoS Neglected Tropical Diseases, 2014, 8, e2946.	3.0	63
11	Evidence for Positive Selection on Drosophila melanogaster Seminal Fluid Protease Homologs. Molecular Biology and Evolution, 2008, 25, 497-506.	8.9	54
12	Systematic Evaluation of Whole Genome Sequence-Based Predictions of Salmonella Serotype and Antimicrobial Resistance. Frontiers in Microbiology, 2020, 11, 549.	3.5	53
13	Parallel evolution and local differentiation in quinolone resistance in Pseudomonas aeruginosa. Microbiology (United Kingdom), 2011, 157, 937-944.	1.8	52
14	Fitness Tradeoffs of Antibiotic Resistance in Extraintestinal Pathogenic Escherichia coli. Genome Biology and Evolution, 2018, 10, 667-679.	2.5	49
15	Efficient prediction of human protein-protein interactions at a global scale. BMC Bioinformatics, 2014, 15, 383.	2.6	32
16	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
17	Phylogenetic incongruence in the Drosophila melanogaster species group. Molecular Phylogenetics and Evolution, 2007, 43, 1138-1150.	2.7	30
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

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19	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
20	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
21	Governing antimicrobial resistance: a narrative review of global governance mechanisms. Journal of Public Health Policy, 2020, 41, 515-528.	2.0	26
22	The mutational landscape of quinolone resistance in Escherichia coli. PLoS ONE, 2019, 14, e0224650.	2.5	25
23	Evolution of protein-protein interaction networks in yeast. PLoS ONE, 2017, 12, e0171920.	2.5	24
24	Don't pull the plug! the <i>Drosophila</i> mating plug preserves fertility. Fly, 2015, 9, 62-67.	1.7	22
25	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
26	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
27	Covariance between Testes Size and Substitution Rates in Primates. Molecular Biology and Evolution, 2014, 31, 1432-1436.	8.9	21
28	Unknown Risk on the Farm: Does Agricultural Use of Ionophores Contribute to the Burden of Antimicrobial Resistance?. MSphere, 2019, 4, .	2.9	20
29	Effects of genotype on rates of substitution during experimental evolution. Evolution; International Journal of Organic Evolution, 2015, 69, 1772-1785.	2.3	14
30	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
31	The effect of environmental heterogeneity on the fitness of antibiotic resistance mutations in Escherichia coli. Evolutionary Ecology, 2020, 34, 379-390.	1.2	13
32	Temporally Variable Selection on Proteolysis-Related Reproductive Tract Proteins in Drosophila. Molecular Biology and Evolution, 2012, 29, 229-238.	8.9	12
33	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
34	Genetic evidence for mixed broods and extra-pair matings in a socially monogamous biparental cichlidÂfish. Behaviour, 2015, 152, 1507-1526.	0.8	11
35	In Silico Engineering of Synthetic Binding Proteins from Random Amino Acid Sequences. IScience, 2019, 11, 375-387.	4.1	10
36	An sRNA Screen for Reversal of Quinolone Resistance in <i>Escherichia coli</i> . G3: Genes, Genomes, Genetics, 2020, 10, 79-88.	1.8	9

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37	Sexual Behavior: A Seminal Peptide Stimulates Appetites. Current Biology, 2006, 16, R256-R257.	3.9	8
38	Selection on the <i>Drosophila</i> seminal fluid protein Acp62F. Ecology and Evolution, 2013, 3, 1942-1950.	1.9	8
39	Mode of action of nisin on Escherichia coli. Canadian Journal of Microbiology, 2020, 66, 161-168.	1.7	8
40	Complete Genome Sequences of Citrobacter braakii Strains GTA-CB01 and GTA-CB04, Isolated from Ground Beef. Genome Announcements, 2015, 3, .	0.8	6
41	Predicting novel protein-protein interactions between the HIV-1 virus and homo sapiens. , 2016, , .		6
42	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
43	Immortal coils: Conserved dimerization motifs of the Drosophila ovulation prohormone ovulin. Insect Biochemistry and Molecular Biology, 2010, 40, 303-310.	2.7	3
44	COMPASS: the COMPletely Arbitrary Sequence Simulator. Bioinformatics, 2017, 33, 3101-3103.	4.1	3
45	Hybrid Nanopore-Illumina Assemblies for Five Extraintestinal Pathogenic Escherichia coli Isolates. Microbiology Resource Announcements, 2021, 10, .	0.6	3
46	Evolution of Drosophila seminal proteins and their networks. , 2012, , 144-152.		3
47	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
48	Draft Genome Sequence of Hafnia paralvei Strain GTA-HAF03. Genome Announcements, 2015, 3, .	0.8	0
49	Quantifying polymorphism and divergence from epigenetic data: a framework for inferring the action of selection. Frontiers in Genetics, 2015, 6, 190.	2.3	0