Frederic Bertels

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/585012/publications.pdf Version: 2024-02-01



4

#	Article	IF	CITATIONS
1	Transposable elements promote the evolution of genome streamlining. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, 20200477.	4.0	14
2	How sequence populations persist inside bacterial genomes. Genetics, 2021, 217, .	2.9	10
3	Parallel Evolution of HIV-1 in a Long-Term Experiment. Molecular Biology and Evolution, 2019, 36, 2400-2414.	8.9	25
4	Repeated Phenotypic Evolution by Different Genetic Routes in Pseudomonas fluorescens SBW25. Molecular Biology and Evolution, 2019, 36, 1071-1085.	8.9	18
5	Estimating the mutational fitness effects distribution during early HIV infection. Virus Evolution, 2018, 4, vey029.	4.9	12
6	Dissecting HIV Virulence: Heritability of Setpoint Viral Load, CD4+ T-Cell Decline, and Per-Parasite Pathogenicity. Molecular Biology and Evolution, 2018, 35, 27-37.	8.9	37
7	Recombination of Virulence Genes in Divergent Acidovorax avenae Strains That Infect a Common Host. Molecular Plant-Microbe Interactions, 2017, 30, 813-828.	2.6	15
8	Discovering Complete Quasispecies in Bacterial Genomes. Genetics, 2017, 206, 2149-2157.	2.9	21
9	Identification and Characterization of Domesticated Bacterial Transposases. Genome Biology and Evolution, 2017, 9, 2110-2121.	2.5	13
10	Bistability in a Metabolic Network Underpins the De Novo Evolution of Colony Switching in Pseudomonas fluorescens. PLoS Biology, 2015, 13, e1002109.	5.6	78
11	Automated Reconstruction of Whole-Genome Phylogenies from Short-Sequence Reads. Molecular Biology and Evolution, 2014, 31, 1077-1088.	8.9	399
12	Adaptive Divergence in Experimental Populations of <i>Pseudomonas fluorescens</i> . V. Insight into the Niche Specialist Fuzzy Spreader Compels Revision of the Model <i>Pseudomonas</i> Radiation. Genetics, 2013, 195, 1319-1335.	2.9	48
13	Genomic Analysis of the Kiwifruit Pathogen Pseudomonas syringae pv. actinidiae Provides Insight into the Origins of an Emergent Plant Disease. PLoS Pathogens, 2013, 9, e1003503.	4.7	247
14	Curiosities of REPINs and RAYTs. Mobile Genetic Elements, 2011, 1, 262-301.	1.8	20
15	Within-Genome Evolution of REPINs: a New Family of Miniature Mobile DNA in Bacteria. PLoS Genetics, 2011, 7, e1002132.	3.5	45
16	Global identification of yeast chromosome interactions using Genome conformation capture. Fungal Genetics and Biology, 2009, 46, 879-886.	2.1	109
17	The perils of plenty: what are we going to do with all these genes?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 3893-3902.	4.0	81

18 Convergent evolution as an indicator for selection during acute HIV-1 infection. , 0, 1, .

2