Geoffrey L Winsor

List of Publications by Year in descending order

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29 papers

8,005 citations

304743 22 h-index 28 g-index

31 all docs

31 docs citations

31 times ranked 12132 citing authors

#	Article	IF	CITATIONS
1	CARD 2020: antibiotic resistome surveillance with the comprehensive antibiotic resistance database. Nucleic Acids Research, 2020, 48, D517-D525.	14.5	1,605
2	IslandViewer 4: expanded prediction of genomic islands for larger-scale datasets. Nucleic Acids Research, 2017, 45, W30-W35.	14.5	1,251
3	InnateDB: systems biology of innate immunity and beyondâ€"recent updates and continuing curation. Nucleic Acids Research, 2013, 41, D1228-D1233.	14.5	1,073
4	Enhanced annotations and features for comparing thousands of <i>Pseudomonas </i> pseudomonas genome database. Nucleic Acids Research, 2016, 44, D646-D653.	14.5	929
5	Pseudomonas Genome Database: improved comparative analysis and population genomics capability for Pseudomonas genomes. Nucleic Acids Research, 2011, 39, D596-D600.	14.5	558
6	InnateDB: facilitating systemsâ€level analyses of the mammalian innate immune response. Molecular Systems Biology, 2008, 4, 218.	7.2	330
7	Newly introduced genomic prophage islands are critical determinants of in vivo competitiveness in the Liverpool Epidemic Strain of <i>Pseudomonas aeruginosa</i> . Genome Research, 2009, 19, 12-23.	5 . 5	317
8	IslandViewer 3: more flexible, interactive genomic island discovery, visualization and analysis: Figure 1 Nucleic Acids Research, 2015, 43, W104-W108.	14.5	316
9	The Burkholderia Genome Database: facilitating flexible queries and comparative analyses. Bioinformatics, 2008, 24, 2803-2804.	4.1	245
10	Pseudomonas Genome Database: facilitating user-friendly, comprehensive comparisons of microbial genomes. Nucleic Acids Research, 2009, 37, D483-D488.	14.5	220
11	Contribution of the PhoP-PhoQ and PmrA-PmrB Two-Component Regulatory Systems to Mg 2+ -Induced Gene Regulation in Pseudomonas aeruginosa. Journal of Bacteriology, 2006, 188, 3995-4006.	2.2	188
12	Construction of a mini-Tn <i>>5-luxCDABE</i> mutant library in <i>Pseudomonas aeruginosa</i> PAO1: A tool for identifying differentially regulated genes. Genome Research, 2005, 15, 583-589.	5 . 5	150
13	Clinical utilization of genomics data produced by the international Pseudomonas aeruginosa consortium. Frontiers in Microbiology, 2015, 6, 1036.	3.5	144
14	Decreasing antibiotic use, the gut microbiota, and asthma incidence in children: evidence from population-based and prospective cohort studies. Lancet Respiratory Medicine, the, 2020, 8, 1094-1105.	10.7	138
15	Pseudomonas aeruginosa Genome Database and PseudoCAP: facilitating community-based, continually updated, genome annotation. Nucleic Acids Research, 2004, 33, D338-D343.	14.5	129
16	OrtholugeDB: a bacterial and archaeal orthology resource for improved comparative genomic analysis. Nucleic Acids Research, 2013, 41, D366-D376.	14.5	73
17	Curating the innate immunity interactome. BMC Systems Biology, 2010, 4, 117.	3.0	68
18	Sexual reproduction in Daphnia pulex (Crustacea: Cladocera): observations on male mating behaviour and avoidance of inbreeding. Freshwater Biology, 2002, 47, 441-450.	2.4	44

#	Article	IF	CITATIONS
19	Bacterial cyclic diguanylate signaling networks sense temperature. Nature Communications, 2021, 12, 1986.	12.8	35
20	High-throughput detection of RNA processing in bacteria. BMC Genomics, 2018, 19, 223.	2.8	33
21	Composition and Associations of the Infant Gut Fungal Microbiota with Environmental Factors and Childhood Allergic Outcomes. MBio, 2021, 12, e0339620.	4.1	31
22	Interleukin-4 and IFN-gammaDifferentially Stimulate Macrophage Chemoattractant Protein-1 (MCP-1) and Eotaxin Production by Intestinal Epithelial Cells. Journal of Interferon and Cytokine Research, 2000, 20, 299-308.	1.2	27
23	Monocyte Chemoattractant Protein-1 Production by Intestinal Epithelial CellsIn Vitro:A Role for p38 in Epithelial Chemokine Expression. Journal of Interferon and Cytokine Research, 2001, 21, 223-230.	1.2	19
24	PSORTdb 4.0: expanded and redesigned bacterial and archaeal protein subcellular localization database incorporating new secondary localizations. Nucleic Acids Research, 2021, 49, D803-D808.	14 . 5	18
25	Enabling genomic island prediction and comparison in multiple genomes to investigate bacterial evolution and outbreaks. Microbial Genomics, 2022, 8, .	2.0	10
26	The Pseudomonas aeruginosa whole genome sequence: A 20th anniversary celebration. Advances in Microbial Physiology, 2021, 79, 25-88.	2.4	7
27	AB569, a nontoxic chemical tandem that kills major human pathogenic bacteria. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 4921-4930.	7.1	6
28	Mining the Pseudomonas Genome. Methods in Molecular Biology, 2014, 1149, 417-432.	0.9	4
29	Transcriptome comparison of dengue-susceptible and -resistant field derived strains of Colombian Aedes aegypti using RNA-sequencing. Memorias Do Instituto Oswaldo Cruz, 2021, 116, e200547.	1.6	2