Daniel Hurley

List of Publications by Year
in descending order

22 papers	600 citations	687363 13 h-index	642732 23 g-index
25 all docs	25 docs citations	25 times ranked	1039 citing authors

#	Article	IF	Citations
1	SalmonellaÃ¢â‚¬â€œHost Interactions Ã¢â‚¬â€œ Modulation of the Host Innate Immune System. Frontiers in Immunology, 2014, 5, 481.	4.8	124
2	A Model System for Studying the Transcriptomic and Physiological Changes Associated with Mammalian Host-Adaptation by Leptospira interrogans Serovar Copenhageni. PLoS Pathogens, 2014, 10, e1004004.	4.7	101
3	Whole-Genome Sequencing-Based Characterization of 100 Listeria monocytogenes Isolates Collected from Food Processing Environments over a Four-Year Period. MSphere, 2019, 4, .	2.9	82
4	Characterisation of multidrug-resistant Shiga toxin-producing Escherichia coli cultured from pigs in China: co-occurrence of extended-spectrum Î²-lactamase- and mcr-1-encoding genes on plasmids. International Journal of Antimicrobial Agents, 2016, 48, 445-448.	2.5	44
5	Complete genetic analysis of a Salmonella enterica serovar Indiana isolate accompanying four plasmids carrying mcr-1, ESBL and other resistance genes in China. Veterinary Microbiology, 2017, 210, 142-146.	1.9	38
6	Exploring the Genome and Phenotype of Multi-Drug Resistant Klebsiella pneumoniae of Clinical Origin. Frontiers in Microbiology, 2017, 8, 1913.	3.5	35
7	Complete Genome Sequence of Clostridium estertheticum DSM 8809, a Microbe Identified in Spoiled Vacuum Packed Beef. Frontiers in Microbiology, 2016, 7, 1764.	3.5	19
8	Potential of known and short prokaryotic protein motifs as a basis for novel peptide-based antibacterial therapeutics: a computational survey. Frontiers in Microbiology, 2014, 5, 4.	3.5	18
9	A novel disrupted mcr-1 gene and a lysogenized phage P1-like sequence detected from a large conjugative plasmid, cultured from a human atypical enteropathogenic Escherichia coli (aEPEC) recovered in China. Journal of Antimicrobial Chemotherapy, 2017, 72, dkw564.	3.0	18
10	Enriching antimicrobial peptides from milk hydrolysates using pectin/alginate food-gels. Food Chemistry, 2021, 352, 129220.	8.2	18
11	Increased Virulence of Bloodstream Over Peripheral Isolates of P. aeruginosa Identified Through Post-transcriptional Regulation of Virulence Factors. Frontiers in Cellular and Infection Microbiology, 2018, 8, 357.	3.9	16
12	Yersinia hibernica sp. nov., isolated from pig-production environments. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 2023-2027.	1.7	15
13	Molecular characterisation of multi-drug resistant Escherichia coli of bovine origin. Veterinary Microbiology, 2020, 242, 108566.	1.9	10
14	Yersinia canariae sp. nov., isolated from a human yersiniosis case. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2382-2387.	1.7	10
15	Characterization of Carbapenem-Resistant Enterobacteriaceae Cultured From Retail Meat Products, Patients, and Porcine Excrement in China. Frontiers in Microbiology, 2021, 12, 743468.	3.5	8
16	Complete Genome Sequence of Leptospira alstonii Serovar Room22 Strain GWTS #1. Genome Announcements, 2016, 4, .	0.8	7
17	Characterisation of Early Positive mcr-1 Resistance Gene and Plasmidome in Escherichia coli Pathogenic Strains Associated with Variable Phylogroups under Colistin Selection. Antibiotics, 2021, 10, 1041.	3.7	7
18	Atypical Salmonella enterica Serovars in Murine and Human Macrophage Infection Models. Infection and Immunity, 2020, 88, .	2.2	6

#	Article	IF	Citations
19	Comparison of Listeria monocytogenes Isolates across the Island of Ireland. Journal of Food Protection, 2014, 77, 1402-1406.	1.7	4
20	Molecular Characterization of Salmonella Serovars Anatum and Ealing Associated with Two Historical Outbreaks, Linked to Contaminated Powdered Infant Formula. Frontiers in Microbiology, 2016, 7, 1664.	3.5	2
21	Draft genome sequences of Salmonella Oslo isolated from seafood and its laboratory generated auxotrophic mutant. Journal of Genomics, 2020, 8, 7-10.	0.9	2
22	Draft Genome Sequence of Escherichia coli 26R 793, a Plasmid-Free Recipient Strain Commonly Used in Conjugation Assays. Genome Announcements, 2016, 4, .	0.8	1