Michael Stanhope

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Evolutionary Genomic and Bacterial Genome-Wide Association Study of Mycobacterium avium subsp. <i>paratuberculosis</i> and Dairy Cattle Johne's Disease Phenotypes. Applied and Environmental Microbiology, 2021, 87, .	3.1	4
2	Genome Based Phylogeny and Comparative Genomic Analysis of Intra-Mammary Pathogenic Escherichia coli. PLoS ONE, 2015, 10, e0119799.	2.5	32
3	Comparative characterization of the virulence gene clusters (lipooligosaccharide [LOS] and capsular) Tj ETQq1 1 Campylobacter species. Infection, Genetics and Evolution, 2013, 14, 200-213.	0.784314 2.3	rgBT /Overlo 42
4	Outcome of infection of C57BL/6 IL-10â^'/â^' mice with Campylobacter jejuni strains is correlated with genome content of open reading frames up- and down-regulated inÂvivo. Microbial Pathogenesis, 2013, 54, 1-19.	2.9	23
5	Gene Repertoire Evolution of Streptococcus pyogenes Inferred from Phylogenomic Analysis with Streptococcus canis and Streptococcus dysgalactiae. PLoS ONE, 2012, 7, e37607.	2.5	37
6	Comparative genomics and the role of lateral gene transfer in the evolution of bovine adapted Streptococcus agalactiae. Infection, Genetics and Evolution, 2011, 11, 1263-1275.	2.3	99
7	Gene content differences across strains of Streptococcus uberis identified using oligonucleotide microarray comparative genomic hybridization. Infection, Genetics and Evolution, 2009, 9, 179-188.	2.3	23
8	Positive selection in penicillin-binding proteins 1a, 2b, and 2x from Streptococcus pneumoniae and its correlation with amoxicillin resistance development. Infection, Genetics and Evolution, 2008, 8, 331-339.	2.3	34
9	The relative frequency of intraspecific lateral gene transfer of penicillin binding proteins 1a, 2b, and 2x, in amoxicillin resistant Streptococcus pneumoniae. Infection, Genetics and Evolution, 2007, 7, 520-534.	2.3	17