

Vincent P Richards

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

22
papers

684
citations

15
h-index

25
g-index

25
ext. papers

918
ext. citations

4.6
avg, IF

3.88
L-index

#	Paper	IF	Citations
22	Phylogenomics and the dynamic genome evolution of the genus <i>Streptococcus</i> . <i>Genome Biology and Evolution</i> , 2014 , 6, 741-53	3.9	91
21	Microbiomes of Site-Specific Dental Plaques from Children with Different Caries Status. <i>Infection and Immunity</i> , 2017 , 85,	3.7	89
20	Comparative genomics and the role of lateral gene transfer in the evolution of bovine adapted <i>Streptococcus agalactiae</i> . <i>Infection, Genetics and Evolution</i> , 2011 , 11, 1263-75	4.5	83
19	A Highly Arginolytic <i>Streptococcus</i> Species That Potently Antagonizes <i>Streptococcus mutans</i> . <i>Applied and Environmental Microbiology</i> , 2016 , 82, 2187-201	4.8	73
18	Species delineation and evolutionary history of the globally distributed spotted eagle ray (<i>Aetobatus narinari</i>). <i>Journal of Heredity</i> , 2009 , 100, 273-83	2.4	51
17	RNA-Seq Reveals Enhanced Sugar Metabolism in Co-cultured with within Mixed-Species Biofilms. <i>Frontiers in Microbiology</i> , 2017 , 8, 1036	5.7	47
16	Comparative characterization of the virulence gene clusters (lipooligosaccharide [LOS] and capsular polysaccharide [CPS]) for <i>Campylobacter coli</i> , <i>Campylobacter jejuni</i> subsp. <i>jejuni</i> and related <i>Campylobacter</i> species. <i>Infection, Genetics and Evolution</i> , 2013 , 14, 200-13	4.5	31
15	Genome characterization and population genetic structure of the zoonotic pathogen, <i>Streptococcus canis</i> . <i>BMC Microbiology</i> , 2012 , 12, 293	4.5	28
14	Genome based phylogeny and comparative genomic analysis of intra-mammary pathogenic <i>Escherichia coli</i> . <i>PLoS ONE</i> , 2015 , 10, e0119799	3.7	25
13	Shared phylogeographical breaks in a Caribbean coral reef sponge and its invertebrate commensals. <i>Journal of Biogeography</i> , 2016 , 43, 2136-2146	4.1	22
12	Patterns of population structure for inshore bottlenose dolphins along the eastern United States. <i>Journal of Heredity</i> , 2013 , 104, 765-78	2.4	21
11	Population gene introgression and high genome plasticity for the zoonotic pathogen <i>Streptococcus agalactiae</i> . <i>Molecular Biology and Evolution</i> , 2019 ,	8.3	19
10	Characterization of the heart transcriptome of the white shark (<i>Carcharodon carcharias</i>). <i>BMC Genomics</i> , 2013 , 14, 697	4.5	19
9	Species Designations Belie Phenotypic and Genotypic Heterogeneity in Oral <i>Streptococci</i> . <i>MSystems</i> , 2018 , 3,	7.6	19
8	Site-Specific Profiling of the Dental Mycobiome Reveals Strong Taxonomic Shifts during Progression of Early-Childhood Caries. <i>Applied and Environmental Microbiology</i> , 2020 , 86,	4.8	17
7	Resolving Phylogenetic Relationships for <i>Streptococcus mitis</i> and <i>Streptococcus oralis</i> through Core- and Pan-Genome Analyses. <i>Genome Biology and Evolution</i> , 2019 , 11, 1077-1087	3.9	14
6	Patterns of population structure and dispersal in the long-lived <i>Bedwood</i> of the coral reef, the giant barrel sponge (<i>Xestospongia muta</i>). <i>Coral Reefs</i> , 2016 , 35, 1097-1107	4.2	14

5	Phylogenetic signature of lateral exchange of genes for antibiotic production and resistance among bacteria highlights a pattern of global transmission of pathogens between humans and livestock. <i>Molecular Phylogenetics and Evolution</i> , 2018 , 125, 255-264	4.1	6
4	Blastocystis Mitochondrial Genomes Appear to Show Multiple Independent Gains and Losses of Start and Stop Codons. <i>Genome Biology and Evolution</i> , 2016 , 8, 3340-3350	3.9	5
3	Deep mitochondrial lineage divergence among populations of the southern stingray (<i>Hypanus americanus</i> (Hildebrand & Schroeder, 1928)) throughout the Southeastern United States and Caribbean. <i>Marine Biodiversity</i> , 2019 , 49, 1627-1634	1.4	4
2	Draft Genome Sequence of an Atypical Strain of <i>Streptococcus pneumoniae</i> Isolated from a Respiratory Infection. <i>Genome Announcements</i> , 2014 , 2,		3
1	Evolutionary genomic and bacteria GWAS analysis of subsp. and dairy cattle Johne's disease phenotypes. <i>Applied and Environmental Microbiology</i> , 2021 ,	4.8	3