Mark S Strom

List of Publications by Year in Descending Order

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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.

31 2,181 21 31 g-index

31 2,367 4.6 4.57 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
31	Vibrio parahaemolyticus risk assessment in the Pacific Northwest: ita not whata in the water. <i>FEMS Microbiology Ecology</i> , 2019 , 95,	4.3	14
30	Comparative Genomic Analysis of and Six Taxonomic Synonyms: A First Look at the Distribution and Diversity of the Expanded Species. <i>Frontiers in Microbiology</i> , 2018 , 9, 1893	5.7	13
29	Genomic evidence of adaptive evolution in emergent Vibrio parahaemolyticus ecotypes. <i>Elementa</i> , 2016 , 4,	3.6	4
28	Environmental influences on the seasonal distribution of Vibrio parahaemolyticus in the Pacific Northwest of the USA. <i>FEMS Microbiology Ecology</i> , 2015 , 91,	4.3	31
27	In situ strain-level detection and identification of Vibrio parahaemolyticus using surface-enhanced Raman spectroscopy. <i>Analytical Chemistry</i> , 2013 , 85, 2630-7	7.8	36
26	Population structure of clinical and environmental Vibrio parahaemolyticus from the Pacific Northwest coast of the United States. <i>PLoS ONE</i> , 2013 , 8, e55726	3.7	83
25	Ecology of Vibrio parahaemolyticus and Vibrio vulnificus in the coastal and estuarine waters of Louisiana, Maryland, Mississippi, and Washington (United States). <i>Applied and Environmental Microbiology</i> , 2012 , 78, 7249-57	4.8	140
24	Climate change and seafood safety: Human health implications. <i>Food Research International</i> , 2010 , 43, 1766-1779	7	83
23	Comparative evolutionary analysis of the major structural subunit of Vibrio vulnificus type IV pili. <i>Molecular Biology and Evolution</i> , 2009 , 26, 2185-96	8.3	10
22	The coastal environment and human health: microbial indicators, pathogens, sentinels and reservoirs. <i>Environmental Health</i> , 2008 , 7 Suppl 2, S3	6	125
21	Genome sequence of the fish pathogen Renibacterium salmoninarum suggests reductive evolution away from an environmental Arthrobacter ancestor. <i>Journal of Bacteriology</i> , 2008 , 190, 6970-82	3.5	49
20	Role of type IV pilins in persistence of Vibrio vulnificus in Crassostrea virginica oysters. <i>Applied and Environmental Microbiology</i> , 2007 , 73, 5041-4	4.8	43
19	Sortase inhibitor phenyl vinyl sulfone inhibits Renibacterium salmoninarum adherence and invasion of host cells. <i>Diseases of Aquatic Organisms</i> , 2007 , 78, 115-27	1.7	15
18	A real-time PCR assay for the rapid determination of 16S rRNA genotype in Vibrio vulnificus. <i>Journal of Microbiological Methods</i> , 2007 , 68, 376-84	2.8	56
17	A Vibrio vulnificus type IV pilin contributes to biofilm formation, adherence to epithelial cells, and virulence. <i>Infection and Immunity</i> , 2005 , 73, 1411-22	3.7	122
16	Efficacy of cellular vaccines and genetic adjuvants against bacterial kidney disease in chinook salmon (Oncorhynchus tshawytscha). <i>Fish and Shellfish Immunology</i> , 2004 , 16, 461-74	4.3	31
15	Sequence polymorphism of the 16S rRNA gene of Vibrio vulnificus is a possible indicator of strain virulence. <i>Journal of Clinical Microbiology</i> , 2003 , 41, 442-6	9.7	135

LIST OF PUBLICATIONS

14	Detection and identification of bacterial pathogens of fish in kidney tissue using terminal restriction fragment length polymorphism (T-RFLP) analysis of 16S rRNA genes. <i>Diseases of Aquatic Organisms</i> , 2002 , 48, 175-85	1.7	52
13	Expression of duplicate msa genes in the salmonid pathogen Renibacterium salmoninarum. <i>Applied and Environmental Microbiology</i> , 2002 , 68, 5480-7	4.8	10
12	Type IV Prepilin Leader Peptidases. <i>The Enzymes</i> , 2002 , 22, 127-159	2.3	2
11	An Aeromonas salmonicida type IV pilin is required for virulence in rainbow trout Oncorhynchus mykiss. <i>Diseases of Aquatic Organisms</i> , 2002 , 51, 13-25	1.7	33
10	Epidemiology and pathogenesis of Vibrio vulnificus. <i>Microbes and Infection</i> , 2000 , 2, 177-88	9.3	352
9	Investigation of the role of type IV Aeromonas pilus (Tap) in the pathogenesis of Aeromonas gastrointestinal infection. <i>Infection and Immunity</i> , 2000 , 68, 4040-8	3.7	36
8	Description and characterization of IS994, a putative IS3 family insertion sequence from the salmon pathogen, Renibacterium salmoninarum. <i>Gene</i> , 2000 , 244, 97-107	3.8	17
7	The type IV leader peptidase/N-methyltransferase of Vibrio vulnificus controls factors required for adherence to HEp-2 cells and virulence in iron-overloaded mice. <i>Infection and Immunity</i> , 1998 , 66, 5659-	68 ⁷	81
6	Structure-function relationship of type-IV prepilin peptidase of Pseudomonas aeruginosaa review. <i>Gene</i> , 1997 , 192, 117-21	3.8	79
5	Cloning of an Aeromonas hydrophila type IV pilus biogenesis gene cluster: complementation of pilus assembly functions and characterization of a type IV leader peptidase/N-methyltransferase required for extracellular protein secretion. <i>Molecular Microbiology</i> , 1996 , 19, 857-69	4.1	60
4	Posttranslational processing of type IV prepilin and homologs by PilD of Pseudomonas aeruginosa. <i>Methods in Enzymology</i> , 1994 , 235, 527-40	1.7	43
3	Structure-function and biogenesis of the type IV pili. <i>Annual Review of Microbiology</i> , 1993 , 47, 565-96	17.5	414
2	The bacterial flora of the forehead and back of Alaskan native villagers in summer and in winter. Journal of Investigative Dermatology, 1984 , 82, 294-7	4.3	7
1	Eight year persistence of individual differences in the bacterial flora of the forehead. <i>Journal of Investigative Dermatology</i> , 1982 , 79, 51-2	4.3	5