

Mark S McClain

List of Publications by Year in descending order

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58
papers

2,648
citations

172386

29
h-index

182361

51
g-index

58
all docs

58
docs citations

58
times ranked

2129
citing authors

#	ARTICLE	IF	CITATIONS
1	Delineation of the pH-Responsive Regulon Controlled by the <i>Helicobacter pylori</i> ArsRS Two-Component System. <i>Infection and Immunity</i> , 2021, 89, .	1.0	17
2	Functional Properties of Oligomeric and Monomeric Forms of <i>Helicobacter pylori</i> VacA Toxin. <i>Infection and Immunity</i> , 2021, 89, e0034821.	1.0	5
3	Enhanced Fitness of a <i>Helicobacter pylori</i> babA Mutant in a Murine Model. <i>Infection and Immunity</i> , 2021, 89, e0072520.	1.0	3
4	Inertial-based Fluidic Platform for Rapid Isolation of Blood-borne Pathogens. <i>Military Medicine</i> , 2021, 186, 129-136.	0.4	1
5	Lipoprotein Processing and Sorting in <i>Helicobacter pylori</i> . <i>MBio</i> , 2020, 11, .	1.8	15
6	Temporal Control of the <i>Helicobacter pylori</i> Cag Type IV Secretion System in a Mongolian Gerbil Model of Gastric Carcinogenesis. <i>MBio</i> , 2020, 11, .	1.8	15
7	Bacterial Energetic Requirements for <i>Helicobacter pylori</i> Cag Type IV Secretion System-Dependent Alterations in Gastric Epithelial Cells. <i>Infection and Immunity</i> , 2020, 88, .	1.0	22
8	Functional Properties of <i>Helicobacter pylori</i> VacA Toxin m1 and m2 Variants. <i>Infection and Immunity</i> , 2020, 88, .	1.0	9
9	Effect of environmental salt concentration on the <i>Helicobacter pylori</i> exoproteome. <i>Journal of Proteomics</i> , 2019, 202, 103374.	1.2	14
10	Role of a Stem-Loop Structure in <i>Helicobacter pylori</i> cagA Transcript Stability. <i>Infection and Immunity</i> , 2019, 87, .	1.0	8
11	Magnetic Extraction of <i>Acinetobacter baumannii</i> Using Colistin-Functionalized $\text{Fe}_3\text{O}_4/\text{Au}$ Core/Shell Composite Nanoclusters. <i>ACS Applied Materials & Interfaces</i> , 2017, 9, 26719-26730.	4.0	10
12	<i>Helicobacter pylori</i> Vacuolating Toxin and Gastric Cancer. <i>Toxins</i> , 2017, 9, 316.	1.5	101
13	Genetic signatures for <i>Helicobacter pylori</i> strains of West African origin. <i>PLoS ONE</i> , 2017, 12, e0188804.	1.1	2
14	An Overview of <i>Helicobacter pylori</i> VacA Toxin Biology. <i>Toxins</i> , 2016, 8, 173.	1.5	155
15	Colistin-Functionalized Nanoparticles for the Rapid Capture of <i>Acinetobacter baumannii</i> . <i>Journal of Biomedical Nanotechnology</i> , 2016, 12, 1806-1819.	0.5	15
16	A Nonoligomerizing Mutant Form of <i>Helicobacter pylori</i> VacA Allows Structural Analysis of the p33 Domain. <i>Infection and Immunity</i> , 2016, 84, 2662-2670.	1.0	19
17	Dynamic Computational Model of Symptomatic Bacteremia to Inform Bacterial Separation Treatment Requirements. <i>PLoS ONE</i> , 2016, 11, e0163167.	1.1	7
18	The Myelin and Lymphocyte Protein MAL Is Required for Binding and Activity of <i>Clostridium perfringens</i> α -Toxin. <i>PLoS Pathogens</i> , 2015, 11, e1004896.	2.1	69

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19	Role of Connexin 43 in <i>Helicobacter pylori</i> VacA-Induced Cell Death. <i>Infection and Immunity</i> , 2014, 82, 423-432.	1.0	37
20	Control of gene expression in <i>Helicobacter pylori</i> using the Tet repressor. <i>Journal of Microbiological Methods</i> , 2013, 95, 336-341.	0.7	13
21	Genome Sequences of Three hpAfrica2 Strains of <i>Helicobacter pylori</i> . <i>Genome Announcements</i> , 2013, 1, .	0.8	11
22	Comparative Genomic Analysis of East Asian and Non-Asian <i>Helicobacter pylori</i> Strains Identifies Rapidly Evolving Genes. <i>PLoS ONE</i> , 2013, 8, e55120.	1.1	27
23	Identification of Amino Acids Important for Binding of <i>Clostridium perfringens</i> Epsilon Toxin to Host Cells and to HAVCR1. <i>Biochemistry</i> , 2012, 51, 7588-7595.	1.2	45
24	The Intermediate Region of <i>Helicobacter pylori</i> VacA Is a Determinant of Toxin Potency in a Jurkat T Cell Assay. <i>Infection and Immunity</i> , 2012, 80, 2578-2588.	1.0	33
25	Coenzyme depletion by members of the aerolysin family of pore-forming toxins leads to diminished ATP levels and cell death. <i>Molecular BioSystems</i> , 2012, 8, 2097.	2.9	18
26	Oligomerization of <i>Clostridium perfringens</i> Epsilon Toxin Is Dependent upon Caveolins 1 and 2. <i>PLoS ONE</i> , 2012, 7, e46866.	1.1	42
27	Analysis of <i>cagA</i> in <i>Helicobacter pylori</i> Strains from Colombian Populations with Contrasting Gastric Cancer Risk Reveals a Biomarker for Disease Severity. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 2237-2249.	1.1	46
28	<i>Helicobacter pylori</i> VacA Induces Programmed Necrosis in Gastric Epithelial Cells. <i>Infection and Immunity</i> , 2011, 79, 2535-2543.	1.0	99
29	<i>Helicobacter pylori</i> Exploits a Unique Repertoire of Type IV Secretion System Components for Pilus Assembly at the Bacteria-Host Cell Interface. <i>PLoS Pathogens</i> , 2011, 7, e1002237.	2.1	144
30	Gene-Trap Mutagenesis Identifies Mammalian Genes Contributing to Intoxication by <i>Clostridium perfringens</i> $\hat{\mu}$ -Toxin. <i>PLoS ONE</i> , 2011, 6, e17787.	1.1	62
31	Identification of Small Molecule Inhibitors of <i>Clostridium perfringens</i> $\hat{\mu}$ -Toxin Cytotoxicity Using a Cell-Based High-Throughput Screen. <i>Toxins</i> , 2010, 2, 1825-1847.	1.5	26
32	Analysis of a $\hat{\mu}$ -helical region in the p55 domain of <i>Helicobacter pylori</i> vacuolating toxin. <i>BMC Microbiology</i> , 2010, 10, 60.	1.3	18
33	Reconstitution of <i>Helicobacter pylori</i> VacA Toxin from Purified Components. <i>Biochemistry</i> , 2010, 49, 5743-5752.	1.2	38
34	Dominant-negative Inhibitors of the <i>Clostridium perfringens</i> $\hat{\mu}$ -Toxin. <i>Journal of Biological Chemistry</i> , 2009, 284, 29446-29453.	1.6	30
35	Reconstruction and functional analysis of altered molecular pathways in human atherosclerotic arteries. <i>BMC Genomics</i> , 2009, 10, 13.	1.2	80
36	Genome sequence analysis of <i>Helicobacter pylori</i> strains associated with gastric ulceration and gastric cancer. <i>BMC Genomics</i> , 2009, 10, 3.	1.2	106

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37	<i>Helicobacter pylori</i> HopQ outer membrane protein attenuates bacterial adherence to gastric epithelial cells. FEMS Microbiology Letters, 2008, 289, 53-58.	0.7	41
38	<i>Helicobacter pylori</i> VacA Subdomain Required for Intracellular Toxin Activity and Assembly of Functional Oligomeric Complexes. Infection and Immunity, 2008, 76, 2843-2851.	1.0	42
39	Crystal structure of the <i>Helicobacter pylori</i> vacuolating toxin p55 domain. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 16293-16298.	3.3	143
40	Functional Analysis of Neutralizing Antibodies against Clostridium perfringens Epsilon-Toxin. Infection and Immunity, 2007, 75, 1785-1793.	1.0	37
41	Random Mutagenesis of Helicobacter pylori vacA To Identify Amino Acids Essential for Vacuolating Cytotoxic Activity. Infection and Immunity, 2006, 74, 6188-6195.	1.0	11
42	Mapping of a Domain Required for Protein-Protein Interactions and Inhibitory Activity of a Helicobacter pylori Dominant-Negative VacA Mutant Protein. Infection and Immunity, 2006, 74, 2093-2101.	1.0	15
43	Protein-Protein Interactions among Helicobacter pylori Cag Proteins. Journal of Bacteriology, 2006, 188, 4787-4800.	1.0	63
44	Helicobacter pylori vacuolating toxin. , 2006, , 468-490.		2
45	Functional Properties of the p33 and p55 Domains of the Helicobacter pylori Vacuolating Cytotoxin. Journal of Biological Chemistry, 2005, 280, 21107-21114.	1.6	68
46	Interactions between p-33 and p-55 Domains of the Helicobacter pylori Vacuolating Cytotoxin (VacA). Journal of Biological Chemistry, 2004, 279, 2324-2331.	1.6	49
47	Essential Role of a GXXXG Motif for Membrane Channel Formation by Helicobacter pylori Vacuolating Toxin. Journal of Biological Chemistry, 2003, 278, 12101-12108.	1.6	144
48	Expression of Helicobacter pylori Vacuolating Toxin in Escherichia coli. Infection and Immunity, 2003, 71, 2266-2271.	1.0	16
49	Association of Helicobacter pylori Vacuolating Toxin (VacA) with Lipid Rafts. Journal of Biological Chemistry, 2002, 277, 34642-34650.	1.6	134
50	Amino-Terminal Hydrophobic Region of Helicobacter pylori Vacuolating Cytotoxin (VacA) Mediates Transmembrane Protein Dimerization. Infection and Immunity, 2001, 69, 1181-1184.	1.0	53
51	Antigenic Diversity among Helicobacter pylori Vacuolating Toxins. Infection and Immunity, 2001, 69, 4329-4336.	1.0	18
52	A 12-Amino-Acid Segment, Present in Type s2 but Not Type s1 Helicobacter pylori VacA Proteins, Abolishes Cytotoxin Activity and Alters Membrane Channel Formation. Journal of Bacteriology, 2001, 183, 6499-6508.	1.0	110
53	Acid activation of Helicobacter pylori vacuolating cytotoxin (VacA) results in toxin internalization by eukaryotic cells. Molecular Microbiology, 2000, 37, 433-442.	1.2	87
54	A Dominant Negative Mutant of Helicobacter pylori Vacuolating Toxin (VacA) Inhibits VacA-induced Cell Vacuolation. Journal of Biological Chemistry, 1999, 274, 37736-37742.	1.6	123

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55	Kinetics and Mechanisms of Extracellular Protein Release by <i>Helicobacter pylori</i> . Infection and Immunity, 1999, 67, 5247-5252.	1.0	31
56	Extracellular Release of Antigenic Proteins by <i>Helicobacter pylori</i> . Infection and Immunity, 1998, 66, 2984-2986.	1.0	88
57	Construction of an alkaline phosphatase fusion-generating transposon, mTn10phoA. Gene, 1996, 170, 147-148.	1.0	8
58	Positive Selection of Mutations in the <i>Helicobacter pylori</i> <i>katA</i> 5' Untranslated Region in a Mongolian Gerbil Model of Gastric Disease. Infection and Immunity, 0, , .	1.0	3