Michael F Minnick

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Sequestration and Scavenging of Iron in Infection. Infection and Immunity, 2013, 81, 3503-3514.	1.0	213
2	Characterization of a two-gene locus from Bartonella bacilliformis associated with the ability to invade human erythrocytes. Infection and Immunity, 1995, 63, 1552-1562.	1.0	104
3	Characterization of Bartonella bacilliformis flagella and effect of antiflagellin antibodies on invasion of human erythrocytes. Infection and Immunity, 1993, 61, 4962-4971.	1.0	99
4	ThelalAInvasion Gene ofBartonella bacilliformisEncodes a (Di)Nucleoside Polyphosphate Hydrolase of the MutT Motif Family and Has Homologs in Other Invasive Bacteria. Biochemical and Biophysical Research Communications, 1999, 256, 474-479.	1.0	76
5	Oroya Fever and Verruga Peruana: Bartonelloses Unique to South America. PLoS Neglected Tropical Diseases, 2014, 8, e2919.	1.3	75
6	A bacterial-induced lectin which triggers hemocyte coagulation in Manducasexta. Biochemical and Biophysical Research Communications, 1986, 137, 729-735.	1.0	63
7	A highly repetitive DNA sequence possibly unique to canids. Gene, 1992, 110, 235-238.	1.0	61
8	Pestilence, persistence and pathogenicity: infection strategies of <i>Bartonella</i> . Future Microbiology, 2009, 4, 743-758.	1.0	61
9	Mitogenic Effect of Bartonella bacilliformis on Human Vascular Endothelial Cells and Involvement of GroEL. Infection and Immunity, 2003, 71, 6933-6942.	1.0	59
10	Hemin-Binding Surface Protein from Bartonella quintana. Infection and Immunity, 2000, 68, 6750-6757.	1.0	57
11	Establishing a Direct Role for the Bartonella bacilliformis Invasion-Associated Locus B (IalB) Protein in Human Erythrocyte Parasitism. Infection and Immunity, 2001, 69, 4373-4381.	1.0	56
12	Experimental Model of Human Body Louse Infection Using Green Fluorescent Protein-Expressing Bartonella quintana. Infection and Immunity, 2001, 69, 1876-1879.	1.0	55
13	Five-Member Gene Family of Bartonella quintana. Infection and Immunity, 2003, 71, 814-821.	1.0	52
14	Proteomic and Immunoblot Analyses of Bartonella quintana Total Membrane Proteins Identify Antigens Recognized by Sera from Infected Patients. Infection and Immunity, 2007, 75, 2548-2561.	1.0	52
15	Strategy for Detection and Differentiation of Coxiella bumetii Strains Using the Polymerase Chain Reactiona. Annals of the New York Academy of Sciences, 1990, 590, 572-581.	1.8	42
16	Cell entry and the pathogenesis of Bartonella infections. Trends in Microbiology, 1996, 4, 343-347.	3.5	41
17	A bacteriophage-like particle from Bartonella bacilliformis. Microbiology (United Kingdom), 2000, 146, 599-609.	0.7	37
18	Identification of outer membrane proteins of Bartonella bacilliformis. Infection and Immunity, 1994, 62, 2644-2648.	1.0	37

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19	Development of a System for Genetic Manipulation of <i>Bartonella bacilliformis</i> . Applied and Environmental Microbiology, 1999, 65, 3441-3448.	1.4	36
20	Environmental Signals Generate a Differential and Coordinated Expression of the Heme Receptor Gene Family of Bartonella quintana. Infection and Immunity, 2006, 74, 3251-3261.	1.0	35
21	Function, Regulation, and Transcriptional Organization of the Hemin Utilization Locus of <i>Bartonella quintana</i> . Infection and Immunity, 2009, 77, 307-316.	1.0	34
22	Colonization of Lutzomyia verrucarum and Lutzomyia longipalpis Sand Flies (Diptera: Psychodidae) by Bartonella bacilliformis, the Etiologic Agent of Carrión's Disease. PLoS Neglected Tropical Diseases, 2015, 9, e0004128.	1.3	32
23	Identification of Novel Small RNAs and Characterization of the 6S RNA of Coxiella burnetii. PLoS ONE, 2014, 9, e100147.	1.1	32
24	A carboxy-terminal processing protease gene is located immediately upstream of the invasion-associated locus from Bartonella bacilliformis. Microbiology (United Kingdom), 1997, 143, 1221-1233.	0.7	30
25	Reversion rates in a leuB auxotroph of Escherichia coli K-12 correlate with ppGpp levels during exponential growth. Microbiology (United Kingdom), 1997, 143, 847-854.	0.7	30
26	gyrA Mutations in Ciprofloxacin-Resistant Bartonella bacilliformis Strains Obtained In Vitro. Antimicrobial Agents and Chemotherapy, 2003, 47, 383-386.	1.4	30
27	Mechanisms by which transcription can regulate somatic hypermutation. Genes and Immunity, 2004, 5, 176-182.	2.2	30
28	Developmental Biology of Coxiella burnetii. Advances in Experimental Medicine and Biology, 2012, 984, 231-248.	0.8	30
29	Characterization of the 16S-23S rRNA intergenic spacer of Bartonella bacilliformis. Gene, 1994, 143, 149-150.	1.0	29
30	Toxic Introns and Parasitic Intein in Coxiella burnetii : Legacies of a Promiscuous Past. Journal of Bacteriology, 2008, 190, 5934-5943.	1.0	29
31	Differential expression of the invasion-associated locus B (ialB) gene of Bartonella bacilliformis in response to environmental cues. Microbial Pathogenesis, 2003, 34, 179-186.	1.3	28
32	Transcriptional Regulation of the Heme Binding Protein Gene Family of Bartonella quintana Is Accomplished by a Novel Promoter Element and Iron Response Regulator. Infection and Immunity, 2007, 75, 4373-4385.	1.0	27
33	Group I Introns and Inteins: Disparate Origins but Convergent Parasitic Strategies. Journal of Bacteriology, 2009, 191, 6193-6202.	1.0	27
34	A DNA-Binding Peroxiredoxin of <i>Coxiella burnetii</i> Is Involved in Countering Oxidative Stress during Exponential-Phase Growth. Journal of Bacteriology, 2010, 192, 2077-2084.	1.0	27
35	Identification of Bartonella using PCR; genus- and species-specific primer sets. Journal of Microbiological Methods, 1997, 31, 51-57.	0.7	24
36	A plasmid-encoded surface protein found in chronic-disease isolates of Coxiella burnetti. Infection and Immunity, 1991, 59, 4735-4739.	1.0	23

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37	Analysis of QpRS-Specific Sequences from Coxiella burnetii. Annals of the New York Academy of Sciences, 1990, 590, 514-522.	1.8	22
38	The Unusual 23S rRNA Gene of <i>Coxiella burnetii</i> : Two Self-Splicing Group I Introns Flank a 34-Base-Pair Exon, and One Element Lacks the Canonical ΩG. Journal of Bacteriology, 2007, 189, 6572-6579.	1.0	21
39	II. Correlations between secondary structure stability and mutation frequency during somatic hypermutation. Molecular Immunology, 2008, 45, 3600-3608.	1.0	20
40	Mutation-Driven Divergence and Convergence Indicate Adaptive Evolution of the Intracellular Human-Restricted Pathogen, Bartonella bacilliformis. PLoS Neglected Tropical Diseases, 2016, 10, e0004712.	1.3	19
41	Analysis of the Caenorhabditis elegans innate immune response to Coxiella burnetii. Innate Immunity, 2017, 23, 111-127.	1.1	19
42	Analysis of the cbhE' plasmid gene from acute disease-causing isolates of Coxiella burnetii. Gene, 1991, 103, 113-118.	1.0	18
43	Mutations in <i>Bartonella bacilliformis gyrB</i> Confer Resistance to Coumermycin A ₁ . Antimicrobial Agents and Chemotherapy, 1998, 42, 2906-2913.	1.4	18
44	Genetics of <i>Coxiella burnetii</i> : on the path of specialization. Future Microbiology, 2011, 6, 1297-1314.	1.0	16
45	I. VH gene transcription creates stabilized secondary structures for coordinated mutagenesis during somatic hypermutation. Molecular Immunology, 2008, 45, 3589-3599.	1.0	15
46	Proteins of <i>Bartonella bacilliformis</i> : Candidates for Vaccine Development. International Journal of Peptides, 2015, 2015, 1-5.	0.7	14
47	Identification of novel MITEs (miniature inverted-repeat transposable elements) in Coxiella burnetii: implications for protein and small RNA evolution. BMC Genomics, 2018, 19, 247.	1.2	14
48	A CsrA-Binding, <i>trans</i> -Acting sRNA of <i>Coxiella burnetii</i> Is Necessary for Optimal Intracellular Growth and Vacuole Formation during Early Infection of Host Cells. Journal of Bacteriology, 2019, 201, .	1.0	14
49	Transformation of Bartonella bacilliformis by electroporation. Canadian Journal of Microbiology, 1994, 40, 782-786.	0.8	13
50	Pentamidine inhibits Coxiella burnetii growth and 23S rRNA intron splicing in vitro. International Journal of Antimicrobial Agents, 2010, 36, 380-382.	1.1	13
51	Bartonella bacilliformis GroEL: Effect on Growth of Human Vascular Endothelial Cells in Infected Cocultures. Annals of the New York Academy of Sciences, 2005, 1063, 286-298.	1.8	12
52	Tissue site and modification of a bacteria-induced coagulation protein from Manduca sexta. Insect Biochemistry, 1988, 18, 637-644.	1.8	11
53	Cloning, functional expression, and complementation analysis of an inorganic pyrophosphatase fromBartonella bacilliformis. Canadian Journal of Microbiology, 1997, 43, 734-743.	0.8	11

A unique Coxiella burnetii lipoprotein involved in metal binding (LimB). Microbiology (United) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 To 0.7

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55	The roles of transcription and genotoxins underlying p53 mutagenesis in vivo. Carcinogenesis, 2011, 32, 1559-1567.	1.3	10
56	Nucleotide sequence and comparison of the 5S ribosomal RNA genes ofRochalimaea henselae, R.quintanaandBrucella abortus. Nucleic Acids Research, 1993, 21, 2518-2518.	6.5	9
57	Bartonella Interactions with Host Cells. Sub-Cellular Biochemistry, 2000, 33, 97-123.	1.0	9
58	Nucleotide sequence analysis of the 23S ribosomal RNA-encoding gene of Bartonella bacilliformis. Gene, 1995, 162, 75-79.	1.0	8
59	Human vascular endothelial cells express epithelial growth factor in response to infection by Bartonella bacilliformis. PLoS Neglected Tropical Diseases, 2020, 14, e0008236.	1.3	8
60	Regulation and synthesis of selected bacteria-induced proteins in Manduca sexta. Insect Biochemistry and Molecular Biology, 1992, 22, 321-331.	1.2	7
61	Nucleotide sequence of the 5S ribosomal RNA gene ofBartonella bacilliformis. Nucleic Acids Research, 1993, 21, 1036-1036.	6.5	7
62	Ribozyme Stability, Exon Skipping, and a Potential Role for RNA Helicase in Group I Intron Splicing by Coxiella burnetii. Journal of Bacteriology, 2011, 193, 5292-5299.	1.0	7
63	Kinetic models reveal the in vivo mechanisms of mutagenesis in microbes and man. Mutation Research - Reviews in Mutation Research, 2013, 752, 129-137.	2.4	7
64	Novel small RNAs expressed by Bartonella bacilliformis under multiple conditions reveal potential mechanisms for persistence in the sand fly vector and human host. PLoS Neglected Tropical Diseases, 2020, 14, e0008671.	1.3	7
65	A Unique Group I Intron in <i>Coxiella burnetii</i> Is a Natural Splice Mutant. Journal of Bacteriology, 2009, 191, 4044-4046.	1.0	6
66	Bartonella. , 2015, , 1911-1939.		6
67	The Intervening Sequence of Coxiella burnetii: Characterization and Evolution. Frontiers in Cellular and Infection Microbiology, 2016, 6, 83.	1.8	6
68	Laboratory Maintenance of <i>Bartonella quintana</i> . Current Protocols in Microbiology, 2008, 10, Unit 3C.1.1-3C.1.13.	6.5	6
69	Characterization and expression analysis of the groESL operon of Bartonella bacilliformis. Gene, 2005, 359, 53-62.	1.0	5
70	The Genus Bartonella. , 2006, , 467-492.		5
71	Evolution of coordinated mutagenesis and somatic hypermutation in VH5. Molecular Immunology, 2011, 49, 537-548.	1.0	4
72	Gene duplication and deletion, not horizontal transfer, drove intra-species mosaicism of Bartonella henselae. Genomics, 2020, 112, 467-471.	1.3	4

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73	Cloning and sequence analysis of a hemolysin-encoding gene from Pseudomonas paucimobilis. Gene, 1993, 130, 57-63.	1.0	3
74	Characterization of the fMet initiator tRNA gene of Bartonella bacilliformis. Gene, 1993, 131, 151-152.	1.0	2
75	Bartonelloses. , 2006, , 454-462.		2
76	Bartonella. , 2002, , 2115-2136.		2
77	Coxiella. , 2015, , 1941-1972.		1
78	Virulence Determinants of Bartonella bacilliformis. , 2002, , 197-211.		0

Virulence Determinants of Bartonella bacilliformis. , 2002, , 197-211. 78