

Robert A Heinzen

List of Publications by Year in descending order

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83
papers

12,119
citations

61857

43
h-index

62479

80
g-index

86
all docs

86
docs citations

86
times ranked

18348
citing authors

#	ARTICLE	IF	CITATIONS
1	Guidelines for the use and interpretation of assays for monitoring autophagy (3rd edition). <i>Autophagy</i> , 2016, 12, 1-222.	4.3	4,701
2	Lounging in a lysosome: the intracellular lifestyle of <i>Coxiella burnetii</i> . <i>Cellular Microbiology</i> , 2007, 9, 829-840.	1.1	1,560
3	Complete genome sequence of the Q-fever pathogen <i>Coxiella burnetii</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 5455-5460.	3.3	506
4	Host cell-free growth of the Q fever bacterium <i>Coxiella burnetii</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 4430-4434.	3.3	363
5	Temporal Analysis of <i>Coxiella burnetii</i> Morphological Differentiation. <i>Journal of Bacteriology</i> , 2004, 186, 7344-7352.	1.0	277
6	Dot/Icm Type IVB Secretion System Requirements for <i>Coxiella burnetii</i> Growth in Human Macrophages. <i>MBio</i> , 2011, 2, e00175-11.	1.8	214
7	Comparative Genomics Reveal Extensive Transposon-Mediated Genomic Plasticity and Diversity among Potential Effector Proteins within the Genus <i>Coxiella</i> . <i>Infection and Immunity</i> , 2009, 77, 642-656.	1.0	197
8	Isolation from Animal Tissue and Genetic Transformation of <i>Coxiella burnetii</i> Are Facilitated by an Improved Axenic Growth Medium. <i>Applied and Environmental Microbiology</i> , 2011, 77, 3720-3725.	1.4	191
9	Developmental biology of <i>Coxiella burnetii</i> . <i>Trends in Microbiology</i> , 1999, 7, 149-154.	3.5	181
10	<i>Coxiella burnetii</i> Phase I and II Variants Replicate with Similar Kinetics in Degradative Phagolysosome-Like Compartments of Human Macrophages. <i>Infection and Immunity</i> , 2010, 78, 3465-3474.	1.0	140
11	A <i>Rickettsia</i> WASP-like protein activates the Arp2/3 complex and mediates actin-based motility. <i>Cellular Microbiology</i> , 2004, 6, 761-769.	1.1	137
12	The <i>Coxiella burnetii</i> Ankyrin Repeat Domain-Containing Protein Family Is Heterogeneous, with C-Terminal Truncations That Influence Dot/Icm-Mediated Secretion. <i>Journal of Bacteriology</i> , 2009, 191, 4232-4242.	1.0	137
13	The <i>Coxiella burnetii</i> Cryptic Plasmid Is Enriched in Genes Encoding Type IV Secretion System Substrates. <i>Journal of Bacteriology</i> , 2011, 193, 1493-1503.	1.0	134
14	<i>Coxiella burnetii</i> Inhibits Apoptosis in Human THP-1 Cells and Monkey Primary Alveolar Macrophages. <i>Infection and Immunity</i> , 2007, 75, 4263-4271.	1.0	125
15	Maturation of the <i>Coxiella burnetii</i> parasitophorous vacuole requires bacterial protein synthesis but not replication. <i>Cellular Microbiology</i> , 2003, 5, 469-480.	1.1	122
16	Virulent <i>Coxiella burnetii</i> does not activate human dendritic cells: Role of lipopolysaccharide as a shielding molecule. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 8722-8727.	3.3	122
17	Genetic Diversity of the Q Fever Agent, <i>Coxiella burnetii</i> , Assessed by Microarray-Based Whole-Genome Comparisons. <i>Journal of Bacteriology</i> , 2006, 188, 2309-2324.	1.0	122
18	Dynamics of Actin-Based Movement by <i>Rickettsia rickettsii</i> in Vero Cells. <i>Infection and Immunity</i> , 1999, 67, 4201-4207.	1.0	112

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19	<i>Coxiella burnetii</i> inhabits a cholesterol-rich vacuole and influences cellular cholesterol metabolism. <i>Cellular Microbiology</i> , 2006, 8, 496-507.	1.1	108
20	Ultrastructure of <i>Rickettsia rickettsii</i> Actin Tails and Localization of Cytoskeletal Proteins. <i>Infection and Immunity</i> , 2000, 68, 4706-4713.	1.0	104
21	Serological Evidence of Human Infection with the Protozoan <i>Neospora caninum</i> . <i>Vaccine Journal</i> , 1999, 6, 765-767.	2.6	101
22	Two Systems for Targeted Gene Deletion in <i>Coxiella burnetii</i> . <i>Applied and Environmental Microbiology</i> , 2012, 78, 4580-4589.	1.4	99
23	Characterization of a <i>Coxiella burnetii</i> <i>ftsZ</i> Mutant Generated by <i>Himar1</i> Transposon Mutagenesis. <i>Journal of Bacteriology</i> , 2009, 191, 1369-1381.	1.0	94
24	Candidate Antigens for Q Fever Serodiagnosis Revealed by Immunoscreening of a <i>Coxiella burnetii</i> Protein Microarray. <i>Vaccine Journal</i> , 2008, 15, 1771-1779.	3.2	92
25	Sustained Activation of Akt and Erk1/2 Is Required for <i>Coxiella burnetii</i> Antiapoptotic Activity. <i>Infection and Immunity</i> , 2009, 77, 205-213.	1.0	88
26	<i>Coxiella burnetii</i> effector protein subverts clathrin-mediated vesicular trafficking for pathogen vacuole biogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E4770-9.	3.3	85
27	Right on Q: genetics begin to unravel <i>Coxiella burnetii</i> host cell interactions. <i>Future Microbiology</i> , 2016, 11, 919-939.	1.0	84
28	Proteome and Antigen Profiling of <i>Coxiella burnetii</i> Developmental Forms. <i>Infection and Immunity</i> , 2007, 75, 290-298.	1.0	80
29	Advances in Genetic Manipulation of Obligate Intracellular Bacterial Pathogens. <i>Frontiers in Microbiology</i> , 2011, 2, 97.	1.5	79
30	<i>Coxiella burnetii</i> Effector Proteins That Localize to the Parasitophorous Vacuole Membrane Promote Intracellular Replication. <i>Infection and Immunity</i> , 2015, 83, 661-670.	1.0	79
31	Rapid Typing of <i>Coxiella burnetii</i> . <i>PLoS ONE</i> , 2011, 6, e26201.	1.1	76
32	Sustained Axenic Metabolic Activity by the Obligate Intracellular Bacterium <i>Coxiella burnetii</i> . <i>Journal of Bacteriology</i> , 2008, 190, 3203-3212.	1.0	71
33	Nitric Oxide Inhibits <i>Coxiella burnetii</i> Replication and Parasitophorous Vacuole Maturation. <i>Infection and Immunity</i> , 2002, 70, 5140-5147.	1.0	69
34	<i>Coxiella</i> type IV secretion and cellular microbiology. <i>Current Opinion in Microbiology</i> , 2009, 12, 74-80.	2.3	66
35	Adaptive immunity to the obligate intracellular pathogen <i>Coxiella burnetii</i> . <i>Immunologic Research</i> , 2009, 43, 138-148.	1.3	65
36	Complementation of Arginine Auxotrophy for Genetic Transformation of <i>Coxiella burnetii</i> by Use of a Defined Axenic Medium. <i>Applied and Environmental Microbiology</i> , 2016, 82, 3042-3051.	1.4	64

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37	Genetic mechanisms of <i>Coxiella burnetii</i> lipopolysaccharide phase variation. <i>PLoS Pathogens</i> , 2018, 14, e1006922.	2.1	60
38	Rickettsial Actin-Based Motility. <i>Annals of the New York Academy of Sciences</i> , 2003, 990, 535-547.	1.8	58
39	Fusogenicity of the <i>Coxiella burnetii</i> Parasitophorous Vacuole. <i>Annals of the New York Academy of Sciences</i> , 2003, 990, 556-562.	1.8	57
40	Specificity of <i>Legionella pneumophila</i> and <i>Coxiella burnetii</i> Vacuoles and Versatility of <i>Legionella pneumophila</i> Revealed by Coinfection. <i>Infection and Immunity</i> , 2005, 73, 4494-4504.	1.0	55
41	Bacterial Colonization of Host Cells in the Absence of Cholesterol. <i>PLoS Pathogens</i> , 2013, 9, e1003107.	2.1	55
42	Comparative DNA Microarray Analysis of Host Cell Transcriptional Responses to Infection by <i>Coxiella burnetii</i> or <i>Chlamydia trachomatis</i> . <i>Annals of the New York Academy of Sciences</i> , 2003, 990, 701-713.	1.8	54
43	Life on the Outside: The Rescue of <i>Coxiella burnetii</i> from Its Host Cell. <i>Annual Review of Microbiology</i> , 2011, 65, 111-128.	2.9	52
44	Transcriptional Profiling of <i>Coxiella burnetii</i> Reveals Extensive Cell Wall Remodeling in the Small Cell Variant Developmental Form. <i>PLoS ONE</i> , 2016, 11, e0149957.	1.1	50
45	β -Barrel proteins tether the outer membrane in many Gram-negative bacteria. <i>Nature Microbiology</i> , 2021, 6, 19-26.	5.9	46
46	Elevated Cholesterol in the <i>Coxiella burnetii</i> Intracellular Niche Is Bacteriolytic. <i>MBio</i> , 2017, 8, .	1.8	44
47	Essential Role for the Response Regulator PmrA in <i>Coxiella burnetii</i> Type 4B Secretion and Colonization of Mammalian Host Cells. <i>Journal of Bacteriology</i> , 2014, 196, 1925-1940.	1.0	43
48	Developmental transitions of <i>Coxiella burnetii</i> grown in axenic media. <i>Journal of Microbiological Methods</i> , 2014, 96, 104-110.	0.7	43
49	Comparative virulence of diverse <i>Coxiella burnetii</i> strains. <i>Virulence</i> , 2019, 10, 133-150.	1.8	41
50	A method for purifying obligate intracellular <i>Coxiella burnetii</i> that employs digitonin lysis of host cells. <i>Journal of Microbiological Methods</i> , 2008, 72, 321-325.	0.7	40
51	Interactions between the <i>Coxiella burnetii</i> parasitophorous vacuole and the endoplasmic reticulum involve the host protein ORP1L. <i>Cellular Microbiology</i> , 2017, 19, e12637.	1.1	38
52	The <i>Coxiella burnetii</i> Parasitophorous Vacuole. <i>Advances in Experimental Medicine and Biology</i> , 2012, 984, 141-169.	0.8	37
53	Antibody-mediated immunity to the obligate intracellular bacterial pathogen <i>Coxiella burnetii</i> is Fc receptor- and complement-independent. <i>BMC Immunology</i> , 2009, 10, 26.	0.9	34
54	<i>Coxiella burnetii</i> Expresses a Functional Δ^24 Sterol Reductase. <i>Journal of Bacteriology</i> , 2010, 192, 6154-6159.	1.0	34

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55	<i>Coxiella burnetii</i> RpoS Regulates Genes Involved in Morphological Differentiation and Intracellular Growth. <i>Journal of Bacteriology</i> , 2019, 201, .	1.0	33
56	Gene Inactivation in <i>Coxiella burnetii</i> . <i>Methods in Molecular Biology</i> , 2014, 1197, 329-345.	0.4	30
57	Bringing Culture to the Uncultured: <i>Coxiella burnetii</i> and Lessons for Obligate Intracellular Bacterial Pathogens. <i>PLoS Pathogens</i> , 2013, 9, e1003540.	2.1	28
58	Host-microbe interaction systems biology: lifecycle transcriptomics and comparative genomics. <i>Future Microbiology</i> , 2010, 5, 205-219.	1.0	27
59	Infection of Human Monocyte-Derived Macrophages With <i>Coxiella burnetii</i> . , 2008, 431, 189-200.		25
60	Sec-mediated secretion by <i>Coxiella burnetii</i> . <i>BMC Microbiology</i> , 2013, 13, 222.	1.3	25
61	Whole-Genome Sequence of <i>Coxiella burnetii</i> Nine Mile RSA439 (Phase II, Clone 4), a Laboratory Workhorse Strain. <i>Genome Announcements</i> , 2017, 5, .	0.8	24
62	High-Content Imaging Reveals Expansion of the Endosomal Compartment during <i>Coxiella burnetii</i> Parasitophorous Vacuole Maturation. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 48.	1.8	23
63	Contributions of lipopolysaccharide and the type IVB secretion system to <i>Coxiella burnetii</i> vaccine efficacy and reactogenicity. <i>Npj Vaccines</i> , 2021, 6, 38.	2.9	22
64	Efficient Method of Cloning the Obligate Intracellular Bacterium <i>Coxiella burnetii</i> . <i>Applied and Environmental Microbiology</i> , 2007, 73, 4048-4054.	1.4	20
65	Noncanonical Inhibition of mTORC1 by <i>Coxiella burnetii</i> Promotes Replication within a Phagolysosome-Like Vacuole. <i>MBio</i> , 2019, 10, .	1.8	20
66	Lack of Dendritic Cell Maturation Following Infection by <i>Coxiella burnetii</i> Synthesizing Different Lipopolysaccharide Chemotypes. <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 154-160.	1.8	19
67	Replication of <i>Coxiella burnetii</i> Is Inhibited in CHO K-1 Cells Treated with Inhibitors of Cholesterol Metabolism. <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 123-129.	1.8	16
68	Actin polymerization in the endosomal pathway, but not on the <i>Coxiella</i> -containing vacuole, is essential for pathogen growth. <i>PLoS Pathogens</i> , 2018, 14, e1007005.	2.1	16
69	Robust growth of avirulent phase II <i>Coxiella burnetii</i> in bone marrow-derived murine macrophages. <i>PLoS ONE</i> , 2017, 12, e0173528.	1.1	14
70	A <i>Coxiella burnetii</i> phospholipase A homolog <i>pldA</i> is required for optimal growth in macrophages and developmental form lipid remodeling. <i>BMC Microbiology</i> , 2018, 18, 33.	1.3	13
71	Fractionation of the <i>Coxiella burnetii</i> Parasitophorous Vacuole. <i>Methods in Molecular Biology</i> , 2008, 445, 389-406.	0.4	12
72	Vasodilator-Stimulated Phosphoprotein Activity Is Required for <i>Coxiella burnetii</i> Growth in Human Macrophages. <i>PLoS Pathogens</i> , 2016, 12, e1005915.	2.1	11

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73	Dependency of <i>Coxiella burnetii</i> Type 4B Secretion on the Chaperone IcmS. <i>Journal of Bacteriology</i> , 2019, 201, .	1.0	9
74	Replication of <i>Coxiella burnetii</i> in a Lysosome-Like Vacuole Does Not Require Lysosomal Hydrolases. <i>Infection and Immunity</i> , 2019, 87, .	1.0	8
75	Intracellular Development of <i>Coxiella burnetii</i> . , 2002, , 99-129.		7
76	Draft Genome Sequences of the Avirulent <i>Coxiella burnetii</i> Dugway 7D77-80 and Dugway 7E65-68 Strains Isolated from Rodents in Dugway, Utah. <i>Genome Announcements</i> , 2017, 5, .	0.8	7
77	Preliminary Assessment of Genome Differences between the Reference Nine Mile Isolate and Two Human Endocarditis Isolates of <i>Coxiella burnetii</i> . <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 64-67.	1.8	6
78	<i>Coxiella burnetii</i> Sterol-Modifying Protein Stmp1 Regulates Cholesterol in the Intracellular Niche. <i>MBio</i> , 2022, 13, e0307321.	1.8	6
79	Draft Genome Sequences of Historical Strains of <i>Coxiella burnetii</i> Isolated from Cow's Milk and a Goat Placenta. <i>Genome Announcements</i> , 2017, 5, .	0.8	4
80	A Comprehensive Phenotypic Screening Strategy to Identify Modulators of Cargo Translocation by the Bacterial Type IVB Secretion System. <i>MBio</i> , 2022, 13, e0024022.	1.8	3
81	Draft Genome Sequences of Three <i>Coxiella burnetii</i> Strains Isolated from Q Fever Patients. <i>Genome Announcements</i> , 2017, 5, .	0.8	2
82	Murine Q Fever Vaccination Model Reveals Sex Dimorphism in Early Phase Delayed-Type Hypersensitivity Responses. <i>Frontiers in Immunology</i> , 0, 13, .	2.2	1
83	Exploring the Cause of Human Q Fever: Recent Advances in <i>Coxiella burnetii</i> Research. , 2010, , 75-85.		0