

Jere W McBride

List of Publications by Year in descending order

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96
papers

4,620
citations

87843

38
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114418

63
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99
all docs

99
docs citations

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times ranked

2564
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Ehrlichia</i> SLiM Ligand Mimetic Activates Notch Signaling in Human Monocytes. <i>MBio</i> , 2022, 13, e0007622.	1.8	11
2	<i>Ehrlichia</i> SLiM ligand mimetic activates Hedgehog signaling to engage a BCL-2 anti-apoptotic cellular program. <i>PLoS Pathogens</i> , 2022, 18, e1010345.	2.1	10
3	<i>Ehrlichia chaffeensis</i> TRP120 Is a Wnt Ligand Mimetic That Interacts with Wnt Receptors and Contains a Novel Repetitive Short Linear Motif That Activates Wnt Signaling. <i>MSphere</i> , 2021, 6, .	1.3	15
4	Anaplasmataceae: Dichotomous Autophagic Interplay for Infection. <i>Frontiers in Immunology</i> , 2021, 12, 642771.	2.2	7
5	<i>Ehrlichia</i> TRP effectors: moonlighting, mimicry and infection. <i>Pathogens and Disease</i> , 2021, 79, .	0.8	22
6	Seroprevalence and Genotypic Analysis of <i>Ehrlichia canis</i> Infection in Dogs and Humans in Cauca, Colombia. <i>American Journal of Tropical Medicine and Hygiene</i> , 2021, 104, 1771-1776.	0.6	5
7	Serological evidence of <i>Ehrlichia minasensis</i> infection in Brazilian dogs. <i>Acta Tropica</i> , 2021, 219, 105931.	0.9	6
8	Alpha Enolase 1 Ubiquitination and Degradation Mediated by <i>Ehrlichia chaffeensis</i> TRP120 Disrupts Glycolytic Flux and Promotes Infection. <i>Pathogens</i> , 2021, 10, 962.	1.2	4
9	Immunoreactive Protein Repertoires of <i>Ehrlichia chaffeensis</i> and <i>E. canis</i> Reveal the Dominance of Hypothetical Proteins and Conformation-Dependent Antibody Epitopes. <i>Infection and Immunity</i> , 2021, 89, e0022421.	1.0	5
10	Editorial: The Autophagy Pathway: Bacterial Pathogen Immunity and Evasion. <i>Frontiers in Immunology</i> , 2021, 12, 768935.	2.2	1
11	<i>Ehrlichia chaffeensis</i> and <i>E. canis</i> hypothetical protein immunoanalysis reveals small secreted immunodominant proteins and conformation-dependent antibody epitopes. <i>Npj Vaccines</i> , 2020, 5, 85.	2.9	12
12	Editorial: Wnt Signaling in Immune Cell Regulation During Microbial Infection and Cancer. <i>Frontiers in Immunology</i> , 2020, 11, 1133.	2.2	1
13	<i>Ehrlichia canis</i> TRP36 diversity in naturally infected-dogs from an urban area of Colombia. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101367.	1.1	17
14	<i>Ehrlichia chaffeensis</i> TRP120-mediated ubiquitination and proteasomal degradation of tumor suppressor FBW7 increases oncoprotein stability and promotes infection. <i>PLoS Pathogens</i> , 2020, 16, e1008541.	2.1	24
15	Activation of ASC Inflammasome Driven by Toll-Like Receptor 4 Contributes to Host Immunity against Rickettsial Infection. <i>Infection and Immunity</i> , 2020, 88, .	1.0	16
16	Bacterial Manipulation of Wnt Signaling: A Host-Pathogen Tug-of-Wnt. <i>Frontiers in Immunology</i> , 2019, 10, 2390.	2.2	39
17	<i>Ehrlichia chaffeensis</i> Outer Membrane Protein 1-Specific Human Antibody-Mediated Immunity Is Defined by Intracellular TRIM21-Dependent Innate Immune Activation and Extracellular Neutralization. <i>Infection and Immunity</i> , 2019, 87, .	1.0	12
18	<i>Ehrlichia chaffeensis</i> TRP120 Effector Targets and Recruits Host Polycomb Group Proteins for Degradation To Promote Intracellular Infection. <i>Infection and Immunity</i> , 2018, 86, .	1.0	30

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19	<i>Ehrlichia chaffeensis</i> TRP47 enters the nucleus via a MYND-binding domain-dependent mechanism and predominantly binds enhancers of host genes associated with signal transduction, cytoskeletal organization, and immune response. <i>PLoS ONE</i> , 2018, 13, e0205983.	1.1	15
20	<i>Ehrlichia chaffeensis</i> TRP120 nucleomodulin binds DNA with disordered tandem repeat domain. <i>PLoS ONE</i> , 2018, 13, e0194891.	1.1	27
21	<i>Ehrlichia chaffeensis</i> TRP75 Interacts with Host Cell Targets Involved in Homeostasis, Cytoskeleton Organization, and Apoptosis Regulation To Promote Infection. <i>MSphere</i> , 2018, 3, .	1.3	18
22	<i>Ehrlichia chaffeensis</i> TRP32 Nucleomodulin Function and Localization Is Regulated by NEDD4L-Mediated Ubiquitination. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 7, 534.	1.8	20
23	Tick-Borne Emerging Infections. <i>Clinics in Laboratory Medicine</i> , 2017, 37, 317-340.	0.7	147
24	Engineering of obligate intracellular bacteria: progress, challenges and paradigms. <i>Nature Reviews Microbiology</i> , 2017, 15, 544-558.	13.6	144
25	<i>Ehrlichia chaffeensis</i> TRP120 Moonlights as a HECT E3 Ligase Involved in Self- and Host Ubiquitination To Influence Protein Interactions and Stability for Intracellular Survival. <i>Infection and Immunity</i> , 2017, 85, .	1.0	25
26	<i>Ehrlichia</i> Activation of Wnt-PI3K-mTOR Signaling Inhibits Autolysosome Generation and Autophagic Destruction by the Mononuclear Phagocyte. <i>Infection and Immunity</i> , 2017, 85, .	1.0	29
27	<i>Ehrlichia chaffeensis</i> Tandem Repeat Effector Targets Differentially Influence Infection. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 178.	1.8	28
28	Hacker within! <i>Ehrlichia chaffeensis</i> Effector Driven Phagocyte Reprogramming Strategy. <i>Frontiers in Cellular and Infection Microbiology</i> , 2016, 6, 58.	1.8	38
29	<i>Ehrlichia chaffeensis</i> TRP120 Activates Canonical Notch Signaling To Downregulate TLR2/4 Expression and Promote Intracellular Survival. <i>MBio</i> , 2016, 7, .	1.8	51
30	Rapid identification of ubiquitination and SUMOylation target sites by microfluidic peptide array. <i>Biochemistry and Biophysics Reports</i> , 2016, 5, 430-438.	0.7	12
31	<i>Ehrlichia chaffeensis</i> TRP32 Is a Nucleomodulin That Directly Regulates Expression of Host Genes Governing Differentiation and Proliferation. <i>Infection and Immunity</i> , 2016, 84, 3182-3194.	1.0	33
32	<i>Ehrlichia chaffeensis</i> Exploits Canonical and Noncanonical Host Wnt Signaling Pathways To Stimulate Phagocytosis and Promote Intracellular Survival. <i>Infection and Immunity</i> , 2016, 84, 686-700.	1.0	52
33	Detection of genotype-specific <i>Ehrlichia canis</i> exposure in Brazilian dogs by TRP36 peptide ELISA. <i>Ticks and Tick-borne Diseases</i> , 2016, 7, 142-145.	1.1	16
34	The Prostaglandin E2-EP3 Receptor Axis Regulates <i>Anaplasma phagocytophilum</i> -Mediated NLRC4 Inflammasome Activation. <i>PLoS Pathogens</i> , 2016, 12, e1005803.	2.1	31
35	<i>Ehrlichia chaffeensis</i> Exploits Host SUMOylation Pathways To Mediate Effector-Host Interactions and Promote Intracellular Survival. <i>Infection and Immunity</i> , 2014, 82, 4154-4168.	1.0	66
36	A novel <i>Ehrlichia</i> genotype strain distinguished by the TRP36 gene naturally infects cattle in Brazil and causes clinical manifestations associated with ehrlichiosis. <i>Ticks and Tick-borne Diseases</i> , 2014, 5, 537-544.	1.1	63

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37	Recombinant Ehrlichia P29 protein induces a protective immune response in a mouse model of ehrlichiosis. <i>Vaccine</i> , 2013, 31, 5960-5967.	1.7	8
38	Ehrlichia moonlighting effectors and interkingdom interactions with the mononuclear phagocyte. <i>Microbes and Infection</i> , 2013, 15, 1005-1016.	1.0	32
39	Ehrlichia chaffeensis TRP32 Interacts with Host Cell Targets That Influence Intracellular Survival. <i>Infection and Immunity</i> , 2012, 80, 2297-2306.	1.0	57
40	Molecular basis of antibody mediated immunity against Ehrlichia chaffeensis involves species-specific linear epitopes in tandem repeat proteins. <i>Microbes and Infection</i> , 2012, 14, 1054-1063.	1.0	31
41	Ehrlichia chaffeensis TRP120 Binds a G+C-Rich Motif in Host Cell DNA and Exhibits Eukaryotic Transcriptional Activator Function. <i>Infection and Immunity</i> , 2011, 79, 4370-4381.	1.0	59
42	Ehrlichia chaffeensis Tandem Repeat Proteins and Ank200 are Type 1 Secretion System Substrates Related to the Repeats-in-Toxin Exoprotein Family. <i>Frontiers in Cellular and Infection Microbiology</i> , 2011, 1, 22.	1.8	58
43	Ehrlichia chaffeensis TRP120 Interacts with a Diverse Array of Eukaryotic Proteins Involved in Transcription, Signaling, and Cytoskeleton Organization. <i>Infection and Immunity</i> , 2011, 79, 4382-4391.	1.0	66
44	Tyrosine-Phosphorylated Ehrlichia chaffeensis and Ehrlichia canis Tandem Repeat Orthologs Contain a Major Continuous Cross-Reactive Antibody Epitope in Lysine-Rich Repeats. <i>Infection and Immunity</i> , 2011, 79, 3178-3187.	1.0	24
45	Molecular and cellular pathobiology of Ehrlichia infection: targets for new therapeutics and immunomodulation strategies. <i>Expert Reviews in Molecular Medicine</i> , 2011, 13, e3.	1.6	63
46	Ehrlichia chaffeensis Transcriptome in Mammalian and Arthropod Hosts Reveals Differential Gene Expression and Post Transcriptional Regulation. <i>PLoS ONE</i> , 2011, 6, e24136.	1.1	56
47	Predominance of Ehrlichia chaffeensis in Rhipicephalus sanguineus ticks from kennel-confined dogs in Limbe, Cameroon. <i>Experimental and Applied Acarology</i> , 2010, 50, 163-8.	0.7	33
48	New insights into molecular Ehrlichia chaffeensis-host interactions. <i>Microbes and Infection</i> , 2010, 12, 337-345.	1.0	37
49	Molecular Characterization of Antibody Epitopes of Ehrlichia chaffeensis Ankyrin Protein 200 and Tandem Repeat Protein 47 and Evaluation of Synthetic Immunodeterminants for Serodiagnosis of Human Monocytotropic Ehrlichiosis. <i>Vaccine Journal</i> , 2010, 17, 87-97.	3.2	38
50	Progress and obstacles in vaccine development for the ehrlichioses. <i>Expert Review of Vaccines</i> , 2010, 9, 1071-1082.	2.0	26
51	Human Ehrlichiosis and Anaplasmosis. <i>Clinics in Laboratory Medicine</i> , 2010, 30, 261-292.	0.7	282
52	Mass Spectrometric Analysis of Ehrlichia chaffeensis Tandem Repeat Proteins Reveals Evidence of Phosphorylation and Absence of Glycosylation. <i>PLoS ONE</i> , 2010, 5, e9552.	1.1	19
53	Nuclear Translocated Ehrlichia chaffeensis Ankyrin Protein Interacts with a Specific Adenine-Rich Motif of Host Promoter and Intronic Alu Elements. <i>Infection and Immunity</i> , 2009, 77, 4243-4255.	1.0	94
54	Molecular and clinical evidence of Ehrlichia chaffeensis infection in Cameroonian patients with undifferentiated febrile illness. <i>Annals of Tropical Medicine and Parasitology</i> , 2009, 103, 719-725.	1.6	26

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55	Ehrlichia. , 2009, , 919-937.		2
56	An <i>Ehrlichia chaffeensis</i> Tandem Repeat Protein Interacts with Multiple Host Targets Involved in Cell Signaling, Transcriptional Regulation, and Vesicle Trafficking. <i>Infection and Immunity</i> , 2009, 77, 1734-1745.	1.0	73
57	Major Species-Specific Antibody Epitopes of the <i>Ehrlichia chaffeensis</i> p120 and <i>E. canis</i> p140 Orthologs in Surface-Exposed Tandem Repeat Regions. <i>Vaccine Journal</i> , 2009, 16, 982-990.	3.2	47
58	Ehrlichia. , 2009, , 117-164.		2
59	Emerging Pathogens: Challenges and Successes of Molecular Diagnostics. <i>Journal of Molecular Diagnostics</i> , 2008, 10, 185-197.	1.2	110
60	Genetic and Antigenic Diversities of Major Immunoreactive Proteins in Globally Distributed <i>Ehrlichia canis</i> Strains. <i>Vaccine Journal</i> , 2008, 15, 1080-1088.	3.2	50
61	A Variable-Length PCR Target Protein of <i>Ehrlichia chaffeensis</i> Contains Major Species-Specific Antibody Epitopes in Acidic Serine-Rich Tandem Repeats. <i>Infection and Immunity</i> , 2008, 76, 1572-1580.	1.0	61
62	Enzyme-Linked Immunosorbent Assay with Conserved Immunoreactive Glycoproteins gp36 and gp19 Has Enhanced Sensitivity and Provides Species-Specific Immunodiagnosis of <i>Ehrlichia canis</i> Infection. <i>Vaccine Journal</i> , 2007, 14, 123-128.	3.2	45
63	<i>Ehrlichia canis</i> gp200 Contains Dominant Species-Specific Antibody Epitopes in Terminal Acidic Domains. <i>Infection and Immunity</i> , 2007, 75, 4900-4908.	1.0	25
64	Identification of a Glycosylated <i>Ehrlichia canis</i> 19-Kilodalton Major Immunoreactive Protein with a Species-Specific Serine-Rich Glycopeptide Epitope. <i>Infection and Immunity</i> , 2007, 75, 74-82.	1.0	58
65	<i>Ehrlichia</i> Species in <i>Rhipicephalus sanguineus</i> Ticks in Cameroon. <i>Vector-Borne and Zoonotic Diseases</i> , 2007, 7, 221-227.	0.6	41
66	A preliminary investigation of <i>Ehrlichia</i> species in ticks, humans, dogs, and capybaras from Brazil. <i>Veterinary Parasitology</i> , 2007, 143, 189-195.	0.7	50
67	Restriction and expansion of <i>Ehrlichia</i> strain diversity. <i>Veterinary Parasitology</i> , 2007, 143, 337-346.	0.7	38
68	Differentially Expressed and Secreted Major Immunoreactive Protein Orthologs of <i>Ehrlichia canis</i> and <i>E. chaffeensis</i> Elicit Early Antibody Responses to Epitopes on Glycosylated Tandem Repeats. <i>Infection and Immunity</i> , 2006, 74, 711-720.	1.0	96
69	Analysis of <i>Ehrlichial</i> p28 Gene Expression in a Murine Model of Persistent Infection. <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 420-424.	1.8	7
70	Molecular Characterization of <i>E. canis</i> gp36 and <i>E. chaffeensis</i> gp47 Tandem Repeats among Isolates from Different Geographic Locations. <i>Annals of the New York Academy of Sciences</i> , 2005, 1063, 433-435.	1.8	32
71	<i>Ehrlichial</i> infection in Cameroonian canines by <i>Ehrlichia canis</i> and <i>Ehrlichia ewingii</i> . <i>Veterinary Microbiology</i> , 2005, 111, 59-66.	0.8	61
72	An Immunoreactive 38-Kilodalton Protein of <i>Ehrlichia canis</i> Shares Structural Homology and Iron-Binding Capacity with the Ferric Ion-Binding Protein Family. <i>Infection and Immunity</i> , 2005, 73, 62-69.	1.0	21

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73	Essential Role for Humoral Immunity during Ehrlichia Infection in Immunocompetent Mice. <i>Infection and Immunity</i> , 2005, 73, 8009-8016.	1.0	47
74	Detection of Medically Important Ehrlichia by Quantitative Multicolor TaqMan Real-Time Polymerase Chain Reaction of the dsb Gene. <i>Journal of Molecular Diagnostics</i> , 2005, 7, 504-510.	1.2	192
75	Rickettsia Species Infecting Amblyomma cooperi Ticks from an Area in the State of São Paulo, Brazil, Where Brazilian Spotted Fever Is Endemic. <i>Journal of Clinical Microbiology</i> , 2004, 42, 90-98.	1.8	522
76	Overproduction of TNF- α by CD8+ Type 1 Cells and Down-Regulation of IFN- γ Production by CD4+ Th1 Cells Contribute to Toxic Shock-Like Syndrome in an Animal Model of Fatal Monocytotropic Ehrlichiosis. <i>Journal of Immunology</i> , 2004, 172, 1786-1800.	0.4	115
77	Molecular Evidence for a Spotted Fever Group <i>Rickettsia</i> Species in the Tick <i>Amblyomma longirostre</i> in Brazil. <i>Journal of Medical Entomology</i> , 2004, 41, 533-537.	0.9	114
78	Histologic, Serologic, and Molecular Analysis of Persistent Ehrlichiosis in a Murine Model. <i>American Journal of Pathology</i> , 2004, 165, 997-1006.	1.9	55
79	Ehrlichia chaffeensis: a prevalent, life-threatening, emerging pathogen. <i>Transactions of the American Clinical and Climatological Association</i> , 2004, 115, 375-82; discussion 382-4.	0.9	29
80	Detection of Rickettsia africae in patients and ticks along the coastal region of Cameroon. <i>American Journal of Tropical Medicine and Hygiene</i> , 2004, 71, 363-6.	0.6	29
81	L-selectin and E-selectin expressed on monocytes mediating Ehrlichia chaffeensis attachment onto host cells. <i>FEMS Microbiology Letters</i> , 2003, 227, 303-309.	0.7	23
82	Novel Immunoreactive Glycoprotein Orthologs of Ehrlichia spp.. <i>Annals of the New York Academy of Sciences</i> , 2003, 990, 678-684.	1.8	32
83	Kinetics of Antibody Response to Ehrlichia canis Immunoreactive Proteins. <i>Infection and Immunity</i> , 2003, 71, 2516-2524.	1.0	68
84	Identification and Functional Analysis of an Immunoreactive DsbA-Like Thio-Disulfide Oxidoreductase of Ehrlichia spp.. <i>Infection and Immunity</i> , 2002, 70, 2700-2703.	1.0	40
85	Immunodiagnosis of Ehrlichia canis Infection with Recombinant Proteins. <i>Journal of Clinical Microbiology</i> , 2001, 39, 315-322.	1.8	37
86	Glycosylation of Homologous Immunodominant Proteins of Ehrlichia chaffeensis and Ehrlichia canis. <i>Infection and Immunity</i> , 2000, 68, 13-18.	1.0	72
87	Characterization of the complete transcriptionally active Ehrlichia chaffeensis 28 kDa outer membrane protein multigene family. <i>Gene</i> , 2000, 248, 59-68.	1.0	78
88	A conserved, transcriptionally active p28 multigene locus of Ehrlichia canis. <i>Gene</i> , 2000, 254, 245-252.	1.0	40
89	Molecular Cloning and Characterization of the 120-Kilodalton Protein Gene of <i>Ehrlichia canis</i> and Application of the Recombinant 120-Kilodalton Protein for Serodiagnosis of Canine Ehrlichiosis. <i>Journal of Clinical Microbiology</i> , 2000, 38, 369-374.	1.8	47
90	Primary and anamnestic responses of bovine bronchoalveolar and peripheral blood lymphocyte subsets to aerosolized Pasteurella haemolytica A1. <i>Veterinary Immunology and Immunopathology</i> , 1999, 67, 161-170.	0.5	6

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91	Evidence of <i>Pasteurella haemolytica</i> linked immune complex disease in natural and experimental models. <i>Microbial Pathogenesis</i> , 1999, 26, 183-193.	1.3	9
92	Molecular Cloning of the Gene for a Conserved Major Immunoreactive 28-Kilodalton Protein of <i>Ehrlichia canis</i> : a Potential Serodiagnostic Antigen. <i>Vaccine Journal</i> , 1999, 6, 392-399.	2.6	38
93	Genetic Diversity of the 28-Kilodalton Outer Membrane Protein Gene in Human Isolates of <i>Ehrlichia chaffeensis</i> . <i>Journal of Clinical Microbiology</i> , 1999, 37, 1137-1143.	1.8	51
94	Memory and CD8+ are the predominant bovine bronchoalveolar lymphocyte phenotypes. <i>Veterinary Immunology and Immunopathology</i> , 1997, 58, 55-62.	0.5	8
95	PCR Detection of Acute <i>Ehrlichia Canis</i> Infection in Dogs. <i>Journal of Veterinary Diagnostic Investigation</i> , 1996, 8, 441-447.	0.5	67
96	Detection of Humoral Antigen and Antibody by Enzyme-Linked Immunosorbent Assay in Horses with Experimentally Induced <i>Ehrlichia Equi</i> Infection. <i>Journal of Veterinary Diagnostic Investigation</i> , 1993, 5, 37-39.	0.5	18