Jun Liu

List of Publications by Year in descending order

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71102 95266 5,088 79 41 68 citations h-index g-index papers 82 82 82 5121 citing authors all docs docs citations times ranked

#	Article	IF	CITATIONS
1	Crosstalk between the ancestral type VII secretion system ESX-4 and other T7SS in Mycobacterium marinum. IScience, 2022, 25, 103585.	4.1	5
2	WhiB4 Is Required for the Reactivation of Persistent Infection of Mycobacterium marinum in Zebrafish. Microbiology Spectrum, 2022, 10, e0044321.	3.0	2
3	Structure of mycobacterial ATP synthase bound to the tuberculosis drug bedaquiline. Nature, 2021, 589, 143-147.	27.8	110
4	1H, 13C, and 15N resonance assignments of reduced apo-WhiB4 from Mycobacterium tuberculosis. Biomolecular NMR Assignments, 2021, 15, 99-101.	0.8	1
5	Early innate and adaptive immune perturbations determine long-term severity of chronic virus and Mycobacterium tuberculosis coinfection. Immunity, 2021, 54, 526-541.e7.	14.3	25
6	Xenogeneic Silencing and Bacterial Genome Evolution: Mechanisms for DNA Recognition Imply Multifaceted Roles of Xenogeneic Silencers. Molecular Biology and Evolution, 2021, 38, 4135-4148.	8.9	16
7	DNA binding mechanism of WhiB4 from Mycobacterium tuberculosis. Magnetic Resonance Letters, 2021, 2, 100010.	1.3	0
8	Interaction analysis of Mycobacterium tuberculosis between the host environment and highly mutated genes from population genetic structure comparison. Medicine (United States), 2021, 100, e27125.	1.0	3
9	Intranasal HD-Ad vaccine protects the upper and lower respiratory tracts of hACE2 mice against SARS-CoV-2. Cell and Bioscience, 2021, 11, 202.	4.8	13
10	Preclinical Progress of Subunit and Live Attenuated Mycobacterium tuberculosis Vaccines: A Review following the First in Human Efficacy Trial. Pharmaceutics, 2020, 12, 848.	4.5	8
11	Mycobacterial Virulence Factors: Surface-Exposed Lipids and Secreted Proteins. International Journal of Molecular Sciences, 2020, 21, 3985.	4.1	21
12	Mycobacterium tuberculosis 6C sRNA binds multiple mRNA targets via C-rich loops independent of RNA chaperones. Nucleic Acids Research, 2019, 47, 4292-4307.	14.5	50
13	Transcriptome Changes of Mycobacterium marinum in the Process of Resuscitation From Hypoxia-Induced Dormancy. Frontiers in Genetics, 2019, 10, 1359.	2.3	7
14	Recombinant BCG Overexpressing phoP-phoR Confers Enhanced Protection against Tuberculosis. Molecular Therapy, 2018, 26, 2863-2874.	8.2	15
15	How bacterial xenogeneic silencer rok distinguishes foreign from self DNA in its resident genome. Nucleic Acids Research, 2018, 46, 10514-10529.	14.5	23
16	Transcription factors Rv0081 and Rv3334 connect the early and the enduring hypoxic response of <i>Mycobacterium tuberculosis</i> . Virulence, 2018, 9, 1468-1482.	4.4	20
17	Acetylation of lysine 182 inhibits the ability of <i>Mycobacterium tuberculosis</i> DosR to bind DNA and regulate gene expression during hypoxia. Emerging Microbes and Infections, 2018, 7, 1-12.	6.5	21
18	Modulation of Central Carbon Metabolism by Acetylation of Isocitrate Lyase in Mycobacterium tuberculosis. Scientific Reports, 2017, 7, 44826.	3.3	36

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19	WhiB4 Regulates the PE/PPE Gene Family and is Essential for Virulence of Mycobacterium marinum. Scientific Reports, 2017, 7, 3007.	3.3	26
20	Immunogenicity and Protective Efficacy of a Fusion Protein Tuberculosis Vaccine Combining Five Esx Family Proteins. Frontiers in Cellular and Infection Microbiology, 2017, 7, 226.	3.9	12
21	The Impact of Genome Region of Difference 4 (RD4) on Mycobacterial Virulence and BCG Efficacy. Frontiers in Cellular and Infection Microbiology, 2017, 7, 239.	3.9	56
22	Loss of Lipid Virulence Factors Reduces the Efficacy of the BCG Vaccine. Scientific Reports, 2016, 6, 29076.	3.3	32
23	Variable Virulence and Efficacy of BCG Vaccine Strains in Mice and Correlation With Genome Polymorphisms. Molecular Therapy, 2016, 24, 398-405.	8.2	96
24	A Novel AT-Rich DNA Recognition Mechanism for Bacterial Xenogeneic Silencer MvaT. PLoS Pathogens, 2015, 11, e1004967.	4.7	53
25	Structural basis for targeting the ribosomal protein <scp>S</scp> 1 of <scp><i>M</i></scp> <i>ycobacterium tuberculosis</i> by pyrazinamide. Molecular Microbiology, 2015, 95, 791-803.	2.5	56
26	BCG Vaccines. Microbiology Spectrum, 2014, 2, MGM2-0028-2013.	3.0	30
27	Hiding behind the mycobacterial cell wall. Trends in Microbiology, 2014, 22, 110-112.	7.7	0
28	PhoY2 of Mycobacteria Is Required for Metabolic Homeostasis and Stress Response. Journal of Bacteriology, 2013, 195, 243-252.	2.2	23
29	Mechanism of DNA organization by Mycobacterium tuberculosis protein Lsr2. Nucleic Acids Research, 2013, 41, 5263-5272.	14.5	48
30	Genome Sequencing and Analysis of BCG Vaccine Strains. PLoS ONE, 2013, 8, e71243.	2.5	84
31	Targeting the global regulator Lsr2 as a novel approach for anti-tuberculosis drug development. Expert Review of Anti-Infective Therapy, 2012, 10, 1049-1053.	4.4	17
32	PPE38 Modulates the Innate Immune Response and Is Required for Mycobacterium marinum Virulence. Infection and Immunity, 2012, 80, 43-54.	2.2	81
33	Silencing of foreign DNA in bacteria. Current Opinion in Microbiology, 2012, 15, 175-181.	5.1	96
34	Both Phthiocerol Dimycocerosates and Phenolic Glycolipids Are Required for Virulence of Mycobacterium marinum. Infection and Immunity, 2012, 80, 1381-1389.	2.2	101
35	Activators of Cylindrical Proteases as Antimicrobials: Identification and Development of Small Molecule Activators of ClpP Protease. Chemistry and Biology, 2011, 18, 1167-1178.	6.0	86
36	MMAR_2770, a new enzyme involved in biotin biosynthesis, is essential for the growth of Mycobacterium marinum in macrophages and zebrafish. Microbes and Infection, 2011, 13, 33-41.	1.9	29

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37	Whole-Genome Sequences of Four Mycobacterium bovis BCG Vaccine Strains. Journal of Bacteriology, 2011, 193, 3152-3153.	2.2	28
38	Structural basis for recognition of AT-rich DNA by unrelated xenogeneic silencing proteins. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 10690-10695.	7.1	204
39	Malaria exacerbates experimental mycobacterial infection in vitro and in vivo. Microbes and Infection, 2010, 12, 864-874.	1.9	32
40	CD36 deficiency attenuates experimental mycobacterial infection. BMC Infectious Diseases, 2010, 10, 299.	2.9	48
41	Lsr2 is a nucleoid-associated protein that targets AT-rich sequences and virulence genes in <i>Mycobacterium tuberculosis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 5154-5159.	7.1	192
42	BCG Vaccines: Their mechanisms of attenuation and impact on safety and protective efficacy. Hum Vaccin, 2009, 5, 70-78.	2.4	131
43	Systematic Genetic Nomenclature for Type VII Secretion Systems. PLoS Pathogens, 2009, 5, e1000507.	4.7	233
44	MyBASE: a database for genome polymorphism and gene function studies of Mycobacterium. BMC Microbiology, 2009, 9, 40.	3. 3	14
45	Novel genome polymorphisms in BCG vaccine strains and impact on efficacy. BMC Genomics, 2008, 9, 413.	2.8	86
46	Spontaneous transposition of IS1096 or ISMsm3 leads to glycopeptidolipid overproduction and affects surface properties in Mycobacterium smegmatis. Tuberculosis, 2008, 88, 390-398.	1.9	41
47	Lsr2 of <i>Mycobacterium</i> Represents a Novel Class of H-NS-Like Proteins. Journal of Bacteriology, 2008, 190, 7052-7059.	2.2	109
48	Lsr2 of Mycobacterium tuberculosis is a DNA-bridging protein. Nucleic Acids Research, 2008, 36, 2123-2135.	14.5	84
49	Evidence for Pore Formation in Host Cell Membranes by ESX-1-Secreted ESAT-6 and Its Role in <i>Mycobacterium marinum</i> Escape from the Vacuole. Infection and Immunity, 2008, 76, 5478-5487.	2.2	260
50	Differential productions of lipid virulence factors among BCG vaccine strains and implications on BCG safety. Vaccine, 2007, 25, 8114-8122.	3.8	56
51	Identification of the lipooligosaccharide biosynthetic gene cluster from Mycobacterium marinum. Molecular Microbiology, 2007, 63, 1345-1359.	2.5	79
52	Tuberculosis: Current Treatment and New Drug Development. Anti-Infective Agents in Medicinal Chemistry, 2006, 5, 331-344.	0.6	9
53	The ESAT-6/CFP-10 secretion system of Mycobacterium marinum modulates phagosome maturation. Cellular Microbiology, 2006, 8, 1417-1429.	2.1	149
54	AsnB Is Involved in Natural Resistance of Mycobacterium smegmatis to Multiple Drugs. Antimicrobial Agents and Chemotherapy, 2006, 50, 250-255.	3.2	33

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55	Roles of Lsr2 in Colony Morphology and Biofilm Formation of Mycobacterium smegmatis. Journal of Bacteriology, 2006, 188, 633-641.	2.2	139
56	Mycobacterial Genomes., 2006,, 151-174.		0
57	LosA, a Key Glycosyltransferase Involved in the Biosynthesis of a Novel Family of Glycosylated Acyltrehalose Lipooligosaccharides from Mycobacterium marinum. Journal of Biological Chemistry, 2005, 280, 42124-42133.	3.4	62
58	PimF, a Mannosyltransferase of Mycobacteria, Is Involved in the Biosynthesis of Phosphatidylinositol Mannosides and Lipoarabinomannan. Journal of Biological Chemistry, 2004, 279, 18824-18833.	3.4	52
59	Impact of Methoxymycolic Acid Production by Mycobacterium bovis BCG Vaccines. Infection and Immunity, 2004, 72, 2803-2809.	2.2	31
60	Acquisition of Hrs, an Essential Component of Phagosomal Maturation, Is Impaired by Mycobacteria. Molecular and Cellular Biology, 2004, 24, 4593-4604.	2.3	90
61	Binding of ATP as Well as Tetrahydrofolate Induces Conformational Changes inLactobacillus caseiFolylpolyglutamate Synthetase in Solutionâ€. Biochemistry, 2003, 42, 1537-1543.	2.5	3
62	Mechanism of SMRT Corepressor Recruitment by the BCL6 BTB Domain. Molecular Cell, 2003, 12, 1551-1564.	9.7	251
63	Mycobacterium bovis BCG Vaccines Exhibit Defects in Alanine and Serine Catabolism. Infection and Immunity, 2003, 71, 708-716.	2.2	44
64	pH-induced Conformational Changes of AcrA, the Membrane Fusion Protein of Escherichia coli Multidrug Efflux System. Journal of Biological Chemistry, 2003, 278, 50474-50482.	3.4	39
65	A Rifampin-Hypersensitive Mutant Reveals Differences between Strains of Mycobacterium smegmatis and Presence of a Novel Transposon, IS1623. Antimicrobial Agents and Chemotherapy, 2003, 47, 3208-3213.	3.2	23
66	Central Region of the Human Splicing Factor Hprp3p Interacts with Hprp4p. Journal of Biological Chemistry, 2002, 277, 23764-23772.	3.4	30
67	Nramp1 Modifies the Fusion of Salmonella typhimurium-containing Vacuoles with Cellular Endomembranes in Macrophages. Journal of Biological Chemistry, 2002, 277, 2258-2265.	3.4	73
68	Cell Wall Structure of a Mutant of Mycobacterium smegmatis Defective in the Biosynthesis of Mycolic Acids. Journal of Biological Chemistry, 2000, 275, 7224-7229.	3.4	57
69	A mutant of Mycobacterium smegmatis defective in the biosynthesis of mycolic acids accumulates meromycolates. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 4011-4016.	7.1	93
70	Two Modes of Ligand Binding in Maltose-binding Protein of Escherichia coli. Journal of Biological Chemistry, 1997, 272, 17610-17614.	3.4	58
71	A Mycobacterium smegmatis mutant with a defective inositol monophosphate phosphatase gene homolog has altered cell envelope permeability. Journal of Bacteriology, 1997, 179, 7827-7833.	2.2	61
72	Active efflux of fluoroquinolones in Mycobacterium smegmatis mediated by LfrA, a multidrug efflux pump. Journal of Bacteriology, 1996, 178, 3791-3795.	2.2	148

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73	Mycolic Acid Structure Determines the Fluidity of the Mycobacterial Cell Wall. Journal of Biological Chemistry, 1996, 271, 29545-29551.	3.4	236
74	Fluidity of the lipid domain of cell wall from Mycobacterium chelonae Proceedings of the National Academy of Sciences of the United States of America, 1995, 92, 11254-11258.	7.1	135
75	Denaturant Unfolding of the Ferric Enterobactin Receptor and Ligand-Induced Stabilization Studied by Site-Directed Spin Labeling. Biochemistry, 1995, 34, 14230-14236.	2.5	44
76	A Site-Directed Spin-Labeling Study of Ligand-Induced Conformational Change in the Ferric Enterobactin Receptor, FepA. Biochemistry, 1994, 33, 13274-13283.	2.5	56
77	Permeability properties of a large gated channel within the ferric enterobactin receptor, FepA Proceedings of the National Academy of Sciences of the United States of America, 1993, 90, 10653-10657.	7.1	70
78	Formation of a gated channel by a ligand-specific transport protein in the bacterial outer membrane. Science, 1992, 258, 471-475.	12.6	185
79	BCG Vaccines. , 0, , 49-59.		0