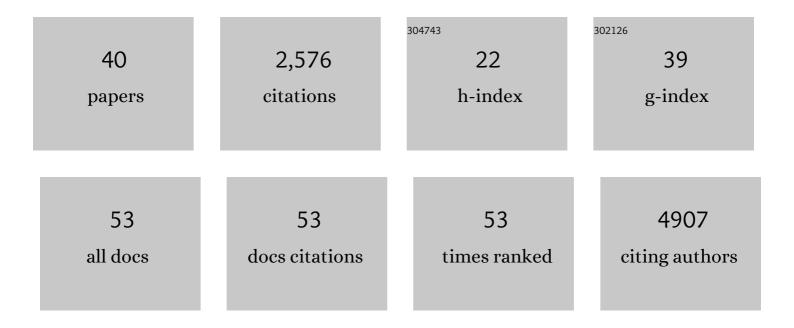
Christina L Stallings

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3711093/publications.pdf Version: 2024-02-01



CHRISTINA | STALLINGS

#	Article	IF	CITATIONS
1	Molecular dissection of RbpA-mediated regulation of fidaxomicin sensitivity in mycobacteria. Journal of Biological Chemistry, 2022, 298, 101752.	3.4	4
2	Perspectives and Advances in the Understanding of Tuberculosis. Annual Review of Pathology: Mechanisms of Disease, 2021, 16, 377-408.	22.4	26
3	Mycobacterium tuberculosis Rv3160c is a TetR-like transcriptional repressor that regulates expression of the putative oxygenase Rv3161c. Scientific Reports, 2021, 11, 1523.	3.3	6
4	UFMylation inhibits the proinflammatory capacity of interferon-γ–activated macrophages. Proceedings of the United States of America, 2021, 118, .	7.1	24
5	A novel class of TMPRSS2 inhibitors potently block SARS-CoV-2 and MERS-CoV viral entry and protect human epithelial lung cells. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	54
6	A Flexible and Deadly Way to Control Salmonella Infection. Immunity, 2020, 53, 471-473.	14.3	1
7	Select autophagy genes maintain quiescence of tissue-resident macrophages and increase susceptibility to Listeria monocytogenes. Nature Microbiology, 2020, 5, 272-281.	13.3	36
8	Editorial overview: Attrition warfare: host cell weapons against intracellular pathogens, and how the pathogens fight back. Current Opinion in Immunology, 2019, 60, vi-ix.	5.5	1
9	CarD contributes to diverse gene expression outcomes throughout the genome of <i>Mycobacterium tuberculosis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 13573-13581.	7.1	26
10	CarD and RbpA modify the kinetics of initial transcription and slow promoter escape of the Mycobacterium tuberculosis RNA polymerase. Nucleic Acids Research, 2019, 47, 6685-6698.	14.5	42
11	Chemical disarming of isoniazid resistance in <i>Mycobacterium tuberculosis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 10510-10517.	7.1	48
12	A platform for glycoengineering a polyvalent pneumococcal bioconjugate vaccine using E. coli as a host. Nature Communications, 2019, 10, 891.	12.8	60
13	Identification of 4-Amino-Thieno[2,3- <i>d</i>]Pyrimidines as QcrB Inhibitors in Mycobacterium tuberculosis. MSphere, 2019, 4, .	2.9	19
14	A promising bioconjugate vaccine against hypervirulent <i>Klebsiella pneumoniae</i> . Proceedings of the United States of America, 2019, 116, 18655-18663.	7.1	116
15	Roles for Autophagy Proteins in Immunity and Host Defense. Veterinary Pathology, 2018, 55, 366-373.	1.7	16
16	Domains within RbpA Serve Specific Functional Roles That Regulate the Expression of Distinct Mycobacterial Gene Subsets. Journal of Bacteriology, 2018, 200, .	2.2	16
17	<i>Irg1</i> expression in myeloid cells prevents immunopathology during <i>M. tuberculosis</i> infection. Journal of Experimental Medicine, 2018, 215, 1035-1045.	8.5	190
18	Bhlhe40 is an essential repressor of IL-10 during <i>Mycobacterium tuberculosis</i> infection. Journal of Experimental Medicine, 2018, 215, 1823-1838.	8.5	95

CHRISTINA L STALLINGS

#	Article	IF	CITATIONS
19	The stringent response and Mycobacterium tuberculosis pathogenesis. Pathogens and Disease, 2018, 76,	2.0	52
20	The impact of ISGylation during Mycobacterium tuberculosis infection in mice. Microbes and Infection, 2017, 19, 249-258.	1.9	40
21	Synthetic (p)ppGpp Analogue Is an Inhibitor of Stringent Response in Mycobacteria. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	47
22	Effects of Increasing the Affinity of CarD for RNA Polymerase on Mycobacterium tuberculosis Growth, rRNA Transcription, and Virulence. Journal of Bacteriology, 2017, 199, .	2.2	15
23	Host response: Inflammation promotes TB growth. Nature Microbiology, 2017, 2, 17102.	13.3	7
24	Rv0004 is a new essential member of the mycobacterial DNA replication machinery. PLoS Genetics, 2017, 13, e1007115.	3.5	21
25	Bacterial Pathogens versus Autophagy: Implications for Therapeutic Interventions. Trends in Molecular Medicine, 2016, 22, 1060-1076.	6.7	136
26	Characterization of phthiocerol and phthiodiolone dimycocerosate esters of M. tuberculosis by multiple-stage linear ion-trap MS. Journal of Lipid Research, 2016, 57, 142-155.	4.2	19
27	Cooperative stabilization of <i>Mycobacterium tuberculosis rrnA</i> P3 promoter open complexes by RbpA and CarD. Nucleic Acids Research, 2016, 44, gkw577.	14.5	29
28	Mycobacterium tuberculosis Transcription Machinery: Ready To Respond to Host Attacks. Journal of Bacteriology, 2016, 198, 1360-1373.	2.2	85
29	Analysis of the contribution of MTP and the predicted Flp pilus genes to Mycobacterium tuberculosis pathogenesis. Microbiology (United Kingdom), 2016, 162, 1784-1796.	1.8	12
30	The Cytosolic Sensor cGAS Detects Mycobacterium tuberculosis DNA to Induce Type I Interferons and Activate Autophagy. Cell Host and Microbe, 2015, 17, 811-819.	11.0	520
31	Unique role for ATG5 in neutrophil-mediated immunopathology during M. tuberculosis infection. Nature, 2015, 528, 565-569.	27.8	317
32	<i>Plasmodium</i> IspD (2-C-Methyl- <scp>d</scp> -erythritol 4-Phosphate Cytidyltransferase), an Essential and Druggable Antimalarial Target. ACS Infectious Diseases, 2015, 1, 157-167.	3.8	42
33	CarD stabilizes mycobacterial open complexes via a two-tiered kinetic mechanism. Nucleic Acids Research, 2015, 43, 3272-3285.	14.5	62
34	Phenotypic complementation of genetic immunodeficiency by chronic herpesvirus infection. ELife, 2015, 4, .	6.0	65
35	<scp>CarD</scp> integrates three functional modules to promote efficient transcription, antibiotic tolerance, and pathogenesis in mycobacteria. Molecular Microbiology, 2014, 93, 682-697.	2.5	31
36	Genome-wide mapping of the distribution of CarD, RNAP σA, and RNAP β on the Mycobacterium smegmatis chromosome using chromatin immunoprecipitation sequencing. Genomics Data, 2014, 2, 110-113.	1.3	14

#	Article	IF	CITATIONS
37	Catalytic and Non-Catalytic Roles for the Mono-ADP-Ribosyltransferase Arr in the Mycobacterial DNA Damage Response. PLoS ONE, 2011, 6, e21807.	2.5	15
38	Is Mycobacterium tuberculosis stressed out? A critical assessment of the genetic evidence. Microbes and Infection, 2010, 12, 1091-1101.	1.9	60
39	CarD Is an Essential Regulator of rRNA Transcription Required for Mycobacterium tuberculosis Persistence. Cell, 2009, 138, 146-159.	28.9	197
40	Exploring the Role of Low-Density Neutrophils During Mycobacterium tuberculosis Infection. Frontiers in Cellular and Infection Microbiology, 0, 12, .	3.9	8